

Nanomaterials for biology and medicine

Edited by Guannan Wang, Xiangzhao Ai and Pengfei Xu

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Nanomaterials for biology and medicine

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Editorial: Nanomaterials for biology and medicine

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KEYWORDS

nanomaterial, nanoprobes, nanoenzyme, biology, medicine

Editorial on the Research Topic Nanomaterials for biology and medicine

The desire for novel technology applications in biology, chemistry, engineering, physics, and medical science has fueled nanomaterials research. Nanomaterials have risen to prominence in technological breakthroughs due to their adjustable thermal, mechanical, electronic, and biological properties and superior performance over bulk equivalents (Baig et al., 2021; Mazari et al., 2021). Accordingly, numerous strategies have been projected to construct various functional structures by the integration of various types of nanomaterials, ranging from metals, metal oxides, alloys, and semiconductive materials to multifarious inorganic and organic polymers (Hao et al., 2010; Wagner et al., 2019). Specifically, nanomaterials offer the virtues as the therapeutic and diagnostic tools, which is owing to their small sizes, design flexibility, large specific surface area, and simple surface modification to enhance avidity for the targeting molecule (Yu et al., 2021; Qiao et al., 2022; Pourmadadi et al., 2023). There has been an explosive development in the use of nanomaterials for biomedical applications to probe biological processes, diagnose and treat medical conditions. The distinguishing characteristics of nanomaterials enable them to preferentially penetrate and be retained by cells and tissues of living organisms, thereby achieving developments in new approaches to target in biological cells and biological tissues. Additionally, the size scale of nanomaterials will provide the inspiration to build complexity into the nanoprobes that endow them versatility with both diagnostic, therapy, and drug delivery function. Moreover, diversified surface modification strategies have broadened the application of nanomaterials for biomedical applications, which adds the property of stability, biocompatibility, biodistribution, solubility, biological or therapeutic effects (Sztandera et al., 2019; Liu et al., 2021).

In this Research Topic, the authors were invited to contribute their research works, which enable the better explanation on the recent advances in diversified applications of nanomaterial in the biology and medicine. Yu et al. have contributed a literature review on the current progress on the nanomaterials' application in diagnosing and treating of glioblastoma (GBM). They discussed the nanomaterials-based nano-diagnosis or treatment mechanisms. Additionally, the advanced application progress of nanomaterial combination diagnostic and therapeutic tools for GBM was summarized. Nano-catalytic therapy, acting as an innovative strategy, has been extensively explored. Shi et al. group synthesized a multi-functional magneto-gold nano-enzyme AuNC@Fe₃O₄ and evaluated their anti-cancer ability in the hepatocellular carcinoma (HCC cells) *in vitro*. The preparative nano-enzyme AuNC@Fe₃O₄ with a small size was characterized using various techniques and demonstrated with high peroxidase (POD)-like activity, good photothermal conversion

efficiency, and can inhibit cell proliferation and enhance cell apoptotic ability in cancer cells, providing a potential anti-cancer method for HCC. Responding to the oxidation state of the microenvironment of bacterial infection, Dorma Momo et al. explored a near-infrared (NIR) photothermal bacterial inactivation by reasonably designing a Metal organic Framework (MOF)-based nano-composite, offering a novel inspiration for constructing precise nano-therapeutic systems. They systematically studied the strong deactivation effect on the Gram-negative and -positive bacteria and intense therapeutic effectiveness on the mouse skin wound infection model of the designed nano-system.

Native and synthetic nanomaterials have drawn a lot of research interests and been projected as the key components for construction of drug delivery systems for healing patients in clinical. Ravelo-Nieto et al. constructed a cellular drug delivery system by using silanized fullerenol and silica nanoparticles (SN) as the nano-structured supports to conjugate potent cell-penetrating agents. The nanobioconjugates showed distinct intracellular trafficking and endosomal escape behavior in the cell lines, which indicated the potentiality to address the challenge of cytoplasmic drug delivery and the development of therapeutic methods for lysosome storage disease. You et al. developed a monodispersed and biocompatible mesoporous SN (MSN) divergent porous channel for loading dapagliflozin (DAPA). They constructed a drug delivery system through the surface-modification of the cardiac-targeting peptides to release drug for the hypoxic and weak acid damaged cardiomyocytes. This MSN-based nanocarriers for the DAPA delivery system can achieve the efficacious cardiac repair and regeneration in vivo. Gui et al. research group exploit a pH responsive antibacterial delivery system (Imi@ZIF-8) for the antibiosis treatment of A. baumannii. They found that at an imipenem concentration of 10 mg/kg, the Imi@ZIF-8 nanosystem manifested the outstanding therapeutic efficacy against A. baumannii in the mice with celiac disease.

Intended as a one-stop reference, this Research Topic provides the reader with the most-up-to-date and comprehensive exploration of a variety of the nanomaterial applications for biology and medicine. Briefly, these nanomaterials can be applied to perspective to advance in intrinsic nano-therapy across the biomedical area, from cancer therapeutics to microbial infection treatment and tissue regeneration. As such, this Research Topic provides the most comprehensive coverage of this intriguing field of study. Currently, this Research Topic has published 11 articles and received over 2,900 downloads and 14,000 readings worldwide.

Author contributions

The author confirms being the sole contributor of this work and has approved it for publication.

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Recent advances in enzyme-related biomaterials for arthritis treatment

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Arthritis is a group of highly prevalent joint disorders, and osteoarthritis (OA) and rheumatoid arthritis are the two most common types. The high prevalence of arthritis causes severe burdens on individuals, society and the economy. Currently, the primary treatment of arthritis is to relieve symptoms, but the development of arthritis cannot be effectively prevented. Studies have revealed that the disrupted balance of enzymes determines the pathological changes in arthritis. In particular, the increased levels of matrix metalloproteinases and the decreased expression of endogenous antioxidant enzymes promote the progression of arthritis. New therapeutic strategies have been developed based on the expression characteristics of these enzymes. Biomaterials have been designed that are responsive when the destructive enzymes MMPs are increased or have the activities of the antioxidant enzymes that play a protective role in arthritis. Here, we summarize recent studies on biomaterials associated with MMPs and antioxidant enzymes involved in the pathological process of arthritis. These enzyme-related biomaterials have been shown to be beneficial for arthritis treatment, but there are still some problems that need to be solved to improve efficacy, especially penetrating the deeper layer of articular cartilage and targeting osteoclasts in subchondral bone. In conclusion, enzyme-related nano-therapy is challenging and promising for arthritis treatment.

KEYWORDS

biomaterials, nano-therapy, arthritis, matrix metalloproteinases, endogenous antioxidant enzymes

Introduction

Arthritis, which is a group of musculoskeletal diseases, is one of the leading causes of disability in the elderly population (Woolf and Pfleger, 2003). Osteoarthritis (OA) and rheumatoid arthritis (RA) are the most prevalent types of arthritis and affected 344 million people and 13 million people, respectively, globally in 2019 (Cieza et al., 2021). OA is characterized by joint degeneration, especially in the knee, and involves multiple joints, such as the hand, hip, knee and foot. A large-scale survey in the

United Kingdom in 2017 showed that the prevalence of OA in adults was 10.7% (Swain et al., 2020-06). The increases in obesity and the ageing population contribute to the prevalence of OA (Briggs et al., 2020-10). RA is an immunization-induced systemic disease characterized by synovial inflammation and joint destruction, and the prevalence of RA is 0.5–1.0% in the US (Palmer et al., 2019).

OA and RA are both inflammatory joint diseases that involve joint and synovial destruction and immune cell infiltration (Zhang et al., 2019-03) and is associated with joint pain, swelling, and limited movement, resulting in a decline in physical function, increased dependence and reduced quality of life. Furthermore, the prevalence of OA and RA is expected to increase significantly as the global population ages. The treatment of arthritis is often a long and complex process due to irreversible damage and the risk of comorbidities, resulting in extremely high medical and economic burdens on society, and these burdens continue to increase globally (Briggs et al., 2020-10).

To date, there is no effective cure for OA or RA. The current interventions include medications, physical therapy, and surgical intervention, all of which are aimed at alleviating symptoms and reducing joint damage and disability. Medications for OA, including topical, oral and intra-articular (IA) injectable drugs, are palliative and limited to controlling symptoms of joint swelling, pain and stiffness (Tschon et al., 2020-06). A randomized clinical trial has even shown that IA corticosteroids may accelerate the destruction of articular cartilage (McAlindon et al., 2017-05). Currently, non-steroidal anti-inflammatory drugs (NSAIDs), glucocorticoids (GCs) and disease-modifying anti-rheumatic drugs (DMARDs) are mainly used in the clinical treatment of RA. The targets of traditional DMARDs are not clear, and approximately 30%–50% of patients respond poorly to these drugs (Sparks, 2019). As a result of the poor bioavailability and short half-lives of anti-rheumatic drugs, prolonged repeated use can cause serious adverse reactions such as vomiting, drug resistance and bone marrow suppression.

Physical therapy for OA and RA includes weight loss, moderate exercise and knee joint distraction. Knee joint distraction can improve symptoms and promote tissue repair in severe knee joint degeneration, but there is frequent infection during the follow-up (Jansen and Mastbergen, 2022-01; van der Woude et al., 2017-01). When conservative treatment is not feasible for end-stage arthritis, surgical intervention, such as total joint replacement, can be considered, but this treatment strategy is related to persistent postsurgical pain and infection (Wylde et al., 2011-03; Chung et al., 2021-11).

Currently, new therapeutic strategies and drugs primarily alleviate symptoms to treat arthritis, and critically unsolved problems, such as how to restore abnormal cellular function in arthritis, should be considered. Cellular activity depends on various proteins, and some of these proteins are important enzymes for physiological and pathological processes. Herein, we summarized the essential enzymes that are involved in pathological changes in arthritis.

Arthritis-related enzymes

The pathological changes in OA and RA are mainly characterized by cartilage destruction and synovial inflammation (Trachana et al., 2019; Scherer et al., 2020). Cell metabolism is often regulated by different enzymes, and abnormal levels of enzymes are typically associated with the occurrence of various diseases. In cartilage, different matrix metalloproteinases (MMPs) are responsible for destroying chondrocytes by degrading collagen and proteoglycans.

1 matrix metalloproteinases linked with arthritis

There is increasing evidence that these inflammatory mediators are involved in the pathogenesis of both OA and RA (Malemud, 2017; van Dalen et al., 2017). Neutrophils, monocytes and macrophages infiltrate cartilage and synovial tissue after inflammation occurs, releasing various inflammatory factors and chemokines, which cause an increase in MMPs.

The destruction or degradation of articular cartilage is regulated by MMPs, which are a family of proteolytic enzymes that hydrolyse extracellular matrix (ECM). Different types of MMPs are involved in degrading proteoglycans and collagens, which are the main components of ECM in cartilage, especially MMP-1, MMP-2, MMP-3, MMP-9 and MMP-13 (Itoh, 2017; Mehana et al., 2019). MMPs can degrade collagen, elastin, and other substances in the ECM of articular cartilage that maintain the structure of cartilage and ultimately destroy the integrity of ECM structure and function.

Under pathological conditions, the expression level of MMP-1 was significantly increased in OA and RA, and this factor degraded ECM collagen and mediated cartilage destruction (Wang et al., 2020a). In cartilage and synovium, MMP-1 expression increased steadily during the progression of OA in a rabbit model of anterior cruciate ligament transection (ACLT) (Wu et al., 2008). MMP-1 could lead to the degeneration of primary collagen (type II collagen) in cartilage, and this effect was irreversible (Macdonald et al., 2018).

The development of OA and RA is associated with the increased secretion and activity of MMP-2 in synovial cells and the joints of RA patients, respectively (Kim et al., 2011; Galasso et al., 2012). Furthermore, MMP-2-sensitive peptide was shown to be specifically released in inflammatory joints *in vitro* and *in vivo*, which might be an important approach for drug-targeted treatment of RA (Yu et al., 2022).

Significantly increased levels of MMP-3 in the serum of OA patients were positively correlated with the severity of knee OA and RA in patients (Ma et al., 2014; Georgiev et al., 2018; Pengas et al., 2018). Furthermore, serum MMP-3 levels were closely

correlated with disease activity scores, suggesting that serum MMP-3 levels could be used as an indicator of structural damage and monitor disease progression (Galil et al., 2016; Tuncer et al., 2019).

MMP-9 was also positively correlated with disease severity in OA patients (Lipari and Gerbino, 2013). A meta-analysis showed that MMP-2 and MMP-9 protein expression levels were significantly higher in the OA group than in the control group, indicating that MMP-2 and MMP-9 are involved in the pathogenesis of OA (Zeng et al., 2015). Multiple studies have shown that the expression of MMP-9 in synovial fluid and synovial cells of RA patients is increased (Silosi et al., 2015; Ma et al., 2019). The degree of inflammation in RA patients correlated with Toll-like receptor 2 (TLR2) expression in peripheral blood monocytes. The increased expression of TLR2 led to the increased expression of MMP-9 (Chen et al., 2015). MMP-9 could participate in the synovial cell-mediated inflammatory response and the degeneration of ECM, especially proteoglycans, which might directly cause joint destruction (Metzger et al., 2012).

MMP-13 is a crucial enzyme leading to the degradation of collagen types I, II and III and the cartilage proteoglycan aggrecans and is considered a significant factor in the pathogenesis of OA (Fosang et al., 1996). MMP-13 attracted much attention due to its obvious overexpression in the articular cartilage of OA patients, but it was almost undetectable in normal adult tissues (Kaneva, 2022). Interfering with the expression of MMP-13 in a surgically induced OA model could efficiently alleviate OA severity (Hoshi et al., 2017). Given its critical role in ECM degradation, MMP-13 has been a promising target in OA treatment (Hu and Ecker, 2021). K/BxN serum-induced arthritis increases MMP-13 expression in C57BL/6 mice, and MMP-13-deficient (MMP-13-/-) mice exhibit reduced inflammation and joint destruction (Singh et al., 2013). In addition, MMP-13 was also associated with the progression of RA, providing crucial predictive information about future structural damage and severity in early RA patients (Tatematsu et al., 2018).

2 Endogenous antioxidant enzymes linked with arthritis

Apart from the direct effect of MMPs on ECM degradation in cartilage and promoting the progression of arthritis, endogenous antioxidants such as superoxide dismutases (SODs), glutathione peroxidase (GPx), catalase (CAT), and glutathione reductase (GR) also affect the occurrence of arthritis by scavenging intracellular reactive oxygen species (ROS) and alleviating cellular oxidative stress.

ROS are key signalling molecules in the progression of inflammatory diseases (Mittal et al., 2014). Under inflammatory conditions, the oxidative stress induced by macrophages, monocytes, and neutrophils leads to the formation of interendothelial junctions, accelerating the crossing of the endothelial barrier and ultimately promoting inflammation (You et al., 2018).

The levels of intra-articular ROS (including H2O2, O2-, OH-, and HOCl) are significantly increased in OA patients, while ROS are maintained at low levels in normal articular tissue (Lepetsos and Papavassiliou, 2016; Yao et al., 2019). The overproduction of ROS causes overoxidation, protein carbonylation, and DNA damage and is considered the primary mechanism of chondrocyte loss and tissue damage (Hosseinzadeh et al., 2016). The associated ROS, including nitric oxide (NO), superoxide anion (O_2^-) and hydrogen peroxide (H_2O_2) , are present in the articular cavities of RA patients in large quantities (Datta et al., 2014). When the local inflammatory response in RA joints is accelerated and ROS levels exceed physiological tolerance, they not only damage proteins, lipids, and nucleic acids but also act as important endogenous signalling regulators that amplify the synovial inflammatory response (Bala et al., 2017; Phull et al., 2018). Li et al. found that ROS significantly promoted the proliferation of RA synovial fibroblasts and the production of inflammatory factors and that inhibiting ROS significantly downregulated the inflammatory factors secreted by RA synovial fibroblasts, ultimately improving RA conditions (Li et al., 2018). Therefore, a potent antioxidant compound that can reduce ROS in inflammatory cells may be a key factor in the treatment of chronic inflammatory diseases.

ROS clearance is regulated by SODs, GPx, CAT and GR (He et al., 2017). CAT and GPx are involved in the decomposition of intracellular hydrogen peroxide and maintain normal ROS levels to reduce toxic reactions. SOD can catalyse O_2^- into O_2 and H_2O_2 . GR catalyses the reduction of glutathione disulfide (GSSG) to the sulfhydryl form of glutathione (GSH), which plays an important role in the tissue oxidative stress response (Deponte, 2013). The levels of SOD, CAT and other antioxidant enzymes in OA chondrocytes were significantly lower than those in normal chondrocytes, indicating that insufficient antioxidant capacity might cause cartilage damage (Zhuang et al., 2018). Unlike the expression pattern of other antioxidant enzymes, the expression of GR was increased in arthritis (Meshkibaf et al., 2019; Idzik et al., 2022).

The proliferation and activation of osteoclasts (OCs) are key factors leading to bone damage and bone metabolism disorders in RA (Auréal et al., 2020). Recent studies have shown a close correlation between bone destruction and oxidative stress in the pathogenesis of RA. ROS promote osteoclast differentiation (Gamal et al., 2018). Decreased expression of SOD, CAT and GPx was found in the ankle joints of RA rats (Ren et al., 2019a). ROS-induced peroxidation is inhibited by antioxidant enzymes, among which superoxide dismutase 3 (SOD3) is the key enzyme that protects cells from oxidative stress (Nguyen et al., 2020). SOD3 reduced proinflammatory cytokines (IL-1 β , IL-2, IL-4, and TNF- α) and the release of MMPs (MMP-2, MMP-3 and MMP-9), ultimately inhibiting inflammatory responses (Xie et al.,



2021). Icariin protects synoviocytes induced by lipopolysaccharide (LPS) by inhibiting ferroptosis by activating the Xc/GPX4 axis (Luo and Zhang, 2021).

Considering the importance of MMPs and oxide reductase associated with ROS in the occurrence of arthritis, biomaterials that target endogenous enzymes have become a hot research topic in recent years. Next, we will introduce the application of biomaterials that are linked with these enzymes.

Nanotherapies that target enzymes in arthritis

Enzymes that play critical roles in arthritis pathology are categorized into two groups according to their expression characteristics: upregulated enzymes and downregulated enzymes, which are listed in Figure 1. Enzyme homeostasis is critical for the human body. Both the upregulated and downregulated expression of these enzymes disrupt the balance of cell metabolism and can cause diseases. Therefore, therapeutic strategies have been designed according to the expression of these enzymes. If the expression of these enzymes is upregulated, nanomaterials can respond and release an effective drug to inhibit pathological changes, or nanomaterials can be fabricated to simulate the effects of downregulated enzymes. Next, we described two different functional enzymes in arthritis treatment.

1 Nanomaterials associated with upregulated enzymes in arthritis treatments

It is well known that MMPs are the main destructive enzymes in chondrocytes that degrade ECM components, such as proteoglycans and collagen networks. Degradation of the ECM leads to functional destruction of chondrocytes and cartilage erosion. Therefore, MMPs have become an important molecular target for studies on the treatment of OA. In particular, MMP-13, a critical protease in chondrocytes, is responsible for the degradation of type II collagen and proteoglycans.

OA is a chronic inflammatory disease. Growing evidence reveals that the changes in the OA microenvironment include excessive inflammation and MMP overexpression (Latourte et al., 2017; Li et al., 2017; Stocco et al., 2019). The microenvironment is an important factor in maintaining joint homeostasis. Long-term inhibition of MMP enzymatic activity may lead to adverse reactions. Therefore, it is necessary to design materials that are highly selective for MMPs and can adapt to the changes in MMP levels *in vivo*. When MMP expression is upregulated, MMP-responsive nanoparticles (NPs) work, and they are inactive when MMPs are at low levels.

The increased expression of MMPs in inflamed tissues may be a promising breakthrough for arthritis therapy. A commercially available, Food and Drug Administration (FDA)-approved molecule known as triglycerol monostearate (TGMS) has been shown to be responsive to MMPs (Wen et al., 2019). MMP-responsive PEGylated lipid NPs (TGMS/DSPE-PEG2000 NPs) can be produced through the coassembly of TGMS and 1,2-distearoyl-sn-glycero-3-phosphoethanolamine-poly (ethylene glycol) (DSPE-PEG2000). Dexamethasone (Dex)-loaded MMP-responsive NPs were obtained by loading TGMS/DSPE-PEG2000 NPs with Dex, and Dex can be rapidly released from the lipid NPs after TGMS is cleaved by MMP-3 and MMP-9. Dex-loaded MMP-responsive NPs significantly reduced the degree of joint swelling and inhibited the production of TNF- α and IL-1 β in the joint (He et al., 2020).

In another study, the nanozyme-like role of the hydrogel form of TGMS(TG-18) was further confirmed in RA treatment. A hydrogel platform that exhibits disassembly and drug release controlled by the concentration of enzymes during arthritis flares was constructed. In this study, a triglycerol monostearate hydrogel (TG-18) loaded with the corticosteroid triamcinolone acetonide (TA) exhibited drug release in response to the increased activities of arthritis-related enzymes *in vitro* (MMP-2, MMP-3, MMP-9) or synovial fluid from patients with RA (Joshi et al., 2018).

In addition to synovial inflammation and joint swelling, obvious cartilage damage and bone erosion are often observed in RA. Synovial macrophages mediate joint inflammation once activated, and OCs are responsible for arthritic bone erosion and resorption of the bone matrix. Both OCs and synovial macrophages express high levels of avß3 integrin, which plays an important role in activated macrophage-dependent inflammation and OC-dependent bone resorption. Macrophages and OCs fail to undergo apoptosis in the RA joint, leading to persistent inflammation and joint destruction. Therefore, inducing OC and macrophage apoptosis in RA joints represents a promising strategy for advanced RA treatment. According to the characteristics of OCs and synovial macrophages, novel CEL-loaded PRNPs (CEL-PRNPs) were synthesized that contained celastrol (CEL), which can induce apoptosis in OCs and macrophages, RGD, which is a ligand of avß3 that targets OCs and inflammatory macrophages, and polyethylene glycol (PEG), which is cleaved by MMP-9. In an adjuvant-induced arthritis rat model, CEL-PRNPs efficiently reduced the number of OCs and inflammatory macrophages and relieved various symptoms, including ankle and paw swelling and bone erosion, in the inflamed joints of AIA rats with advanced arthritis (Deng et al., 2021).

To determine the inflammatory condition and investigate the therapeutic effects of MMP-responsive biomaterials, fluorescence imaging was considered for diagnosis and therapy.

Inflamed cartilage is characterized by MMP-13 overexpression and an acidic microenvironment. Therefore, MMP-13/pH-responsive ferritin nanocages (CMFn) loaded with an anti-inflammatory drug (hydroxychloroquine, HCQ), termed CMFn@HCQ, were constructed for OA imaging and therapy. CMFn is a marker for imaging diagnosis that emits light in response to MMP-13 overexpression. The intensity of CMFn light increases with the severity of OA. However, in normal joints, this compound emits no light. The release of HCQ causes an anti-inflammatory effect in OA joints to reduce synovial inflammation, and the retention time lasts up to 14 days (Chen et al., 2019a).

Cartilage-targeting C-PPL was created by grafting collagen type II-targeting peptides with the sequence WRYGRL onto the polymer poly (2-ethyl-2-oxazoline)-poly (ε-caprolactone) (PPL). Additionally, PPL was conjugated with a specific peptide substrate of the MMP-13 enzyme (H2N-GPLGVRGC-SH) that was labelled with a fluorescent dye (Cy5.5) and was subsequently coupled with the black hole quencher-3 (BHQ-3) that can quench Cy5.5 fluorescence to obtain an MMP-13responsive and pH-sensitive polymer (MR-Cy5.5-BHQ-3-PPL). A cartilage-targeting and OA-specific theragnostic nanoplatform (MRC-PPL) was obtained by the self-assembly of C-PPL and MR-PPL. Finally, MRC-PPL was loaded with the traditional Chinese medicine psoralidin (PSO) to form MRC-PPL@PSO nano-micelles, which specifically target and protect cartilage (Lan et al., 2020). The synthesis and mechanism of MRC-PPL@PSO nano-micelles to treat OA are shown in Figure 2.

In addition to MMP overexpression in arthritis tissue, the intrinsic properties of the OA microenvironment, especially synovial fluid, are also considered when designing novel nanomaterials. The increased activity of the GR enzyme was reported in the synovial fluid of RA and OA patients, and a selectively controlled drug release that is sensitive to the GR enzyme was designed for the treatment of arthritic diseases (Ostalowska et al., 2006; Sredzińska et al., 2009). Polymeric micelles were made of methoxypolyethylene glycol amineglutathione-palmitic acid (mPEG-GSHn-PA) polymers. Dex was loaded into the cores of the polymeric micelles. The release of Dex was slow under physiological conditions, while the presence of the GR enzyme stimulated a burst release via a thiol-disulfide exchange between GSH and GSSG (Lima et al., 2021). The above biomaterials associated with MMPs are listed in Table 1.

2 Nanomaterials associated with downregulated enzymes in arthritis treatments

Apart from the destruction of cartilage tissue induced by the increased expression of MMPs, the decreased expression of oxide reductase associated with ROS showed a similar effect on cartilage. To reduce the expression of oxide reductase, the strategy was to supply these enzymes directly or mimic the activities with special biomaterials.

Supplementation with antioxidant enzymes such as SOD has been shown to be effective in treating arthritis. Chitosan was chemically conjugated with SOD to generate the nanoparticlelike conjugate 6-O-2'-hydroxylpropyltrimethyl ammonium chloride chitosan-SOD (O-HTCC-SOD), which was superior to unmodified SOD in bioavailability, prolonged half-life and

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residence in the rat joint cavity. After IA injection of O-HTCC-SOD into rats with MIA-induced OA, mechanical allodynia was greatly reduced, and changes in the gross morphological and histological lesions of articular cartilage were dramatically inhibited (Wang et al., 2020b).

Although the nanosized conjugate O-HTCC-SOD has exhibited higher enzyme activity and superior membrane permeability to native SOD, natural enzymes are unstable, expensive and difficult to store. Currently, biomaterials called nanozymes have been designed to mimic the effects of these oxide reductases. Nanozymes are a specific kind of nanomaterial that have the activities of intrinsic enzymes and possess unique advantages, such as high efficiency, increased compatibility with specific environments, such as high temperatures and pH variations, cyclic use, and a large surface area, and these materials can be conjugated to multiple ligands to achieve multifunctionality. These features give rise to their promising applications in a variety of fields (Pirmohamed et al., 2010).

Recently, numerous nanomaterials with enzyme-like properties have been discovered for OA treatment, including metals, metal oxides, and carbon-based materials. As a representative metal oxide, cerium oxide has been evaluated in RA treatment. Engineered cerium oxide (CeO₂) nanoparticles (CeONPs), which are also known as nanoceria, have attracted much attention for exhibiting SOD⁻, CAT⁻, and oxidase-like activity (Heckert et al., 2008; Baldim et al., 2018; Kalashnikova et al., 2020). In reduction reactions, SOD catalyses $O_2^{\bullet-}$ into H_2O_2 , which may undergo catalysis by CAT into H_2O .

Given that albumin is a natural protein and scavenging receptors are widely distributed in the inflamed joints of RA, albumin-nanoceria NPs (A-nanoceria) were synthesized by connecting albumin to nanoceria and further conjugated with near-infrared, indocyanine green (ICG) dye. Enzymatic properties and ROS scavenging activities against a monocyte cell line and systemic targeting potential were evaluated in a collagen-induced arthritis (CIA) mouse model. Such a design has the advantages of targeting inflammation, assessing severity, and controlling inflammation with imaging guidance in RA (Kratschmer et al., 1990).

Moreover, carbon-based materials have also exhibited the activities of nanozymes in scavenging ROS. Fullerene (C60) is a

Arthritis	Enzymes-responsive group	Nanomaterial	Platform	Components	Responsive enzymes	Drug	Ref
RA	TGMS	Dex-loaded TGMS/DSPE- PEG2000	NPs	1.TGMS	MMP-3	Dex	He et al. (2020)
				2.Dex	MMP-9		
				3.PEG2000			
				4.DSPE			
RA	TGMS	TA-loaded TG-18 hydrogel	hydrogel	1.TGMS	MMP-2	ТА	Joshi et al.
				2.TA	MMP-3		(2018)
					MMP-9		
RA	PEG	CEL-PRNPs	NPs	1.celastrol (CEL)	MMP-9		Deng et al
				2.RGD			(2021)
				3.PEG			
				4.PLGA			
OA	H2N-GPLGVRGC-SH	CMFn@HCQ	nanocages	1.MMP-13 cleavble peptide	MMP-13	HCQ	Chen et a (2019a)
				2.HCQ			
				3.collagen type II targeting peptides			
				4.BHQ3			
				5.Cy5.5			
				6.ferrritin			
OA	H2N-GPLGVRGC-SH	MRC-PPL@PSO	micelles	1.MMP-13 cleavble peptide	MMP-13	PSO	Lan et al. (2020)
				2.PSO			
				3.collagen type II targeting peptides			
				4.PPL			
				5.Cy5.5			
				6.BHQ-3			
OA	GSSG	Dex-loaded mPEG-	micelles	1.PEG	GR enzyme	Dex	Lima et al
		GSHn-PA		2.GSH			(2021)
				3.PA			
				4.Dex			

TABLE 1 The biomaterials that target the upregulated enzymes in arthritis.

spherical carbon molecule with a unique cage structure that functions as a free-radical scavenger. Apart from inhibiting ROSinduced catabolism in cartilage, fullerene also decreases friction on the cartilage surface and subsequently prevents the further development of cartilage degeneration. With these advantages, fullerene has been used to synthesize biomaterials for the treatment of arthritis. For example, fullerene-like MoS₂ (F-MoS₂) NPs are efficient lubricants and antioxidants for artificial synovial fluid. These NPs possess intrinsic dualenzyme-like activity, mimicking SOD and CAT under physiological conditions (pH 7.4, 25°C) and regulating the ROS level in artificial synovial fluid containing HA (Chen et al., 2019b).

Prussian blue (PB) has been approved by the U.S. FDA as a commonly used dye and medicine due to its excellent biocompatibility and biosafety. The peroxidase, CAT, and

SOD activities of PBzymes mediate the scavenging of \bullet OH, \bullet OOH, and H₂O₂, exhibiting outstanding anti-inflammatory and antioxidative bioactivities (Long et al., 2016; Zhang et al., 2016; Dacarro et al., 2018; Qin et al., 2018).

A hollow PBzyme (HPBzyme) with a mesopore structure and a high specific surface area was produced that could remodel the OA microenvironment by mitigating the inflammatory response, protecting against chondrocyte ECM degradation, and exhibiting therapeutic efficacy *in vivo* (Hou et al., 2021).

PB has also been integrated into other therapeutic approaches, such as exosomes and ultrasound, for arthritis treatment. Low-density ultrasound is a noninvasive biophysical treatment that can reduce joint swelling and inflammation in OA models (Iwabuchi et al., 2014). The combined therapeutic effects of PB and low-density ultrasound on animal OA by scavenging oxygen free radicals



was investigated. It was found that this treatment could significantly remove ROS, alleviate ROS-induced apoptosis, and reduce the degeneration of articular cartilage (Zuo et al., 2021). Furthermore, neutrophil-derived exosomes engineered with ultrasmall PB nanoparticles (uPB-Exo) have been shown to be effective in treating RA. uPB-Exo selectively accumulated in activated fibroblast-like synoviocytes and acted as mimics of SOD2 and NOX2 in inflamed joints of RA *in vivo*, subsequently neutralizing proinflammatory factors, alleviating inflammatory synovitis and protecting against cartilage damage in an advanced RA mouse model (Zhang et al., 2022).

Selenium (Se) is an essential dietary nutrient and has been reported to have lower serum concentrations in RA patients than healthy individuals (Yu et al., 2016). Supplementation with Se is controversial in the treatment of arthritis is controversial due to its toxicity. Nanosized Se is known to have superior antioxidant effects and reduced toxicity (Malhotra et al., 2016). In a rat RA model, SeNPs exhibited potent anti-inflammatory effects and promoted the expression of CAT, SOD and GPX (Ren et al., 2019b).

Ultrasound, which is a noninvasive biophysical therapy and a common mode of sonodynamic therapy (SDT), can strongly penetrate inflammatory tissues and kill inflammatory cells, thus reducing synovial hyperplasia and minimizing oxidative damage to surrounding normal tissues. SDT is hampered by the hypoxic microenvironment of RA caused by fibroblast-like synoviocyte (FLS) proliferation. Rhodium NP (Rh) nanozymes with concave-cube shapes could compensate for the deficiency of ultrasound therapy by exhibiting the activities of POD and CAT, which generate O_2 and •OH to alleviate hypoxia. In addition to its

remarkable sonosensitive properties, the antibacterial drug sparfloxacin (SPX) can reside for a long time in joint tissues after systemic administration, which makes it possible to target the abnormal proliferation of FLSs in synovial tissue in the joint and block the development of RA. A small glycoprotein rich in cysteine known as SPARC is overexpressed in the synovial fluid and synovium from RA patients and increased in mice with CIA (Liu et al., 2019). SPARC has high affinity for human serum albumin (HAS) (Park et al., 2019). Therefore, HSA-modified Rh/ SPX nanozyme was fabricated for RA treatment by combining the advantages and characteristics of these components (Li et al., 2021). The preparation of Rh-SPX/HSA and its related mechanisms in the treatment of RA are shown in Figure 3.

In addition to the combination of ultrasound and nanozymes to treat arthritis, a promising technique that combines nearinfrared (NIR) with nanozymes for the treatment of OA was proposed. Epigallocatechin gallate (EGCG)-coated Au-Ag nanojars (E@Au-Ag) were produced based on the POD-like activity of Au-Ag and the scavenging of oxygen free radicals by EGCG, which is sensitive to NIR. These multifunctional enzyme-like nanomaterials can repair mitochondrial damage, promote cartilage migration, and reduce chondrocyte apoptosis (Xu et al., 2022). Biomaterials associated with antioxidant enzymes for arthritis treatment are listed in Table 2.

Discussion

OA and RA are both inflammatory diseases. RA is a systemic disease that affects joints all over the body, especially the overloaded knee joints, and affects normal movement (Radu and Bungau, 2021). OA is a local joint disease, which is common in patients with metabolic syndrome, trauma, and aging (Whittaker et al., 2021). In comparing OA and RA, a striking similarity in gene expression is found. For example, the increased levels of MMPs and the decreased expression of antioxidant enzymes occur in OA and RA, but the differences also exist. MMP-9 is the main enzyme that causes RA while MMP-13 is reported to be the most important enzyme for the development of OA. Meanwhile, in terms of pathological changes, the proliferation of synovial tissue and blood vessels in RA was more obvious than that in OA. Macrophages distributed in synovial tissue and osteoclasts from subchondral bone were the main sources of inflammation, ultimately leading to the destruction of cartilage. Therefore, chondrocytes, osteoclasts and macrophages have been the main targets for arthritis treatment with different biomaterials.

RA is a systemic inflammatory disease, and joint destruction is generally more intense than that in OA. Compared to IA injection, oral drug delivery for arthritis causes severe side effects. Recently, pain has been primarily controlled with corticosteroids and hyaluronic acid via IA injection. It is possible to deliver high drug concentrations directly to osteoarthritic joints through

Arthritis	Nanomaterial	Platform	Components	Enzyme mimics	Drug	Ref
RA	A-nanoceria-ICG	NPs	1.Albumin	SOD	metal oxides	Kratschmer et al. (1990
			2.cerium oxide	CAT		
			3.ICG	POD		
OA	F-MoS ₂	NPs	1.Fullerene	SOD	carbon-based materials	Chen et al. (2019b)
			2.MoS2	CAT		
OA	HPBzyme	NPs	Prussian blue	POD	Prussian blue	Hou et al. (2021)
				CAT		
				SOD		
OA	PBNPs	NPs	Prussian blue	POD	Prussian blue	Zuo et al. (2021)
				CAT		
				SOD		
RA	uPB-Exo	NPs	1.neutrophil-derived exosomes	SOD2	Prussian blue	Zhang et al. (2022)
			2.Prussian blue	NOX2		
RA	SeNPs	NPs	Selenium	SOD	metal	Ren et al. (2019b)
				CAT		
				GPx1		
RA	HSA-modified Rh/SPX	nanocube	1.human serum albumin (HSA)	POD	noble metal	Li et al. (2021)
			2.Sparfloxacin (SPX)	CAT		
			3.Rhodium (Rh)			
OA	E@Au-Ag	nano-jars	1.EGCG	POD	noble metal	Xu et al. (2022)
			2.Au-Ag			

TABLE 2 Biomaterials that mimic the downregulated enzymes in arthritis.

direct IA delivery. The administration of IA corticosteroids efficiently reduces articular pain and synovitis, but high concentrations of corticosteroids can also damage chondrocyte metabolism, causing changes in ECM composition and articular cartilage structure. A novel treatment for arthritis is urgently needed.

Enzymes are involved in various physiological reactions and participate in the proteolytic degradation of proteins and complex regulatory signalling pathways. Aberrant expression of these enzymes in the human body plays a critical role in pathological processes, especially inflammatory reactions. Different types of MMPs were upregulated by inflammatory factors and subsequently degrade the ECM. In addition to MMPs, ROS also participate in the development of arthritis. The generation of ROS is inhibited by endogenous antioxidants such as SOD, CAT, GPX, and heme oxygenase (HO-1). Despite the complex pathological process of arthritis, different types of arthritis including OA and RA share the common features: the increased levels of MMPs and the decreased expression of antioxidant enzymes. Hence, it is extremely feasible to design nanomaterials based on these enzymes as molecular targets for arthritis therapy.

Although nanomaterials have the advantages of high biocompatibility and bioavailability due to their structural and functional characteristics, the biosafety of nanomaterials cannot be ignored (Chen et al., 2021-09). Nanomaterials enter the body through ingestion, injection, inhalation and skin contact and subsequently accumulate in organs through blood flow, affecting the structure and function of organs (Ai et al., 2011-01). For arthritis treatment, intra-articular injection of enzyme-related biomaterials can guarantee the controlled release and targeted therapy without affecting other tissues or organs through blood circulation. Natural polymers are more suitable and safer for clinical application due to their biodegradation. Especially, hyaluronic acid from cartilage tissue has been commonly used for biomaterial. It is a promising strategy for arthritis treatment through discovering more biologically active materials from the human body in the future and combining them with drugs to regulate the expression of the enzymes mentioned above.

Given that cartilage and the synovium are affected in arthritis, various NPs that target these upregulated or downregulated enzymes mainly act on these sites, especially macrophages from the synovium and OCs from the subchondral bone. Both macrophages and osteoclasts are inflammatory cells with the same receptor on the surface of the membrane and release inflammatory factors. Therefore, biomaterials that target these inflammatory cells or chondrocytes are the current options for arthritis treatment. For the treatment of inflammatory arthritis, nano-drug delivery technologies that respond to subchondral enzymes are rare. There are technical challenges, such as how to penetrate the cartilage and reach the deep layer to target OCs that destroy the subchondral bone. Second, aside from MMPs and endogenous reductase, many enzymes are also involved in the pathological processes of arthritis. The expression of cyclooxygenase-2 (COX-2) in joints has also been linked to synovial inflammation in arthritis, and COX-2 inhibitors (celecoxib) have been frequently used and have shown therapeutic benefits in arthritis. Synergistic treatments targeting several enzymes may obtain better results. Finally, avoiding rapid clearance after IA injection is critical for maintaining drug concentrations and guaranteeing efficacy.

It should be noted that the current studies regarding enzyme-related biomaterials in the field of arthritis are not numerous; nanotherapies are extremely challenging and are also promising based on the molecular mechanism underlying arthritis.

Author contributions

Conceptualization, X-LX and D-FD; writing—original draft preparation, X-HL, J-YD; writing—review and editing, Z-HZ, -CW; visualization, Y-JS; All authors have read and agreed to publish the manuscript.

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Conflict of interest

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Sonochemical synthesis of a copper reduced graphene oxide nanocomposite using honey and evaluation of its antibacterial and cytotoxic activities

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The combination of graphene-based materials and inorganic nanoparticles for the enhancement of the nanomaterial properties is extensively explored nowadays. In the present work, we used a sonochemical method to synthesize a copper/reduced graphene oxide (Cu/RGO) nanocomposite using Australian honey and vitamin C as capping and reducing agents, respectively. The honey-mediated copper/reduced graphene oxide (H/Cu/RGO) nanocomposite was then characterized through UV-visible, XRD, HRTEM, and FTIR analysis. The copper nanoparticles (Cu-NPs) in the nanocomposite formed uniform spherical shapes with a size of 2.20 \pm 0.70 nm, which attached to the reduced graphene oxide (RGO) layers. The nanocomposite could suppress bacterial growth in both types of bacteria strains. However, in this study, the nanocomposite exhibited good bactericidal activity toward the Gram-positive bacteria than the Gramnegative bacteria. It also showed a cytotoxic effect on the cancer colorectal cell line HCT11, even in low concentrations. These results suggested that the H/Cu/ RGO nanocomposite can be a suitable component for biomedical applications.

KEYWORDS

sonochemical method, copper/reduced graphene oxide nanocomposite, honey, antibacterial, cytotoxicity assay

Introduction

In the past 50 years, pathogenic bacteria have caused a plethora of diseases in the human population. Some of these major emerging bacteria include *Staphylococcus aureus* (*S. aureus*), *Escherichia coli* (*E. coli*), *Clostridium difficile*, *Campylobacter spp.*, and *Helicobacter pylori* (Vouga and Greub, 2016). Although some commensal bacteria such as *Lactobacillus* and *Bifidobacterium* may exist as part of the human microbiota,

which might play beneficial roles in maintaining homeostasis (Wang et al., 2017), many of these bacteria could also cause various illnesses such as liver diseases, infection, respiratory diseases, gastrointestinal malignancy, and metabolic disorders (Wang et al., 2017). The emergence of bacterial antimicrobial resistance (AMR) has even aggravated this issue and poses a major threat to global health. According to a recent systematic analysis, the six leading pathogens responsible for death associated with AMR are *E. coli, S. aureus, Klebsiella pneumoniae, Pseudomonas aeruginosa (P. aeruginosa), Streptococcus pneumoniae*, and Acinetobacter baumannii which are the combination of both Gram-positive and Gram-negative strains (Murray et al., 2022).

The second main cause of death in the United States is cancer, and among different cancer types, lung cancer is the leading cause of cancer mortality (Siegel et al., 2022). For both sexes, the highest incidence of cancer is led by breast cancer, followed by prostate and lung/bronchus cancers (Siegel et al., 2022). Similarly, chemo- and/or radio-resistance presented by the patient's tumor remains the main barrier to effectively eradicating tumor from the body. On top of this, off-target side effects suffered by the patients due to cancer therapy are another obstacle (Miller et al., 2022). Hence, it is vital to look for a novel anticancer drug or therapeutic strategy to treat cancer more effectively.

Recently, nanomaterial has been widely explored for their special properties to mitigate these problems. There are various factors that may affect the biological activities of the nanomaterial, such as the shape, size, electronic structure, surface properties, and some additional factors related to the interaction conditions between the materials and the target cells (Sengupta et al., 2019). Copper nanoparticles (Cu-NPs) have been widely assessed for their properties. Aside from its low-cost production, copper also exhibits good thermal and electrical conductivity, and biological and antimicrobial activities (Zhou et al., 2019; Noman et al., 2020; Merugu et al., 2021). Recently, the United States Environmental Protection Agency recognized copper as the first solid antimicrobial material (Ouyang et al., 2013; Arendsen et al., 2019). However, researchers found that pure metallic Cu-NPs are difficult to obtain as copper tends to oxidize easily when exposed to the air, and it will also tend to agglomerate without proper protection (Rostami-Tapeh-Esmaeil et al., 2021). Hence, the usage of green material as a capping agent or stabilizer has gained researchers' attention since it is reported to produce monodispersed pure Cu-NPs by a fast and green method (Nagar and Devra, 2018).

Graphene is a unique structure that attracts great attention due to its interesting physical and chemical characteristics (Luo et al., 2020), including large surface area, good conductivity, and high thermal properties (Ouyang et al., 2013). Graphene and its derivatives are used in extensive applications such as electronic devices (Moozarm Nia et al., 2017), energy storage (Rawal et al., 2020), and biomedical applications (Kumar et al., 2017). The presence of oxygen functional groups such as epoxide, carboxyl, and hydroxyl in the structure of the graphene oxide (GO) and reduced graphene oxide (RGO) makes them suitable for the production of nanocomposites (Gan et al., 2019a). These groups act as bioactive molecules that could functionalize the graphene sheet with other materials such as metal and metal oxide nanoparticles (Gan et al., 2019b; Jang et al., 2020).

Graphene oxide can be decorated with some materials through chemical reduction (in situ), hydrothermal, and electrochemical processes, and through the attachment of the premade nanoparticles to the graphene surface (ex situ) to form graphene-based nanomaterials (Sarkar and Dolui, 2015; Yin et al., 2015; Iranshahi and Iranshahi, 2022; Thy et al., 2022). Luo et al. (2020) used refluxed process in reduced graphene oxide/copper synthesizing the nanocomposites (RGO/Cu-NCs) in an oil bath at 100°C for 24 h with hydrazine hydrate. This method involved a long processing time and hazardous material. In fact, nowadays, researchers are interested in using simple and green materials to synthesize the Cu/RGO nanocomposite since it is a much eco-friendlier method. Rios et al. (2019) used an in situ reduction method to produce reduced graphene oxide/ copper nanoparticles (RGO/Cu-NPs) in the presence of ascorbic acid for 12 h at 80°C. Fahiminia et al. (2019) synthesized Cu/RGO nanocomposites using plant extract (Euphorbia cheiradenia Boiss) and applied it as a catalyst for dye removal. Yang et al. (2019) produced cuprous oxide/ reduced graphene oxide (Cu2O-RGO) nanocomposites through chemical reduction by using polyethylene glycol (PEG) and ascorbic acid with the addition of sodium hydroxide (NaOH), and used them for the antibacterial study.

Indeed, Tu et al. (2021) reported that the Cu/RGO nanocomposite exhibited better biological activity compared to the reduced graphene oxide (RGO) alone. Generally, the combination of RGO and copper ions happened by the cation- π interaction between copper ions and π -electrons that coming from the aromatic rings of RGO (Xu et al., 2019; Yan et al., 2019; Ismail et al., 2021; Tu et al., 2021). This functionalized RGO could enhance the antibacterial activity where both participated in killing the bacteria cells through electrostatic interaction between the positive charge of copper ions from the nanocomposite and the negatively charged membranes of bacteria (Sanchez-Lopez et al., 2020). RGO could also kill bacteria through the sharp edge of its structure (Prasad et al., 2017). The aggregation of the RGO due to the π - π stacking would have a hydrophobic structure which is known to give strong bacteria absorption that could help in better releasing copper ions and attacking the bacteria cell efficiently (Szunerits and Boukherroub, 2016). This will show excellent results in antibacterial activity compared to the copper ion and RGO alone. Up until now, few studies for anticancer using Cu/RGO nanocomposite were reported. Kodous et al. (2022) found that Cu/RGO nanocomposites produced by using the ultrasonication method could inhibit human breast cancer cells (MCF-7 cancer cells).

Honey is considered a green material since it is a non-toxic substance that possesses rich sugar source carbohydrate components (Balasooriya et al., 2017). It is also a simple material that does not have to undergo any extraction process, unlike plants and microorganisms. Most importantly, it was also reported for its biological activity properties and its potential as a capping agent (Ismail et al., 2019). Eucalyptus flower species is usually one of the main sources of nectar for the Australian honeybees (Apis mellifera) to produce honey. The source of nectar, the combination of proteins secreted by the bee for the honey-ripening process, and protein from plant pollen will affect the honey's chemical composition. According to Beiranvand et al. (2021), the major component in pure Australian honey was carbohydrates, which could act as a capping and reducing agent. However, the chemical component such as carbohydrate in honey is considered a weak reducing agent so it needs another booster to enhance the reducing process of the nanoparticles, and for this, vitamin C (ascorbic acid) was chosen in this study since it is also a green material. Hence, in this work, we produced a honey-mediated copper reduced graphene oxide (H/Cu/RGO) nanocomposite using a sonochemical method, where Australian honey and ascorbic acid were served as capping and reducing agents, respectively, during the process. The sample was analyzed by using UVvisible, XRD, HRTEM, and FTIR, and it was then tested for antibacterial and cytotoxicity properties.

Materials and methods

Materials

The source of honey was from the Capilano Honey Limited (Australia). Standard graphene oxide (water dispersion, 4 mg/ ml) was purchased from Graphene (U-Malaya). Copper II nitrate trihydrate (Cu(NO₃)₂.3H₂O, AR grade), ascorbic acid (C₆H₈O₆, AR grade), and sodium hydroxide (NaOH) were purchased from R&M Chemical, United Kingdom. All the chemicals were of analytical grade without further purification. Two Gram-positive bacteria, methicillin-resistant Staphylococcus aureus (MRSA, clinical isolate) and Enterococcus faecalis (E. faecalis, ATCC 33186), and two Gram-negative bacteria, Escherichia coli (E. coli, ATCC 11775) and Pseudomonas aeruginosa (P. aeruginosa, ATCC 10145), were used for antibacterial assessment. They were cultured and maintained in sterile Mueller-Hinton agar and broth media (Becton Dickinson, United States). The colorectal cancer cell line HCT116 (ATCC CCL-247) and human normal colon cell CCD112 (ATCC CRL-1541) were used for the cytotoxicity assay.

Synthesis of reduced graphene oxide (RGO) and honey-mediated copper/ reduced graphene oxide (H/Cu/RGO) nanocomposite

RGO was produced through the reduction of graphene oxide. For this, 2 ml of GO was added to 50 ml of deionized water. After vigorous stirring for around 30-40 min, 7.5 ml ascorbic acid (1 M) was slowly added to the solution using a dropper, while the sample was treated with ultrasonic irradiation for 10 min with a fixed setting parameter (amplitude 80%, pulse on 1s and pulse off 1 s). The mixture was then cooled down to room temperature (RT), centrifuged, washed with distilled water, and dried in the oven. The H/Cu/RGO nanocomposite was synthesized following the method by Zhang et al. (2016) with some modifications. In brief, Australian honey (15 w/v %) was dissolved in 50 ml of 0.025 M Cu(NO₃)₂.3H₂O, and the pH was adjusted between pH 7 to 8. Afterward, 1 ml of the RGO (1.5 mg/ ml) was mixed with the combination of honey and copper salt solution under continuous stirring at RT for 30-40 min. The mixture solution was then treated with ultrasonic irradiation for 10 min by adding 7.5 ml of 1 M ascorbic acid dropwise simultaneously. The compound was finally cooled down to RT, centrifuged, washed with distilled water, and dried in the oven.

Characterizations of the H/Cu/RGO nanocomposite

The synthesis H/Cu/RGO nanocomposite was determined using ultraviolet-visible (UV-vis) spectroscopy (UV-2600, SHIMADZU) in the range of 220–800 nm. X-ray diffraction (XRD, Philips, X'pert, Cu Ka) was used to analyze the structure of the H/Cu/RGO nanocomposite in the range of 5° -80° (20). The size and the shape of the nanocomposite were evaluated by using high-resolution transmission electron microscopy (HRTEM, JEM-2100F). Fourier transform infrared (FTIR) spectra were obtained using an attenuated total reflectance (ATR) IRTracer-100 spectrophotometer (Shimadzu, Malaysia). The spectra were set within a range of 400–4,000 cm⁻¹.

Antibacterial activity

To determine the minimum inhibitory concentration (MIC) values, the broth micro-dilution method was used for the H/Cu/ RGO nanocomposite against Gram-positive (MRSA and *E. faecalis*) and Gram-negative (*E. coli* and *P. aeruginosa*) bacteria using the Clinical and Laboratory Standards Institute (CLSI) protocols. For this, a single colony of fresh bacterial culture (12–18 h) was isolated from the Mueller–Hinton agar



(MHA) plates and inoculated into the Mueller-Hinton broth (MHB). The culture was grown overnight (16–18 h) prior to the experiments. The next day, the bacterial concentration was standardized to an optical density (OD) of 600 nm (approximately 1×10^8 CFU/ml) with MHB. Two-fold serial dilutions of the H/Cu/RGO nanocomposite were prepared in 96well plates to get the final test concentrations of 0, 7.8, 15.6, 31.3, 62.5, 125, 250, 500, and 1,000 $\mu g/ml$ per well. Thereafter, 10 μl of bacterial suspension equivalent to 106 CFU/ml of exponentially growing bacterial cells were added to the wells followed by 18 h of incubation at $35 \pm 2^{\circ}$ C. The plate was then read for absorbance at 600 nm using a microplate reader (GloMax Discover Instrument, Promega). The percentage of cell viability was calculated using Equation (1), and the minimum inhibitory concentration which inhibits 50% bacterial growth (MIC_{50}) value was then determined.

%Viability = OD of sample well (mean)/OD of control well (mean) \times 100. (1)

Cytotoxic effect of the H/Cu/RGO nanocomposite

Cell proliferation assay (Promega) was used to determine the cytotoxic properties of the H/Cu/RGO nanocomposite. Briefly, 5×10^3 human colorectal cancer cell line HCT116 and human normal colon cell CCD112 were seeded in a 96-well plate (100 µL/well) and incubated at 37°C overnight in a 5% CO₂ humidified

incubator. The next day, 2-fold serially diluted nanocomposites (500, 250, 125, 62.5, 31.3, 15.6, 7.8, and 0 µg/ml) were added into the wells (100 µl/well). After 72 h incubation at 37°C in a 5% CO₂ humidified incubator, the wells were treated with 20 µl MTS (3-(4,5-dimethylthiazol-2-yl)-5-(3-carboxymethoxyphenyl)-2-(4-sulfophenyl)-2H-tetrazolium) reagent followed by an additional 3 h incubation at 37°C in the 5% CO₂ incubator. Optical density (OD) was then measured at 490 nm using a multimode microplate reader (Tecan). The dose–response graph was plotted by calculating the percent of cell viability using Eq. 1, and half maximal inhibitory concentration (IC₅₀) was then calculated.

Results and discussion

Synthesis of the H/Cu/RGO nanocomposite

As illustrated in Scheme 1, we used copper nitrate solution as a precursor for Cu-NPs synthesis. To accelerate the process, NaOH was added to form an intermediate which is copper hydroxide $Cu(OH)_2$. The pH of the solution was controlled between pH 7 to 8 since it is the preferred environment to produce smaller sizes of pure Cu-NPs. According to Amjad et al., when the pH increased (between pH 6 and pH 10), the size of nanoparticles decreased (Rajesh et al., 2016; Amjad et al., 2021). Since the aim of this study is to produce pure metallic Cu-NPs, the pH needs to be in a basic medium. The reduction process of



the compound was furthered with the addition of the ascorbic acid as a reducing agent and assistance of ultrasonic irradiation to enhance the reaction process. Honey acts as a capping agent to control the size and shape of the nanoparticles in the solution. The nanocomposite was then tested against the bacteria and cancer cell line to observe its biological activities.

Meanwhile, Eqs 2–5 described the possible chemical formation of H/Cu/RGO nanocomposite.

$$Cu^{2+}(aq) + Honey(1) \xrightarrow[T=25\circ C]{Stirring} [Cu(Honey)]^{2+}(aq), \quad (2)$$
$$[Cu(Honey)]^{2+}(aq)$$

$$+ 2OH^{-}(aq) \xrightarrow{\text{Stirring}}_{T=25 \circ C} [Cu(OH)_{2}/\text{Honey}](s), \qquad (3)$$

$$\left[Cu(OH)_2 / Honey \right] (aq) + GO(aq) \xrightarrow{\text{Stirring}}_{T=25 \circ C} \left[Cu(OH)_2 / Honey \right]$$

$$-GO(aq),$$
(4)

$$[Cu(OH)_2/Honey] - GO(aq)$$

$$\begin{array}{c} + 2H_2O(1) & \xrightarrow{C_6H_8O_6} \\ & & \\ &$$

The formation of the reaction process of the $Cu(OH)_2/$ honey complex (Eq. 2) resulted in a blue-colored solution as previously reported study (Ismail et al., 2019). Cu(OH)₂ was obtained by adding the sodium hydroxide (NaOH), and it acts as nuclei during the process. A mutarotation process occurs during this phase since OH⁻ in the solution could change the α -glucose into β -glucose by opening the chain structure and forming the aldehyde group (-CHO) (Upadhyay and Kumar, 2017; Alejandro et al., 2017). This aldehyde group with the presence of the energy from the ultrasonic irradiation process was then oxidized by the complex copper ions to form the gluconic acid. This initiates the nucleation and growth of the Cu-NPs in the solution. In addition, the Cu²⁺ ion growth could also occur at the nucleation site of the GO substrate as the reduction of Cu2+ takes place through galvanic displacement and redox reaction (Eq. 4). The GO and copper ions were further reduced to H/Cu/RGO nanocomposites with the presence of ascorbic acid and the assistance of the ultrasonic irradiation process. It could be concluded that GO acts as a substrate and could also help as a reducing agent for the copper ions (Zhang et al., 2016). The GO sheets might bind with the copper ion by electrostatic interaction of the copper ion and through the cation-pi (cation-) interaction of the benzene ring with the cation (Cu²⁺) (Alayande et al., 2020).

Characterization of GO, RGO, and H/Cu/ RGO nanocomposite

Figure 1 shows the UV-vis spectra of the honey, GO, RGO, and H/Cu/RGO nanocomposite samples. The absorption peak of the honey appeared around 277 nm due to the origin and age of the honey itself (Posudin, 2016; Zhang et al., 2016). The $\pi - \pi^*$



transition of the C=C bond and $n-\pi^*$ transition of the C=O bond could be seen in Figure 1B for GO at 238 and 305 nm, respectively. The shifting peak at 260 nm to a higher wavelength and the peak around 305 nm disappeared for the RGO (Figure 1C) indicating the reduction of GO to RGO by the ascorbic acid. This phenomenon occurred because of the restoration of the aromatic system conjugation and the decrease of the carboxyl groups in the RGO layer (Navya Rani et al., 2019; Kang et al., 2020). The higher the conjugation degree, the lower the energy required in order to produce the electronic transition, and hence, the peak is shifted to a higher value associated with less energy involved (Rios et al., 2019). Figure 1D depicts the surface plasmon resonance of metallic phase copper nanoparticles (Cu-NPs) exhibited at 569 nm which proved the reduction of Cu^{2+} to Cu^0 occurs during the synthesis process (Fahiminia et al., 2019).

XRD analysis was conducted for the honey, GO, RGO, and H/Cu/RGO nanocomposite. Figure 2A shows that the XRD diffraction pattern of honey at $2\theta = 17.64^{\circ}$ with a broad peak. The shifting of the diffraction peak in Figures 2B,C for GO and RGO from 9.43° to 24.87° indicated that the reduction of GO to

RGO occurred under sonication treatment with the presence of the ascorbic acid. The interspacing distance between the layer of the GO and RGO was calculated by using Bragg's law equation as in Eq. 6

$$n\lambda = 2d\sin\theta,$$
 (6)

where n = 1, λ is the wavelength of the X-ray beam (0.154 nm), d is the distance between adjacent GO or RGO sheets, and θ is Bragg's angle. The values of interspacing distance were 0.937 and 0.357 nm, respectively, for GO and RGO. The decrement of the value shows that the formation of RGO occurs according to the previous study (Rana et al., 2018). The changes in the interspacing distance reveal the exfoliation of the RGO layer happened after the reduction process and the decrease of the oxygenated functional groups on the surface (Kumar et al., 2019). For the H/Cu/RGO nanocomposite (Figure 2D), three diffraction peaks at $2\theta = 43.4^{\circ}$, 50.5°, and 74.4° could be assigned to the (111), (200), and (220) crystal planes corresponding to the cubic structure of Cu which signified the formation of metallic copper on the RGO. These diffraction peaks of Cu were matched with the standard reference of the metallic Cu for



the cubic structure which is JCPDS 04-0836. The peak for RGO could not be seen in the XRD pattern of the H/Cu/RGO nanocomposite, which could be related to the aggregation and restacking layer of the RGO with the insertion of the Cu-NPs in the nanocomposite (Guo et al., 2016; Kumar et al., 2019; Chen et al., 2020).

In the HRTEM images (Figure 3), GO showed a fine-layer structure like a sheet, while RGO revealed a wrinkled structure. This phenomenon is related to the reduction of the GO to RGO, where the GO layer was exfoliated and tended to be decreased in size as it was treated with the ultrasonic. The thermal treatment through the ultrasound irradiation process can lead to the wrinkling of the RGO due to the reduction of the amount of oxygen-containing functional groups during sheet exfoliation (Rana et al., 2018; Khan et al., 2020). The Cu-NPs in the nanocomposite formed uniform spherical particles with a size of 2.20 \pm 0.70 nm on the RGO layers. The exfoliation of the RGO layers into smaller scale with a fine particle of Cu-NPs that either are decorated between or on the surface of RGO layers happened. It concludes that Cu-NPs can bind to the graphene-based materials to form Cu/RGO nanocomposite (Zhu et al., 2017; Menazea and Ahmed, 2020).

Through the FTIR spectra (Figure 4), the honey peak (Figure 4A) illustrated a strong and broad peak at $3,291 \text{ cm}^{-1}$ related to a hydroxyl group (-OH) stretching vibration which also

might overlap with the -NH stretching vibration of primary amine protein (Boldeiu et al., 2019). Two weak peaks appeared at 2,924 cm⁻¹ and 2,883 cm⁻¹, matched to the C-H stretching bands of the aldehyde group of glucose. Carbonyl group (-C=O) stretching vibration of protein could be seen at 1,636 cm⁻¹, and the peak at 1,427 $\rm cm^{-1}$ and 1,334 $\rm cm^{-1}$ were related to C-H bending and C-O bending of glucose. While, at 1,017 cm⁻¹, the peak correlated to C-O-C stretching, C-O stretching, and C-N stretching amine of glucose, fructose, and protein in honey. The GO spectrum (Figure 4B) demonstrated O-H stretching vibration with a broad peak between 3,200 cm⁻¹ to 3,600 cm⁻¹. The peaks at 1734, 1,618, 1,394, 1,161, and 1,033 cm⁻¹ were correlated to the C=O stretching vibration of carbonyl groups presented in the GO sheet, C=C skeletal vibration, the sp3 C-H stretching vibration of saturated carbon, the epoxy C-O stretching vibration, and the alkoxyl C-O stretching vibration, respectively (Nguyen et al., 2019; Sengupta et al., 2019). However, in RGO, the disappearance of the carboxyl group at 1,734 cm⁻¹ and sp³ C-H stretching vibration of saturated carbon at 1,394 cm⁻¹ demonstrated the reduction of GO to RGO during the synthesis process. For the Cu/RGO nanocomposite (Figure 4D), the C=C vibration of the graphene skeleton peak could be observed at 1,539 cm⁻¹, indicating that the GO was reduced to form the Cu/RGO nanocomposite (Navya Rani et al., 2019). In addition, the





TABLE 1 MIC ₅₀	values	of H	/Cu/RGO	nanocomposite	against	four
bacteria strains						

Sample	MIC ₅₀ of sample (µg/ml)					
	Bacterial strains					
H/Cu/RGO nanocomposite	MRSA	E. faecalis	P. aeruginosa	E. coli		
	67.96	6.12	>1,000	134.16		



weak peak band area around 900 cm⁻¹ in Figure 4D could be due to the shifting of the C-O/C-N stretching of protein and carbohydrate of honey biomolecules presented in the H/Cu/RGO nanocomposite (Ismail et al., 2019).

Antibacterial activity of the H/Cu/RGO nanocomposite

Antibacterial activity of the H/Cu/RGO nanocomposite toward bacterial strains was tested using the minimum inhibitory concentration (MIC) assay, where the selected Gram-positive (MRSA and *E. faecalis*) and Gram-negative (*P. aeruginosa* and *E. coli*) bacteria were found to be affected by the nanocomposite. Figure 5 illustrates that the inhibition activity of the nanocomposite was better toward Grampositive bacteria compared to the Gram-negative strains. As shown in Table 1, the lowest MIC₅₀ value was detected toward *E. faecalis*, where the nanocomposite could inhibit the growth of less than a quarter of the bacteria at a low concentration of 6.12 µg/ml, while for MRSA, the MIC₅₀ was detected at the concentration of 67.96 µg/ml. For Gramnegative strains, the MIC₅₀ was at 134.16 µg/ml and greater than 1,000 µg/ml for *E. coli* and *P. aeruginosa*, respectively.

The value of Cu-NPs as antibacterial agents has been studied for a long time (Lv et al., 2020). However, the H/Cu/RGO nanocomposite gave better antibacterial activity performance compared to the Cu-NPs only. In a prior study where honeymediated Cu-NPs was tested on E. faecalis and E. coli, the values of MIC₅₀ were 15.6 µg/ml and 250 µg/ml, respectively, which were higher than the value gained from the nanocomposite in current work. This is most possibly due to the toxic effects of Cu-NPs and the RGO that influenced the bacteria cells. It might be due to the Cu-NPs that affect bacteria by the generation of reactive oxygen species, lipid peroxidation, protein oxidation, and DNA degradation through liberating nascent Cu ions from the Cu-NP surface (Chatterjee et al., 2014). The cell membrane stress due to the graphene sheet layer structure itself is also possibly among the factors for the bactericidal activity to change (Prasad et al., 2017).

A possible explanation for the variation in antibacterial activity against different bacterial strains can be related to the difference in the bacteria cell envelope (Sriramulu et al., 2020). In contrast to Gram-positive strains which consist of the layers of peptidoglycan, Gram-negative bacteria, besides the inter thin peptidoglycan cell wall, are an outer membrane surrounded by containing lipopolysaccharide (LPS), which can act as an additional protection shield for the cell. It could be one of the possible reasons why P. aeruginosa could tolerate the H/ Cu/RGO nanocomposite, even at high concentrations.

Cytotoxic effect of the H/Cu/RGO nanocomposite

Cytotoxic activity of the nanocomposite was tested in both normal and cancer colorectal cell lines (Figure 6), where the H/ Cu/RGO nanocomposite showed a higher cytotoxic effect compared to RGO, even at low concentrations. This could be due to the combination of Cu-NPs and RGO that enhanced the properties of the cytotoxic activity. This enhancement might also be attributed to the size of the Cu-NPs attached to the RGO, which is smaller in size with a spherical shape that makes them easier to interact with the cells and kill them. Similar to the

Samples	IC ₅₀ of sample (µg/ml)				
	CCD 112 (normal cell)	HCT 116 (cancer cell)			
RGO	33.70	141.50			
H/Cu/RGO nanocomposite	2.14	7.7			

TABLE 2 IC₅₀ of the RGO and nanocomposite toward normal and cancer cell lines of the colorectal cell.

antibacterial study, in comparison with our previous work, the H/Cu/RGO nanocomposite showed higher anticancer action (IC₅₀–7.7 µg/mL as shown in Table 2) than honey-mediated Cu-NPs without RGO (IC₅₀–46.11 µg/ml) in HCT116 cells (Ismail et al., 2019). A previous study reported that Cu-NPs killed SW480 human colon cancer cells at an IC₅₀ value of 68 µg/ml by inducing reactive oxygen species (ROS)-mediated apoptosis (Ghasemi et al., 2022). As this is the first study reporting potential anticancer action of the H/Cu/RGO nanocomposite, their exact mechanisms of cancer cell killing are not known and warrant further investigations.

In both graphs (Figures 6A,B), at a concentration of 31.2 µg/ ml, the nanocomposite killed both cells at 100%, which indicates that this compound is not selective toward cancer cells. This could be seen in Table 2, where both RGO and H/Cu/RGO nanocomposite are non-selective toward cancer cells. Thus, further modifications are needed to enhance the selectivity of the nanocomposite toward cancer cells. For example, the nanocomposite can be conjugated to target-specific aptamers, peptides, antibodies, or other ligands to allow the nanocomposite to specifically bind to the surface molecules of cancer cells and enhance the drug localization, retention effect, and cellular uptake (Sutradhar and Amin, 2014; Martinelli et al., 2019).

Conclusion

In conclusion, the green synthesized Cu/RGO nanocomposite using honey and ascorbic acid as capping and reducing agents, respectively, resulted in small-sized and spherical-shaped Cu-NPs attached to the RGO sheet. Our data proved that the agglomeration of copper could be prevented by combining the Cu-NPs with graphene-based materials in the presence of eco-friendly capping and reducing agents. The nanocomposite revealed good antibacterial and cytotoxicity activities, making them suitable for biomedical applications. However, the nanocomposite needs to be further studied to improve its specificity toward cancerous cells.

Data availability statement

The original contributions presented in the study are included in the article/Supplementary Material; further inquiries can be directed to the corresponding authors.

Author contributions

NI: writing-original draft, conceptualization, methodology, investigation, formal analysis, data curation, validation, resources, and data curation. KS: funding acquisition, methodology, conceptualization, investigation, resources, formal analysis, data curation, supervision, and writing-review and editing. SM: methodology, formal analysis, and validation. hirofumi hara: resources. S-YT: resources, funding acquisition, methodology, formal analysis, investigation, data curation, and writing-review and editing. HM: review and editing, supervision, and financial support of research publication.

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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A metal-organic framework nanocomposite with oxidation and near-infrared light cascade response for bacterial photothermal inactivation

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Photothermal treatment is an effective and precise bacterial disinfection method that can reduce the occurrence of bacterial drug resistance. However, most conventional photothermal treatment strategies have the problem that the photothermal response range does not match the infection area. Herein, a metal-organic framework (MOF) nanocomposite responding to the oxidation state of the bacterial infection microenvironment was constructed for near-infrared (NIR) photothermal bacterial inactivation. In this strategy, the MOF was used as a nanocarrier to load tetramethylbenzidine (TMB) and horseradish peroxidase (HPR). The high oxidation state of the bacterial infection microenvironment can trigger the enzyme-catalyzed reaction of the nanocomposite, thereby generating oxidation products with the NIR photothermal effect for bacterial disinfection. The synthesis and characterization of the nanocomposite, oxidation state (H_2O_2) response effect, photothermal properties, and antibacterial activities were systematically studied. This study provides a new idea for building a precision treatment system for bacterial infection.

KEYWORDS

metal-organic framework, cascade response, antibacterial materials, photothermal therapy, nanodrug

Introduction

Bacterial infection seriously threatens human life and health. As a traditional medicine for treating bacterial infections, antibiotics have saved countless lives. However, the use of antibiotics will lead to the emergence of bacterial resistance, which greatly reduces the therapeutic effect of antibiotics and even makes antibiotics ineffective (Laxminarayan et al., 2013; Mamun et al., 2021). The abuse of antibiotics in recent years has accelerated the emergence of bacterial drug resistance. Unfortunately, the speed at which we develop new antibiotics is far lower than the speed at which bacterial drug resistance develops (Hutchings et al., 2019). According to the World Health Organization (WHO), around 7 00,000 people die of drug-resistant bacterial infections every year worldwide. If effective measures are not taken, it is estimated that 10 million people will die of drug-resistant bacterial infections every year by 2050. In the face of such a severe situation, on one hand, it is necessary to accelerate the development of antibiotics and meanwhile avoid the abuse of antibiotics; on the other hand, it is necessary to develop new antibacterial strategies.

Nanoparticle-mediated physical stimulation therapy is a promising bacterial therapy strategy that can partially replace antibiotics (Ji et al., 2022; Wang et al., 2022). In such a strategy, special nanoparticles are utilized as antennas to convert physical stimulation (e.g., light, magnetic, X-ray, and ultrasound) into heat energy or free radicals for bacterial inactivation (Jia and Zhao, 2021; Zhang et al., 2022b; Deng et al., 2022; Ji et al., 2022). For example, most of the noble metal nanoparticles, nano-carbon materials, magnetic nanomaterials, some nanopolymers, etc. can be heated under light, magnetic, ultrasonic, or other physical stimulations to generate high temperature for bacterial inactivation; photosensitizers and nanosemiconductor materials (such as titanium dioxide, bismuth vanadate, and quantum dots) can generate free radicals under light, X-ray, or even ultrasound irradiation for bacterial disinfection (Karami et al., 2021; Du et al., 2022; Fan et al., 2022). Among these strategies, the photothermal strategy has obvious advantages such as easy access to light sources, high bacterial inactivation efficiency, and low toxic side effects. In addition, photothermal treatment is not easy to induce bacterial resistance (Zhang et al., 2020; Mu et al., 2022). Therefore, in recent years, photothermal antibacterial therapy has attracted wide attention, and many progresses have been achieved in this field (Han et al., 2020). Achieving high antibacterial efficiency is no longer a major problem of photothermal methods. Improving the accuracy of treatment is currently a development trend in this field. Although modifying targeted molecules such as antibodies and antimicrobial peptides can improve the accuracy of nanophotothermal therapy to a certain extent, the modification also brings

high cost problems. Using the special microenvironment of the lesion site to construct a responsive photothermal treatment strategy is an effective means to improve the accuracy of treatment. Such a strategy is expected to be a good solution to improve the accuracy of nanoparticlemediated photothermal therapy.

In this study, a metal-organic framework (MOF) nanocomposite responding to the oxidation state of the bacterial infection microenvironment was constructed for near-infrared (NIR) photothermal bacterial inactivation. In this strategy, MOF (UiO-66) was used as a nanocarrier to load tetramethylbenzidine (TMB) and horseradish peroxidase (HPR) (Scheme 1). The high oxidation state of the bacterial infection microenvironment can trigger the enzyme-catalyzed reaction of the nanocomposite, thereby generating oxidation products with the NIR photothermal effect for bacterial disinfection.

Materials and methods

General information

1,4-Dicarboxybenzene, 70% zirconium propoxide [Zr(OnPr) 4] solution in 1-propanol, and tetramethylbenzidine (TMB) were purchased from Shanghai Aladdin Bio-Chem Technology Co., Ltd. Horseradish peroxidase (HPR) was purchased from Sigma-Aldrich. The transmission electron microscopy (TEM) image was captured with a 120-KV JEM-1400 microscope with a Gatan Rio16 digital camera. The sample for TEM was prepared by dropping the dilute UiO-66 solution onto carbon-coated copper grids. Powder X-ray diffraction (XRD) patterns were recorded with a Bruker D8 diffractometer (Bruker, Germany). UV-vis adsorption spectra were detected using a UV-1900 spectrometer (SHIMADZU, Japan). Fluorescence images were captured using a NIB900-FL fluorescent microscope with a Nexcan-T6CCD digital camera (Nexcope, China). A homemade 900-nm NIR light source was used for NIR light irradiation, and the power density was measured with a power density meter. A colony counter icount 11 (Xun Shu, China) was used to count colonyforming units.

Synthesis of the metal-organic framework and the nanocomposite

MOF UiO-66 was synthesized according to the previous literature with some slight modifications (DeStefano et al., 2017). A measure of 3.5 ml of DMF, 2 ml of acetic acid (2.1 g, 35 mmol), and 30.5 μ l of a 70% zirconium propoxide [Zr(OnPr)4] solution (in 1-propanol) (26 mg, 0.079 mmol) were mixed in a 10-ml scintillation vial. The solution was heated in an oil bath at 130°C for 2 h and then allowed to cool to room temperature. The color of

the mixture changed from colorless to yellow during heating. To the solution, 37.5 mg of 1,4-dicarboxybenzene was added, and after sonication for 30 s, the solution was stirred at room temperature for 18 h. Then, the MOF was separated by centrifugation and washed several times with DMF and water and finally dispersed in water for further use. The nanocomposite (UiO-66@TMB-HRP, UTH) was synthesized by simple incubation of UiO-66 with TMB and HRP. Briefly, to a 5 ml solution of 5 mg/ml UiO-66, the TMB stock solution was added with a final concentration of 0.5 mM; after incubation for 3 h, 25 U HRP was added and stirred at 4°C for another 8 h. After then, the nanocomposite was separated by centrifugation, washed several times with water, and finally dispersed in water for further use.

Photothermal measurement

The photothermal effect of UTH in different conditions under 900 nm light (0.5 w/cm²) irradiation was measured by using a thermal imaging camera. The heating and cooling temperature changes were recorded, and the photothermal conversion efficiency (η) was calculated according to the following equations:

$$\eta = \frac{hs (T_{max} - T_{suur}) - Q_0}{I (1 - 10^{-A_{900}})},$$
(1)

$$\tau_s = \frac{m_d c_d}{hs},\tag{2}$$

$$Q_{s0} = hs \left(T_{max.water} - T_{suur} \right), \tag{3}$$

where τ_s was observed by linearly fitting the plot of the cooling time versus $-\text{Ln}\theta$. m_d is the mass of the UTH solution, and C_d is the heat capacity of water (4.2 J g⁻¹ K⁻¹). T_{max} is the equilibrium temperature; T_{surr} is the surrounding ambient temperature; $T_{\text{max}, \text{ water}}$ is the maximum temperature of the heated water.

Antibacterial test

Escherichia coli (E. coli) and Staphylococcus aureus (S. aureus) were selected as Gram-positive and -negative model strains, respectively. Monocolonies of the bacteria on a solid agar plate were transferred to 2 ml LB medium and shaken under 150 rpm at 37°C for 12 h. In the photothermal antibacterial experiments, the bacterial solution (with an optical density at 600 nm of 0.5) was mixed with UTH and 1 mM H_2O_2 was added; after incubation for 5 min, the mixture was irradiated under 900-nm light for another 5 min. After then, the treated bacterial solution was diluted and transferred to the solid agar plate. After being placed in an incubator at 37°C for 12 h, the plates were photographed and colonies were counted.

Therapeutic effect against mouse skin wound infection

Kunming mice were used for skin wound infection model fabrication, which has been approved by the Ethics Committee of Animal Experiments in Zhejiang Sci-Tech University, and all procedures followed the guidelines for animal experiments in Zhejiang Sci-Tech University. The hair of the mouse's quilt was removed with depilation cream. A small piece of the back skin was cut off to construct a wound model. A measure of 10 μ l of the *S. aureus* solution with the OD600 of 1 was dropped to the wound for the infection. Then, 10 μ l of UTH was dropped to the infected wound once a day for the first 3 days, followed by irradiation with 900-nm light for 5 min. Photographs of the wound were taken every day to record the changes.

Results and discussion

Synthesis and characteristics

Considering the porous characteristics and high stability, zrmof (UiO-66) was chosen as the nanocarrier for TMB and HRP loading. The prepared UiO-66 is milky white and has good stability and dispersion in water. TEM characterization results show that UiO-66 has a size of about 100 nm and good dispersion (Figure 1A). The crystalline and phase information were investigated by powder X-ray diffraction (XRD), and the patterns are shown in Figure 1B. The appearance of sharp peaks in the XRD patterns indicates that UiO-66 has good crystallinity. The size and zeta potential of UiO-66 and UiO-66 loaded with TMB and HRP (UiO-66@TMB-HRP, UTH) were measured with a DLS machine. The main size was around 110 nm; after TMB and HRP loading, the obtained nanocomposite UTH showed a larger size of around 210 nm (Figure 1C). UiO-66 shows a positive zeta potential of 26.3 mV; after loading with TMB and HRP, the zeta potential increased to 46.1 mV (Figure 1D). The positive zeta potential of UiO-66 is ascribed to the positive charge of the Zr⁴⁺ cation. Both TMB and HRP are positive structures; therefore, after the loading, the zeta potential of the nanoparticles increased to 46.1 mV. The pore structure and the intermolecular interactions of aromatic molecules are mostly responsible for the loading of TMB. The coordination between the Zr⁴⁺ cation of UiO-66 and chelating groups (e.g., -COOH and -SH) of HRP would be the reasons for HRP loading. The changes in the size and zeta potential clearly indicated the successful preparation of the nanocomposite UTH. Moreover, the presence of the characteristic peaks of TMB and HRP in the absorption spectrum of the nanocomposite also indicated the successful loading of TMB and HRP by UiO-66 (Supplementary Figure S1).



Characteristics of the nanocomposite preparation. TEM (A) image and XRD patterns (B) of the MOF (UiO-66), the size distribution (C), and zeta potential (D) of the MOF and the obtained nanocomposite UiO-66@TMB-HRP (UTH).



Response characteristics to H₂O₂

Then, we studied the response characteristics of the nanocomposite to H_2O_2 (Figure 2). The nanocomposite UTH contains both the enzyme (HRP) and the substrate (TMB); in the presence of H_2O_2 , the HRP will catalyze H_2O_2 to generate an intermediate that can oxidize TMB to a colored state. As

expected, the color of the nanocomposite solution changes from light milky to dark turquoise, and the absorption spectra clearly indicate the generation of the oxidized product of TMB (Figure 2A). After being treated with H_2O_2 , two strong absorption peaks around 650 nm and near 900 nm appeared. The intensity of the absorption peak increases with the increase in the concentration of the nanocomposite (Figure 2A). Even



when the concentration of H_2O_2 is as low as 0.1 mM, it can obviously cause discoloration of the nanocomposite in a short time, indicating that the nanocomposite has a high sensitivity to H_2O_2 (Figure 2B). It is worth mentioning that the photothermal effect of nanomaterials is often closely related to the intensity of their absorption peaks, which means that the oxidized nanocomposites will likely produce a strong photothermal effect under near-infrared light (900 nm) irradiation.

Photothermal effect

As shown in Figures 3A,B, under 900-nm light irradiation, the solution of 0.2 mg/ml UTH showed a very slight temperature increase. However, in the presence of 1 mM H_2O_2 , even a low concentration of UTH (0.05 mg/ml) could be sharply heated up by a 900-nm light irradiation. The heating rate and the maximum temperature increased with the increase in the concentration of nanocomposites. The temperature of the solution with 0.2 mg/ml UTH and 1 mM H_2O_2 can reach above 45°C within 3 min with

900 nm light irradiation, clearly indicating the excellent photothermal effect. It is worth mentioning that the concentration of H₂O₂ in the bacterial infection area is usually about 1 mM. In addition, during light irradiation, the local temperature of the nanoparticle surface is much higher than the solution temperature. These indicate that the nanoparticles provide the necessary basis for the sensitive response to H2O2 and efficient antibacterial activity in the infected area. Then, we calculated the photothermal efficiency by detecting the heating and cooling rates (Figures 3C,D), and the results showed that the photothermal efficiency of the nanocomposite UTH in the presence of H₂O₂ reached 18%. The photothermal conversion efficiency of most organic nanomaterials is between 20 and 50% (Li et al., 2020). Compared with these photothermal materials, the photothermal efficiency of 18% is slightly lower. Nevertheless, it is worth mentioning that indocyanine Green (ICG), as an organic molecule frequently used in photothermal therapy, has a photothermal efficiency of only 9% (Li et al., 2020). It indicates that 18% is enough for effective photothermal therapy.
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nm light irradiation. (B) Corresponding bacterial viabilities by counting (A). (C) Fluorescence images (stained with SYTO 9 and PI) of *E. coli* before (left) and after (right) being treated with UTH in the presence of H_2O_2 under 900-nm light irradiation.

Antibacterial effect

After verifying the sensitive response to H₂O₂ and the good photothermal effect of UTH, we then investigated the cascade antibacterial effect. We selected E. coli and S. aureus Gram-negative and -positive bacterial models, as respectively. The plate counting method was used to measure the antibacterial efficiency. The results showed that in the presence of only H_2O_2 (1 mM) or UTH (Supplementary Figure S2), NIR light irradiation could not cause significant antibacterial activity. However, in the presence of both H₂O₂ and UTH, NIR light irradiation can cause obvious antibacterial activity, and the antibacterial activity increases with the increase in the concentration of UTH. The IC₅₀ values of UTH on E. coli and S. aureus under NIR light irradiation with 1 mM H_2O_2 were around 150 and 450 µg/ml, respectively (Figures 4A,B). The significant antibacterial effect was observed in both Gram-positive and -negative bacteria, indicating such a cascade nanosystem has broad-spectrum antibacterial properties. Notably, the antibacterial effect of the photothermal system on the two stains is slightly different, and E. coli was more sensitive to the photothermal effect. The different antibacterial efficiencies may be related to the different structures of the two bacteria. S. aureus has a cell wall composed of peptidoglycan, which is relatively stable (Zhang et al., 2022a). However, the surface of E. coli is a cell membrane composed of phospholipid molecules, which is more fragile than the cell wall of S. aureus. Therefore, the photodynamic system has a stronger inactivation efficiency for E. coli. Subsequently, the antibacterial mechanism was primarily discussed by live/dead staining with a LIVE/DEAD bacterial viability kit. In such an assay, the green fluorescence (SYTO 9 dye) indicates the live bacteria and the red fluorescence (PI dye) indicates the cell wall-damaged dead bacteria. As shown in Figure 4C, most of the treated bacteria





have strong red fluorescence, indicating that the cascade nano-system can cause cell wall damage to kill bacteria.

Therapeutic effect against mouse wound infection

Finally, in order to verify the therapeutic effect on real wound infection, we constructed a mouse skin infection model. As shown in Figure 5, after being treated by NIR light irradiation or UTH, the mouse skin wounds infected by *S. aureus* showed obvious symptoms such as suppuration, and the wounds healed slowly. Even 10 days later, the wounds still had obvious dents. However, after being treated with UTH and NIR light irradiation, the wound purulent symptoms were less and the wound healing was faster. After 10 days, the wound basically healed. These results show that the cascade nano-system has a good therapeutic effect on skin wound infection.

Conclusion

In this study, the MOF nanocomposite with the characteristics of H2O2 and NIR light cascade response was successfully constructed, and the photothermal antibacterial activities were verified. The nanocomposites can respond sensitively and quickly to H₂O₂. Due to the catalytic oxidation, the color of the nanocomposite changes to dark turquoise in the presence of H₂O₂; meanwhile, a strong absorption peak in the near-infrared region around 900 nm appeared. The oxidized nanocomposite can convert near-infrared photons into thermal energy with an efficiency of 18%. This nano-system showed a strong inactivation effect on both Gram-negative and -positive bacteria. Under the conditions of 1 mM hydrogen peroxide and 0.5 W/cm² NIR light intensity, the IC₅₀ values of the MOF nanocomposite on E. coli and S. aureus were 150 and 450 µg/ml, respectively. This cascade response nanomedicine also showed a strong therapeutic effect on the mouse skin wound infection model. This study not only provides an effective photothermal antibacterial strategy but also offers a new idea for building precise nano-therapeutic systems.

Data availability statement

The original contributions presented in the study are included in the article/Supplementary material; further inquiries can be directed to the corresponding author.

Ethics statement

The animal study was reviewed and approved by Zhejiang Sci-Tech University.

Author contributions

ZZ: supervision, conceptualization, and writing—original draft preparation. CM, YZ, LL, and LW: methodology and investigation. CM, YZ, WZ, XL, and WB: writing—review and editing. ZZ and XL: funding acquisition.

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Supplementary material

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Precise dapagliflozin delivery by cardiac homing peptide functionalized mesoporous silica nanocarries for heart failure repair after myocardial infarction

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Myocardial infarction (MI) may cause irreversible damage or destroy to part of the heart muscle, affecting the heart's ability and power to pump blood as efficiently as before, often resulting in heart failure (HF). Cardiomyocyte death and scar formation after MI may then trigger chronic neurohormonal activation and ventricular remodeling. We developed a biocompatible and monodispersed mesoporous silica nanoparticles (MSN) divergent porous channel for dapagliflozin (DAPA) loading. After surface modification of the cardiactargeting peptides, the novel drug delivery system was successfully homed, and precisely released drugs for the hypoxic and weak acid damaged cardiomyocytes. Our biocompatible MSN- based nanocarriers for dapagliflozin delivery system could effective cardiac repair and regeneration *in vivo*, opening new opportunities for healing patients with ischemic heart disease in clinical.

KEYWORDS

myocardial infarction, dapagliflozin, nano-targeted drugs delivery system, heart failure repair, hypoxia-inducible factor $1-\alpha$ (HIF- 1α)

1 Introduction

Myocardial infarction (MI) contributes to more than 40% of sudden cardiac deaths and represents the leading cause of morbidity and mortality worldwide (Sathisha et al., 2011; Nigam 2007; Anwar et al., 2016). In myocardial ischemia area, broadscale myocardium tissue are impaired with apoptotic/necrotic cardiomyocytes. Due to the low proliferation capability of myocardial cells, the damaged myocardium tissue is always unable to be effective regeneration and restoration (Zhao, et al., 2022). Meanwhile, complications of heart failure (HF) after MI hospitalization may cause cardiomyocyte apoptosis and scar formation, subsequently triggering chronic neurohormonal activation and ventricular remodeling (Dudas et al., 2011; Roger 2013; Desta et al., 2015; Jenča D et al., 2021). The high prevalence of HF complications after hospital discharge greatly



reduces the quality of life of MI patients. It is estimated that approximately 13% of patients will experience HF complications 30 days post-discharge, with this number increasing to 20-30% in the first year after discharge (Hung et al., 2013; Sulo et al., 2016). Thus, HF is associated with three alarmingly high rates: high incidence rate, high mortality rate, and high rehospitalization rate. Scientists recently reported that the sodium-glucose cotransporter two inhibitor, dapagliflozin (DAPA) is a promising new drug for the treatment of HF. DAPA is approved by the Food and Drug Administration (FDA)-could reduce the risk of cardiovascular death and hospitalization for cardiac deterioration, increase survival rate and improve symptoms of HF in patients with decreasing ejection fraction (McMurray et al., 2019b, a). DAPA can effectively improve cardiac structure and function, weaken cardiac fibrosis and myocardial apoptosis, as well as inhibit inflammatory cytokines (Wang K et al., 2021). While the low selectivity, poor biodistribution and high systematic toxicity of DAPA greatly restrict its further clinical application (Wilczewska ZA, et al., 2012). Furthermore, treatment for MI is expensive and extensive, the prognosis is poor. Frequent episodes of HF and repeated emergency treatment in MI patients further increase the cost of treatment (Cowper et al., 2019). This situation led to the urgent construction of a novel drug delivery system (DDS) for HF repair. The novel DDS can be loaded with DAPA, which improves HF symptoms and increases patient survival. In addition, the novel DDS should be targeted precisely to improve drug utilization.

Novel drug delivery systems have been rapidly developed over the past few decades. Both native and synthetic nanomaterials have been extensively explored and proposed as the main components of DDS (Rodrigues et al., 2016; Swamy and Sinniah 2016; Mohanty et al., 2017). Recently, synthesized nanomaterials, including carbon materials (Bao R et al., 2017; Sun X et al., 2017), dopamine nanoparticles (Chakroborty et al., 2011; Das and Jana 2015), inorganic silica (Yang et al., 2010; Zhu et al., 2011), and polymeric assembly nanohybrids (Syed et al., 2012; Kerr et al., 2022), have been employed as nanocarriers for DDS. Porous silica has been recommended as an ideal DDS based on the biocompatibility of silica's drug embedding ability in the mesoporous channel (Meng et al., 2010; Che et al., 2014; Chen K et al., 2020; Chen YJ et al., 2020). Furthermore, mesoporous silica nanospheres (MSNs) have been widely used for drug delivery because of their optimal surface area for drug loading, their facile synthesis and surface functionalization by anchoring targeting ligands, stimuli-responsive agents and sensing molecules (Sahoo et al., 2014; Sapino et al., 2015; Ugazio et al., 2016; Jafari et al., 2019). MSNs are considered biocompatible with the human body, as indicated by numerous investigations. The therapeutic dose of MSNs for DDS is well below the toxic level for humans, making MSN an excellent option for clinical applications.

Cardiac homing peptide (CHP) is a short peptide that preferentially binds to ischemic myocardium, and its sequence is CSTSMLKAC. CHP was first screened in a study of selective targeting of random peptides to ischemic tissue *in vivo* and was shown to be a safe peptide that did not cause any significant impairment of left ventricular systolic function. CHP has the potential to be used in the development of targeted therapy drugs for ischemic lesions of myocardial tissue (Kanki et al., 2011; Won et al., 2013; Vandergriff et al., 2018).

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Here, we propose a conventional bi-phase approach for MSN synthesis for DAPA loading in heart failure repair. The hydrophobic and cation- π interaction properties of DAPA facilitate the efficient encapsulation of this drug in mesoporous channels (DAPA@MSN). Moreover, a CHP was functionalized on the surface of DAPA@MSN (DAPA@MSN-CHP). After intravenous injection of DAPA@MSN-CHP in a mouse model of HF after MI, this DDS efficiently and precisely accumulated in the HF site. DAPA-loaded MSNs have a slightly negative charge under normal physiological conditions but transform to a positive charge in the intracellular microenvironment of apoptotic cardiomyocytes because of the protonation effect under acidic conditions (Figure 1) (Wang et al., 2020). The results indicate that our DDS can effectively inhibit the apoptosis of cardiomyocytes, leading to viable myocardium preservation and cardiac function augmentation, laying a solid foundation for clinical HF repair.

2 Materials and characterization

Materials: Hexadecyltrimethylammonium bromide (CTAB), decahydronaphthalene, 1-octadecene (ODE) were purchased from Sigma-Aldrich. NaOH, cyclohexane and NH_4NO_3 were obtained from Shanghai Chemical Co., Ltd. Ammonia aqueous solution (28 wt%), tetraethyl orthosilicate (TEOS), (3-Aminopropyl) triethoxysilane (APTES), triethanolamine (TEA), decahydronaphthalene (98%) were purchased from Aladdin Industrial Inc. All chemicals were used as received without further purification.

Characterization: Transmission electron microscopy (TEM) measurements were carried out on a JEM 2100F microscope (Japan) operated at 200 kV. The samples were first dispersed in ethanol and then collected by using copper grids covered with carbon films for measurements. UV-vis–NIR absorption spectra were measured on a Shimadz spectrophotometer (UV-3150) (Japan) with wavelength range of 300 –1,200 nm, unless otherwise specified, all spectra were collected under identical experimental conditions. All animals underwent transthoracic echocardiography under anesthesia at 4 weeks after treatment using a VisualSonics Vevo 2,100 imaging system. The results were obtained by detecting Si content with inductively coupled plasma mass spectrometer in main organs.

2.1 Synthesis of mesoporous silica nanoparticles loading with DAPA and targeting peptides modification

2.1.1 Mesoporous silica fabrication

The mesoporous silica nanoparticles were synthesized as the procedure reported previously. Briefly, 16 ml of (25 wt%) CTAB solution and 90 mg of triethanolamine were added to 20 ml of

water and stirred gently at 60°C for 1 h in flask, then 1.5 ml of TEOS and 1 ml of cyclohexane was added to the solution and kept at 60°C with magnetic stirring for 24 h. The products were collected by centrifuging and washed by water and ethanol for several times, and then were dispersed in 30 ml of acetone and refluxed for 8 h to remove CTAB templates. The final products were washed with ethanol and dried in vacuum at 45°C for 8 h.

2.1.2 Loading of dapagliflozin (DAPA)

DAPA was dissolved in DMSO (2.0 ml). Mesoporous silica nanoparticles (5.0 mg) were added to the solution and the suspension was stirred at room temperature for 48 h. The DAPA molecules could be adsorbed in the mesopore channels. The as-prepared DAPA-loaded mesoporous silica nanoparticles were collected by centrifugation. The amount of the adsorbed guests was determined from the difference between the initial amounts of DAPA by measuring the UV absorbance from the supernatant liquid and quantified from a standard curve.

2.1.3 Cardiac homing peptides modification

To functionalization of DAPA@MSN, 3.0 g of silica was placed in a 100-ml round bottom flask with 50 ml of dry toluene and 2.4 g of 3-aminopropyltriethoxysilane (APTES). The mixture was stirred and refluxed at 110°C for 48 h under the protection of nitrogen. The white 3-aminopropyl functional silica washed with toluene, *n*-hexane and dichloromethane, respectively, and dried at 25°C for 12 h.

2.1.4 DAPA@MSN-CHP fabrication

First, 20 mg CHP, 10 mg EDC and 8 mg NHS were dissolved in 20 ml anhydrous DMF, and incubated at room temperature overnight to activate the carboxyl groups of FA. Then, DAPA@ MSN with NH_2 modification (20 mg) was added. The mixed solution was allowed to react for 12 h, and the DAPA@MSN-CHP were obtained by centrifugation at 12,000 rpm for 20 min, and then washed with methanol three times.

2.2 *In vitro* cellular targeting and cell viability of DAPA@MSN-CHP

2.2.1 Cell viability

All experiments were carried in 96-well plates. Cytotoxicity of MSN-CHP was tested *via* CCK-8 assay. Briefly, the primary cardiomyocytes isolated through enzymatic digestion were seeded into plate at 5×10^3 /well in 100 µL of DMEM/Low glucose (10% FBS, 100 units/mL of penicillin and 100 µg/ml of streptomycin), and incubated for 24 h. Then, 10 µL CCK-8 solution was added followed by12 h incubation. After 12 h incubation, the absorbance of each well at wavelength of 450 nm was measured using a microplate reader. Data were presented as mean \pm SD (n = 3).



FIGURE 2

TEM image (A), magnification TEM image (B) and SEM image of the bi-phase prepared MSN (C). DLS result of size distribution of MSN (D). Zetapotential data of MSN, DAPA@MSN, MSN-CHP and DAPA@MSN-CHP (E). DAPA release curve of DAPA@MSN-CHP under different pH values (F). The corresponding physical and chemical parameters of the DDS (G).

2.2.2 CLSM images of cellular uptake

The cellular uptake of DAPA@MSN-CHP was observed and imaged by confocal laser scanning microscopy (CLSM). Briefly, primary cardiomyocytes were seeded into 6-well plates at a density of 1×10^5 /well with 1 ml of DMEM/Low glucose (10% FBS, 100 units/mL of penicillin and 100 µg/ml of streptomycin) media and incubated for 24 h. After the treatment with DAPA@MSN-CHP, DAPA@MSN at a final concentration of 100 µg/ml, the cardiomyocytes were incubated for 12 h. Then, the cardiomyocytes were washed with PBS for three times and DAPI (1 $\mu g/ml$ in PBS) was used to stain nuclei for 30 min prior to being observed under CLSM.

2.2.3 Western-blot analysis

Cells were collected after 12 h incubation with PBS, DAPA, DAPA@MSN, and DAPA@MSN-CHP. Samples containing equal amounts of protein were electrophoresed using 10% SDS-PAGE, transferred onto polyvinylidene fluoride membranes and then incubated with specific primary antibodies. The blots were reacted with horseradish peroxidaseconjugated secondary antibodies and were detected using the enhanced chemiluminescence system (Santa Cruz Biotechnology, Inc.). The density of the band was quantified by densitometry and exposed to X-ray film (Eastman-Kodak, Rochester, NY, United States) using GAPDH levels as a control.

2.3 *In vivo* HF repair capability of DAPA@ MSN-CHP

2.3.1 In vivo HF repair

Animal experiments were approved by the Animal Research Ethics Committee of Shanghai Seventh People's Hospital. Animal experiments were performed according to the animal use and care regulation and the animal management rules of the ministry of health of the People's Republic of China. C57BL/6 mice (8-10 weeks old) were bought from Shanghai Institute of Family Planning Science Laboratory Animal management Department. (Shanghai, China). All efforts were made to minimize animal suffering. The myocardial ischemia model was constructed based on previously reported works. A total of 70 male 8- to 10-week-old C57 mice were used for this work (Luo L et al., 2016; Snider JC et al., 2021). For the myocardial ischemia model construction, all mice were received permanent coronary artery ligation. Mice were anesthetized with 160 mg/kg barbital sodium via intraperitoneal injection. Typically, before preparing the surgical model, ventilation system after endotracheal intubation is recommend to use when coronary artery ligation can be successfully and routinely performing the within 3 min. Subsequently, a 1.2 cm of small skin cut was conducted over the left chest according previously reported. After the dissection, both pectoral major and minor muscle retracted, then the fourth intercostal space can be successfully observed. Simultaneously, in order to open both the pleural membrane and pericardium, a mosquito clamp was used to cut a small hole at this intercostal space. When the clamp was slightly opened, partial heart section was gently squeezed out through this hole. The coronary artery ligation was carefully located, sutured, and ligated (from its origin \approx 3 mm) by a commonly used 6–0 silk suture. The surgical ligation was confirmed successfully when the anterior wall from left ventricular changed to pale. Immediately, the heart was put back to form the intrathoracic space followed by air manual evacuation and muscle and the skin closure through the previously used purse-string suture. During the recovery durations, the mice were allowed to breathe and carefully monitored, which was usually completed within 3-5 min. The artificial respiratory aid was not allowed during the recovery period. For the DAPA@MSN-CHP induced therapy, the C57BL/6 mice with myocardial ischemia were divided into four groups including the MI, DAPA, DAPA@MSN, and DAPA@MSN-CHP. Meanwhile, health mice were set as control group. Each group contained 4 mice. Each group contained 4 mice. The MI, MI + DAPA, MI + DAPA@MSN, and MI + DAPA@MSN-CHP groups were treated with 100 µL of PBS, DAPA, DAPA@ MSN, and DAPA@MSN-CHP solutions *via* tail injection, respectively one time every day for 3 days. Meanwhile, the control groups were also received tail vein injection of PBS one time every day for 3 days.

2.3.2 Cardiac function assessment

All animals underwent transthoracic echocardiography under anesthesia at 4 weeks after treatment using a VisualSonics Vevo 2,100 imaging system. During ultrasound process, mice were anesthetized with 3% isofluorane *via* a R500-Comapct Small Animal Anesthesia Machine (Shenzhen, China). Hearts were imaged 2D in long-axis views at the level of the greatest left ventricular diameter. Estimation of the function and fractional shortening were determined by measurement from views taken from the infarcted area. All measurements were done in random order, with the surgeon and echocardiographer being blind to the treatment groups.

2.3.3 Histopathological evaluation

Hearts were harvested and cut into 10 μ m-thick tissue sections. H&E and Masson's trichrome staining of normal, MI, DAPA, DAPA@MSN, and DAPA@MSN-CHP was performed. Image analysis related to viable myocardium and scar size was performed using NIH ImageJ software.

3 Results and discussion

3.1 DAPA@MSN-CHP fabrication and characterization

MSNs were fabricated using a novel biphase approach in a cyclohexane and water stratification system (Shen et al., 2014). The biphase stratification approach enables hydrolysis precursors in the interface. It conveniently regulates the nanoparticle assembly in the biphase interface by adding or changing other reactants in two different phases without affecting interfacial tension. The hydrophobic upper layer consisted of 25% tetraethyl orthosilicate (TEOS) solution in cyclohexane. The lower hydrophilic layer was a pure water solution mixed with cationic cetyl trimethylammonium cetyl bromide (CTAB) as the template and surfactant and 25% organic weak base triethanolamine (TEA) as a reducing agent. The dendritic hierarchical mesostructure with monodispersion was obtained via continuous interfacial growth in a facile one-pot strategy for 48 h (Figure 2A). A divergent mesoporous channel could be observed in the magnification transmission electron microscope (TEM) image (Figure 2B). The MSN pore size can be altered by changing the type of hydrophobic solvent in the upper laver.

The surface morphology of the MSN was also investigated using a scanning electron microscope (Figure 2C). The properties of the mesoporous channels and their uniform size were detected. The size distribution of our MSNs were also estimated by



dynamic light scattering (DLS). Results showed that the drug delivery carriers had a narrow unimodal size distribution with a ~96.2 nm peak, suggesting superior dispersity and uniform practical diameter (Figure 2D). Subsequently, the MSN surface was modified with the amino group in the ethanol solution

through aminopropyltriethoxysilane hydrolysis. In the following step, DAPA was encapsulated in the mesoporous channels. Then, CHP could be anchored on the silica surface by EDC/NHS reaction between the MSN amino group and the carboxyl group of the targeting peptides.



Zeta potential of MSN presents ~ -25 mV, contributing to the massive silicon hydroxyl group on the silica surface (Figure 2E). Interestingly, compared with free MSN, DAPA@MSN exhibits a weaker negative charge, mainly due to the positive charge of the amino group modification. DAPA@MSN-CHP has a positive charge in the aqueous solution, mainly because of the amino group consumption after the condensation reaction. The dominant positive charge of our DDS may originate from CHP, which facilitates the endocytosis of the DDS. The corresponding physical and chemical parameters of the DDS are listed in Figure 2G, showing all the sizes fluctuate minimally after drug loading or CHP modification. Loading efficiency meets theoretical efficiency estimations, further demonstrating that MSNs are the ideal nanocarriers for drug delivery. Li ZX et al. reported that small nanoparticles (< 200-300 nm) are normally taken up by cells via the endocytic pathway. Particle size, particle shape, and particle surface charge have a certain impact on the endocytic uptake of cells. Generally, particles with small particle size, rough surface and positively charged surface are more likely to be phagocytosed by cells (Li et al., 2012). The characterization results of the above materials suggest that DAPA@MSN-CHP may have good biocompatibility thanks to its small nanoparticle size, rough mesoporous surface, and positive charge characteristics.

MI leads to changes in the microenvironment at the site of cardiac injury. Constructing acidic pH-responsive DDS based on the weakly acidic microenvironmental characteristics of lesions can improve the value of drug utilization (Li et al., 2021). Finally, we investigated the drug release ability of DAPA@MSN-CHP under different pH conditions. Compared with the normal pH value of ~7.2, 92.1% of DAPA can be released after 24 h incubation in biological buffers with pH values of ~5.5 (Figure 2F). These results suggest that DAPA can continuously release under the weak acid in the HF site after MI which is beneficial to improve the utilization value of drugs.

3.2 *In vitro* investigation of DAPA@ MSN-CHP

Following the excellent performance of DAPA@MSN-CHP, we then estimated the intracellular localization of DDS in cardiomyocytes. DAPA@MSN and DAPA@MSN-CHP were co-loaded with Fluorescein isothiocyanate (FITC) for intuitive



observation under a Confocal Laser Scanning Microscope (CLSM). They were then added to two groups of cardiomyocytes for 12 h before CLSM observation. FITC signals at 550 nm were collected under 488 nm excitation. As shown in Figure 3A, DAPA@MSN-CHP presents stronger fluorescent spots than DAPA@MSN with no targeted protein modification. The fluorescent spots of DAPA@ MSN-CHP were mainly localized in the cytoplasm. At present, the target receptor of CHP and the mechanism of its specific targeting to ischemic tissue are not yet clear, but CHP has been proved to have good targeting delivery ability in a number of studies on ischemic myocardial targeted substance delivery systems. In a gene delivery study, Young-Wook Won et al. used CHP as a guide to achieve increased gene expression in H9C2 cells under hypoxic conditions (Won et al., 2013). Another study reported CHP as a targeting peptide to increase the uptake of peptide-labeled exosomes by oxidatively damaged H9C2 cells (Vandergriff et al., 2018). Our findings also indicate the precise binding ability of CHP to the receptor on the membrane of cardiomyocytes.

Previous reports indicate that hypoxia also affects most sites of heart diseases such as HF, cardiac arrest and heart attack (Jianqiang et al., 2015; Amofa et al., 2017; Hesse et al., 2018). The hypoxic conditions may originate from inadequate blood oxygen concentration delivery following vessel impairment in the damaged tissues. Furthermore, Hypoxia-inducible factor 1- α (HIF-1 α) promotes the adaptation to hypoxia-related stress through increasing oxygen delivery and decreasing oxygen consumption (Anand et al., 2007; Kumar et al., 2020; Nguyen et al., 2021). Accordingly, H9C2 cells were treated with hypoxia, and than the expression level of HIF-1 α was significantly increased (Figure 3B), indicating the successful establishment of myocardial hypoxia model. Based on this, we used the cardiomyocyte hypoxia model to study the effect of DAPA@MSN-CHP *in vitro*.

We then explored cell viability of cardiomyocytes under conditions of both normoxia and hypoxia based on our MSN-CHP. Figure 3C shows negligible cytotoxicity with the DDS concentration as high as 400 µg/ml in both normoxia and hypoxia, compared with a control group. These findings further demonstrate the superior biocompatibility of the MSN. Cell viability of our DDS was also studied under various conditions. Our DDS promoted significantly higher cell activity in hypoxia (~77%). In contrast, only ~51%, 61%, 70% of hypoxic cells survived in the solvent, DAPA, and DAPA@ MSN incubation conditions, respectively (Figure 3D). This result derives from the efficient and precise release of our DDS. DAPA successfully transported and gradually released the drug in the weak acid intracellular microenvironment of HF cells in DAPA@ MSN-CHP. Simultaneously, western-blot results demonstrated that our DDS improved the expression of cell survival-promoting factor (BCL-2) while reducing the expression of apoptotic factors Caspase-3 and BAX (Figure 3E). DAPA has been found to be effective in reducing cardiotoxicity and inhibiting apoptosis in vitro (Chang et al. 2022). Our results suggesting that the precise targeted delivery of DDS enhanced the intervention effect of DAPA on hypoxia-induced cardiomyocyte apoptosis.

3.3 *In vivo* HF repair capability of DAPA@ MSN-CHP

Since the targeting mechanism of CHP is not yet clear, the *in vitro* results are insufficient to demonstrate the targeting ability of DAPA@MSN-CHP. Therefore, we investigated HF repair efficiency using an MI mice model. The MI model was constructed with a temporary ligation of the left anterior descending coronary artery for 0.5 h. These mice were divided into four groups: pure MI, free DAPA, DAPA@MSN and DAPA@MSN-CHP groups, with normal mice as the control group (Figure 4A). Body distribution of DAPA@MSN-CHP results shows that the novel DDS had the highest cardiac-targeting efficiency compared with DAPA and DAPA@MSN (Supplementary Figure S1). Studies have reported changes in



protein expression and protein distribution in cardiomyocytes under ischemia and hypoxia conditions (Liu et al., 2014), suggesting that the unknown receptor protein of CHP may increase its binding to CHP through these changes to play a specific targeting role. In the study of Young-Wook Won et al. and Adam Vandergriff et al., CHP as a component of a drug delivery system was also reported to play an active role in targeting MI tissue *in vivo* (Won et al., 2013; Vandergriff et al., 2018).

Studies have shown that DAPA reduces heart failure exacerbations and improves symptoms in patients with heart failure and reduced ejection fraction (Kosiborod et al., 2020). After 1 month, the therapeutic efficiency of DDS was evaluated by detailed echocardiography analysis. The heart functions were carefully estimated by conventional M-mode echocardiography. Compared with the healthy group, the interventricular septum of the MI group without any administration became weak as septal amplitude decreased, proving the persistent and serious avascular necrosis in myocardium tissue. Compared with the no treatment condition, the morphology of the myocardial layer gradually improved in the echocardiogram image of the DAPA and DAPA@MSN groups. DAPA@MSN-CHP treatment significantly restored hearts, as shown by comparison with healthy hearts.

The conventional indicators of heart function were subsequently evaluated. The ejection fraction values of the left ventricle in normal, MI, DAPA, DAPA@MSN and DAPA@ MSN-CHP groups were 69.2%, 22.3%, 44.6%, 50.8% and 63.1%, respectively (Figure 4B). Similar trend was also detected in the ultrasound photographs of whole heart that MI + DAPA@MSN-CHP displayed most effective MI repair capability with highest ejection fraction values (Supplementary Figure S2). The fractional shortening values of the left ventricle in healthy, MI, DAPA, DAPA@MSN and DAPA@MSN-CHP groups were 42.3%, 12.3%, 24.8%, 29.7% and 37.6%, respectively (Figure 4C). In contrast with the MI, DAPA, and DAPA@MSN groups, the DAPA@MSN-CHP treatment resulted in optimal performance in restoring both ejection fraction and fractional shortening. There was no significant difference between the normal group and the DAPA@MSN-CHP groups, demonstrating the superior repair capability of the DDS.

3.4 Histopathological evaluation of DAPA@MSN-CHP for HF repair

In the process of heart failure, pathological changes in cardiac tissue occur, mainly manifested as myocardial remodeling and inflammatory infiltration (de Boer et al., 2019). After various administrations, heart sections were stained with hematoxylin and eosin (H & E). The DAPA@MSN-CHP administration group was similar to the healthy mice and demonstrated a normal histoarchitecture with intact myocardial membranes, similar oval nuclei and regular cross striations (Figure 5A). In

contrast, other groups showed histological alterations with myocardial separation and few scattered inflammatory cells. These atypical pathological phenomena were mainly caused by inflammatory leukocytes and wavy fibers. Masson's trichrome staining of heart tissue was also used to validate the associated collagen deposits and fibrotic transformation associated with MI. Pure MI, DAPA, and DAPA@MSN showed thick and dense interstitial collagen fibrils (blue) related to necrotic myocytes. DAPA@MSN-CHP and control groups exhibited normal heart histoarchitecture with slight interstitial collagen fibrils stained with blue (Figure 5B). All histopathological examination results demonstrated the superior therapeutic efficiency of MSN-based DDS to repair cardiac function following MI.

3.5 Biocompatibility evaluation in vivo

Finally, we herein evaluate the systematic toxicity of our MSN-based DDS via H&E tissue staining analysis and blood biochemistry assessment. The tissue of main orans including heart, liver, lung, spleen and kidney were dissected after the tail vein injection of MSN-CHP and then they were stained by H&E. As shown in Figure 6A, there was ignorable tissue damage in MI, DDS treated group, as compared with the control group (normal mice), testifying the superior biocompatibility of our MSN-based nanocarriers. For the blood biochemistry analysis, as displayed as liver functional makers including ALT (alanine aminotransferase), AST (asparatate aminotransferase), ALP (alkaline phosphatase), TBIL (total bilirubin), BUN (Blood urea nitrogen) and CRE (creatinine), no any variations of hepatic toxicity were found after MSN-CHP administration (Figure 6B). As indicator for heart muscle damage, CK (creatine kinase) value in the blood of MSN based DDS treated mice was also maintained at normal level (Figure 6B). The above results provide a validation that MSN-based nanocarriers bore no remarkable side effect in vivo.

4 Conclusion

We developed a straightforward approach to fabricate biocompatible and monodispersed MSNs of negligible toxicity. Their large surface area facilitated the DAPA encapsulation in the mesoporous channel, and the nanocarrier surface was then functionalized for cardiac-targeting peptides. The DDS was highly efficient in MI region in HF, and the DAPA could be precisely released in the hypoxic, apoptotic and weak acid intracellular environment. DAPA@MSN-CHP demonstrated optimal therapeutic efficiency toward MI model mice, restoring the MI hearts and making them comparable with those of the healthy group. MSN-based DDS provides a possible application of precise and effective repair of MI in clinical.

Data availability statement

The original contributions presented in the study are included in the article/Supplementary Material, further inquiries can be directed to the corresponding authors.

Ethics statement

The animal study was reviewed and approved by the Animal Research Ethics Committee of Shanghai Seventh People's Hospital.

Author contributions

Conceptualization, LY, QW and YM; methodology, LY, QW and YM; software, YL and HY; validation, LY, QW and YM; formal analysis, YL; investigation, HY; resources, LY; data curation,QW; writing-original draft preparation, LY, QW and YM; writing-review and editing, LX and ML; visualization, YM; supervision, LY; project administration, LX; funding acquisition, ML. All authors have read and agreed to the published version of the manuscript.

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Supplementary material

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Application of nanomaterials in diagnosis and treatment of glioblastoma

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Diagnosing and treating glioblastoma patients is currently hindered by several obstacles, such as tumor heterogeneity, the blood-brain barrier, tumor complexity, drug efflux pumps, and tumor immune escape mechanisms. Combining multiple methods can increase benefits against these challenges. For example, nanomaterials can improve the curative effect of glioblastoma treatments, and the synergistic combination of different drugs can markedly reduce their side effects. In this review, we discuss the progression and main issues regarding glioblastoma diagnosis and treatment, the classification of nanomaterials, and the delivery mechanisms of nanomedicines. We also examine tumor targeting and promising nano-diagnosis or treatment principles based on nanomedicine. We also summarize the progress made on the advanced application of combined nanomaterial-based diagnosis and treatment tools and discuss their clinical prospects. This review aims to provide a better understanding of nano-drug combinations, nano-diagnosis, and treatment options for glioblastoma, as well as insights for developing new tools.

KEYWORDS

glioblastoma, glioma, tumor targeting, nanomaterial, blood-brain barrier

1 Introduction

Glioblastoma (GBM) is a general term for tumors derived from glial cells and neuronal cells. It is the most common malignant tumor in the brain, accounting for 40%–50% of all intracranial tumors (Chen et al., 2017). The World Health Organization (WHO) classifies gliomas by cell type: astrocytoma, glioblastoma, and oligodendroglioma, and by malignancy grade (WHO I, II, III, IV). In 2021, the WHO CNS5 updated the molecular biomarkers for different tumor types, bringing more benefits and meaningful guidance to clinical practice. The WHO CNS5 has taken a new approach to classifying gliomas, glioneuronal tumors, and neuronal tumors, dividing them into six families: adult-type diffuse gliomas, pediatric-type diffuse low-grade gliomas, pediatric-type diffuse high-grade gliomas, circumscribed astrocytic gliomas, glioneuronal and neuronal tumors, and ependymomas (Louis et al., 2021). At present, temozolomide chemotherapy combined with radiotherapy after maximum feasible resection is the primary clinical

treatment for adult primary GBM. With the emergence of many targeted therapies, bevacizumab and programmed death -1 (PD-1) have also been used in the standardized treatment of GBM (Khasraw et al., 2012; Cloughesy et al., 2019; Zhao M. et al., 2020; Detti et al., 2021). However, GBMs grow infiltratively and are not clearly demarcated from normal tissues. They are, therefore, difficult to completely remove surgically and have high recurrence rates. Meanwhile, the blood-brain barrier (BBB), a highly selective semipermeable structural and chemical barrier (Cai et al., 2018), hinders drug delivery to the brain. Thus, drugs rarely reach effective therapeutic concentrations at the tumor site, limiting the effect of radiotherapy and chemotherapy. Due to these obstacles, GBMs have high mortality rates. To overcome this, nanomaterials have been used as a new treatment modality. They have many interesting characteristics, such as small size and targetable transport, making them good delivery tools for drugs, genes, or proteins across cells or the BBB.

Nanomaterials are materials with at least one dimension in the three-dimensional space in the nanoscale or composed of such materials as basic units. Nanomaterials have small particle sizes, controllable texture, and strong plasticity. In GBM treatment, nanomaterials are mainly used as carriers for radiotherapy and chemotherapy drugs (Zhao M. et al., 2020), helping drugs cross the BBB and maintain the necessary blood drug concentration. Additionally, nanomaterials can cause tumor cell necrosis by affecting the tumor microenvironment (Zanganeh et al., 2019). For example, a nanocomposite drug consisting of a polyglycerol functionalized doxorubicincontaining nanodiamond was designed to re-adjust the inhibitory glioblastoma immune microenvironment. This nanocomposite provided GBM with immunosuppressive microenvironment by activating autophagy. This rebuilding promoted the anti-GBM immune response and strengthened the activation of dendritic cells (Li et al., 2019).

The primary purpose of nanomaterials is to improve the solubility, stability, and effective concentration of drugs and reduce their systemic toxicity (Zhao M. et al., 2020). The main nanomaterials used in current research are nanoparticles (NPs). They can be inorganic, polymeric, or bionic. Each type has its advantages. Inorganic NPs, for example, are highly stable, and their physical and chemical properties can be adjusted by using suitable materials and sizes (Bharti et al., 2019). In addition, polymeric nanoparticles can encapsulate drugs by electrostatic or covalent bonding, maintaining their blood concentration and thereby improving the bioavailability of drugs. Most importantly, modified polymeric NPs can aggregate at specific sites. Biomimetic nanocomposites have higher biological stability and can escape the immune system, allowing drugs to act on the target continuously without being cleared by the liver and kidney. Besides nanoparticles, other nanomaterials (e.g., liposomes, quantum dots, cellular and extracellular vesicles, or virus-like particles) are frequently used in GBM treatment research.

The emergence of nanomaterials has brought opportunities for the diagnosis and treatment of GBM, and their various characteristics can help overcome the current clinical challenges of GBM. Here, we review the current progress on the applications of nanomaterials in GBM treatment research to provide insights for developing new nano-drug combinations, nano-diagnosis, and GBM treatment schemes.

2 Blood brain barrier and delivery mechanism of nanomaterials

2.1 Blood brain barrier

The blood-brain barrier (BBB) is a specialized structure within the central nervous system that acts as a physical and metabolic barrier restricting transport between the blood and neural tissues. It consists of brain microvascular endothelial cells, pericytes, astrocytes, neurons, and a basement membrane. This physical barrier protects the brain and maintains the stability of the intracranial environment. The excellent barrier properties of the BBB protect the brain from harmful macromolecules and pathogens present in the blood (Xie et al., 2019). However, this barrier also hinders traditional drug delivery and affects drug efflux. Currently, most smallmolecule drugs and almost all macromolecular drugs (e.g., recombinant proteins, therapeutic antibodies, and nucleic acids) cannot cross the BBB (Baratta, 2018). Thus, more and more researchers pay attention to nanomaterials, as shown in many clinical trials on the diagnosis and treatment of GBM (Dong, 2018). The delivery mechanisms of nanomaterials can be divided into passive targeting and active targeting (Zhao M. et al., 2020).

2.2 Passive targeting

In passive targeting, a drug of a specific size is injected through the BBB into the abnormal, porous vascular endothelium of the tumor. Since the tumor vascular endothelium lacks appropriate drainage, the drug remains in the tumor area for a long time. This phenomenon is also known as the enhanced permeability and retention (EPR) effect and was first proposed by Matsumura and Maeda in 1986 (Matsumura and Maeda, 1986). However, current studies have shown that the EPR effect is unstable and varies significantly among different tumors. Besides, the EPR effect achieved in rodent models cannot be reproduced clinically because human tumors have heterogeneity or lack fenestrations in the tumor endothelium, acidic and anoxic areas, low and heterogeneous pericyte coverage, and high interstitial fluid pressure induced by a dense extracellular matrix (Danhier, 2016).

2.3 Active targeting

Active targeting is a non-invasive approach that involves transporting drugs to target organs using site-specific ligands. In particular, drug-loaded nanocarriers that can target brain capillary endothelial cells and brain tumor cells show potential in oncology (Béduneau et al., 2007). In active targeting, nanocarriers enter cells by taking advantage of ligand-receptor interactions. So far, active targeting nanomaterials have been applied to the diagnosis and treatment of various malignant tumors, such as liver cancer, lung cancer, and lymphoma (Falgàs et al., 2020; Kaps and Schuppan, 2020). Developing active targeting nanomaterials able to cross the BBB requires understanding how to use the brain capillary endothelium. Nevertheless, the active targeting of the BBB represents a promising non-invasive strategy for improving antiglioblastoma drug delivery (Miranda et al., 2017). Active targeting can be further subdivided into adsorptive-mediated, carrier-mediated, receptor-mediated, and cell-mediated delivery, which target cells in different ways.

2.3.1 Adsorptive-mediated delivery

Adsorptive-mediated endocytosis (AMT) can deliver drugs to the brain through the BBB by allowing cationic molecules to bind to and be adsorbed onto the surface of the endothelial cell lumen. Adsorption-mediated transcytosis is initiated by the electrostatic interaction between positively charged ligands and negatively charged cell membranes. Nanocarriers then enter cells through clathrin-dependent endocytosis (Patel and Patel, 2017). Cationic proteins combined with cellpenetrating peptides (CPPs) can improve transport. With their short amino acid sequence, CPPs can interact with cell membranes and pass through cell membranes by energydependent and energy-independent mechanisms. Studies have shown that cationic CPPs have significantly more flux to the brain parenchyma than amphiphilic CPPs in vivo (Komin et al., 2017). Due to electrostatic interactions, cationic CPPs are easily captured by intracellular organelles to some extent. The drug-CPP linkage type substantially affects their ability to cross the BBB. Liu et al. (2014) compared the effects of amide, maleimide, and disulfide linkers linking the endorphin 1 to the CPP synB3 on BBB crossing efficiency. The disulfide linkage was the most efficient, and it was able to release the free drug in the brain. The adsorption-mediated transcytosis combined with a targeting strategy can effectively improve the EPR effect and reduce nonspecific uptake. Srimanee et al. (2018) developed a non-covalent CPPtargeting peptide (CPP-TP) complex with the CPP PepFect 14 and a hexaglutamate-modified angiopep-2 (ANG), as a targeting peptide. This complex showed enhanced penetration ability and glioblastoma cell specificity as an siRNA carrier. During the last decades, nanoparticles with various compositions have been developed, such as polymeric

nanoparticles (PPs), gold NPs, gadolinium NPs, selenium NPs, or protein-based NPs (Gupta and Sharma, 2019).

2.3.2 Carrier-mediated delivery

Carrier-mediated delivery is initiated by combining a designed nanocarrier and a specific transporter protein (Bourganis et al., 2018). This drug delivery system consists of nanocarriers (such as liposomes, NPs, polymeric micelles, dendrimers, or polymersomes) and ligands for various receptors, including transferrin receptor (TfR), lactoferrin receptor (LfR), low-density lipoprotein receptor (LDLR), and folate receptor (FR) (Chen et al., 2014; Wang et al., 2015). Ligand-modified drug carriers deliver drugs to the receptor-containing target cells like "guided missiles" (Liu et al., 2021).

Based on transport direction and substrate, transport modes can be divided into three categories: 1) The system pumping blood into the brain, which transports essential nutrients to the brain, including glucose, amino acids, and nucleotides. 2) The drug efflux pumps expelling exogenous substances out of the peripheral circulation to prevent them from entering the brain. 3) The efflux system from the brain to the blood, which mainly removes metabolic waste and neurotoxic substances in the interstitial fluid of the brain. Among them, the pumping system is the breakthrough point of carrier-mediated delivery. Nanocarriers going through the pumping system are generally designed as nutritional analogs with a high affinity for transporters, so their molecular weight is generally small. Cellmediated delivery can be used in anticancer therapy (Naik et al., 2021). For example, Naik et al. (2021) developed doxorubicincontaining liposomes and confirmed that conjugating these liposomes with a ligand mimic increased their antiproliferative activity on cancer cells overexpressing the corresponding receptor.

2.3.3 Receptor-mediated delivery

Targeting receptors that mediate endocytosis allows more robust targeting than with adsorption-mediated and carriermediated delivery because of the high specificity of the ligandreceptor interaction. In one study, cationic liposomes loaded with temozolomide were encapsulated in a multilayer crown of plasma proteins with a natural affinity for the folic acid (FA) receptor, which is highly expressed in the BBB (Tang et al., 2021). In an in vitro BBB model, these cationic liposomes with multilayered biomolecular crowns exhibited high ingestion by endothelial cells of human umbilical vein, which promoted the anticancer effect of temozolomide in U-87 MG cells (Arcella et al., 2018). Mram Alho et al. (Ramalho et al., 2018) developed stable polylactic acid-co-glycolic acid nanoparticles functionalized with the OX26 monoclonal antibody for the delivered transferrin receptor. These nanoparticles temozolomide, an anti-glioma agent.

Up to now, researchers have used many ligands of receptors on the BBB or glioma cells as targeting moieties for BBB



crossover and/or glioma-targeted drug delivery, such as peptides (Wang et al., 2015). In addition, more and more studies have proved that BBB/glioma-specific targeting nanocarriers can help drugs selectively target glioma cells, increasing their therapeutic efficiency while reducing systemic toxicity (Wang et al., 2017; Ramalho et al., 2018). Receptor-mediated drug delivery may allow membrane-impermeable drugs to penetrate target cells and activate natural signaling cascades (McPherson et al., 2001).

Besides directly targeting tumor surface receptors, the receptor-mediated pathway includes three important steps, including the formation of ligand-receptor complexes, transport through the cytoplasm of endothelial cells, and extracellular secretion outside the base of the BBB (Azarmi et al., 2020). In the second step, the lysosomal system may threaten the integrity of the drug. This can be bypassed by using cationic molecules and pH-sensitive drug carriers (Shir et al., 2006). However, receptor-mediated transfer also faces some problems. The high affinity leads to strong interactions between multiligand receptors on the lumen side of the BBB, limiting the entry of therapeutic molecules into the brain parenchyma. By contrast, using ligands with lower affinities can lead to higher drug release into the brain, but this requires administering higher doses, which is usually not applicable (Clark and Davis, 2015).

2.3.4 Cell-mediated delivery

In recent years, cell-mediated transcytosis has received increasing attention because immunogenicity and instability can hinder the use of antibodies and peptides (Yu et al., 2016). Neural stem cells, mesenchymal stem cells, neutrophils, macrophages, and exosomes, among others, have an intrinsic tumor-homing ability, allowing them to target malignant GBMs for drug delivery (Cho et al., 2019). Neutrophils have been widely studied in the treatment of brain tumors, especially for the treatment of postoperative recurrent tumors. For example, inspired by the ability of macrophages to cross the BBB, Tingting et al. encapsulated catalase into silica nanoparticles to produce a nanoplatform called CAT@SiO2-ICG (CSI). Next, they further encapsulated CSI into AS1411 aptamer-modified macrophage exosomes to form CSI@Ex-A (Wu et al., 2022) (Figure 1). Similarly, recent studies have confirmed that cellmediated delivery systems can contribute to the clinical treatment of gliomas. Xue et al. (2017) have shown that neutrophils carrying liposomes that contain paclitaxel, can penetrate the brain and inhibit GBM recurrence in mice whose tumors have been surgically removed. Similarly, one study demonstrated that the dendritic cell-mediated delivery doxorubicin-polyglycerol-nanodiamond of composites stimulated GBM cells immunogenicity and elicited an antiglioblastoma immune response (Li et al., 2018). These researches have revealed the feasibility of cell-mediated delivery for GBM treatments and laid the foundation for a translational study of this therapeutic paradigm to improve clinical outcomes in patients with malignant brain tumors.

2.4 Nano-assisted GBM diagnosis

Besides treating tumors, nanomaterials can be used in combination with imaging modalities, such as computed tomography (CT), functional magnetic resonance imaging (FMRI) (Richiardi et al., 2011), and positron emission tomography (PET) (Fink et al., 2015) to increase the

sensitivity and accuracy of tumor detection. Moreover, ultrasound (Imbault et al., 2017) and fluorescence imaging (Li et al., 2017) have been widely used in the clinical diagnosis and treatment of GBM. First, under normal physiological conditions, the physical and chemical barriers of BBB can efficiently transport necessary particles to human brain and selectively discharge harmful or excessive materials (Wu et al., 2019). Under pathological situations, practically all kinds of macromolecular drugs (proteins, antibodies, peptides, developers, etc.,) and small molecular drugs can pass through the BBB (Umlauf and Shusta, 2019). Although multiple brain tumor diagnosis strategies exist, their various limitations have affected their efficacy in the diagnosis of GBM. Second, the perpetual drainage and circulation of blood-cerebrospinal fluid and interstitial fluid stop most macromolecules from entering the bloodstream and diagnostic drugs from entering the brain. Moreover, tumor-acquired characteristics of the brain prevent drug from penetrating into tumors. In particular, tumor-induced endothelial cell tight junction damage increases drug penetration, leading to a heterogeneous distribution (Groothuis, 2000; Roose et al., 2003; Arvanitis et al., 2020).

Magnevist, an extensively used clinical contrast agent, plays an important role in evaluating tumors and their recurrence in magnetic resonance imaging (MRI) (McDannold et al., 2012; Kim, 2020). However, due to its short half-life, maintaining a sufficient concentration in the tumor requires repeated high-dose injections. Vascular leakage (Zhao H. L. et al., 2020), pseudoprogression (Brandsma et al., 2008), and pseudoreaction (Batchelor et al., 2007; Dhermain et al., 2010) after radiotherapy or anti-angiogenesis therapy also affect MRI accuracy. Besides, magnetic resonance spectroscopy and PET are often used to quantify and describe the development of cerebral tumors. Integrated with tracers and positron radionuclides in existence, they will achieve evaluable and reliable tissue autoradiography (Kratochwil et al., 2019), providing important information needed in diagnosis on tumors. But it is the requirements for accurate diagnosis of cerebral tumors that still cannot meet. In response to these hurdles (substandard specificity and accuracy, ephemeral contrast agent half-life, large requirement for imaging, etc.,), and to improve imaging sensitivity in diagnosis, nanoparticles have been developed.

2.4.1 Nanomaterials for MRI

The properties of nanomaterials greatly help overcome the difficulties faced by traditional radiographic agents. For example, Self-assembled nanoparticles of amphiphilic gadolinium chelates show extremely high Gd3 + loading capacity for enhanced imaging (Othman et al., 2011). Coupling gadolinium to interleukin 13 (Li et al., 2015), an arginine-glycine-aspartate (RGD) peptide (Zhan et al., 2010; Sun et al., 2014; Richard et al., 2017), an epidermal growth factor receptor (EGFR) mutant antibody (EGFRvIII) (Hadjipanayis et al., 2010) or anti-gd2 antibody (Shah et al., 2013) can markedly enhance

tumor targeting. Gadolinium metal fullerene nanoparticles have a cage surface charged by amino (-NH3+), showing incredible 1H MR relaxation. These data suggest that composite nanoprobes can serve as alternatives to magnetic resonance contrast agents (Lajous et al., 2018). The clinical applications of nanodiagnostic agents consisting of other magnetic materials especially iron and manganese, have also been researched. Superparamagnetic iron oxide nanoparticles (SPIO) have been paid tremendous clinical attention due to their magnetic properties. For example, the tungsten-doped iron oxide crystal (WFe) contrast agent has a sterling T1-weighted effect. The WFe nanoparticles possessed high average T1, 22% shorter than that of ferritin at the injection site (Clavijo Jordan et al., 2014). In addition, the superparamagnetism of SPIO is beneficial to the aggregation of the magnetic target. For example, the synergistic release of SPIO by focused ultrasound and magnetic targeting notably increases its accumulation in the brain parenchyma (Lee et al., 2019). In addition, modifying SPIO with a brain tumor-targeting peptide cRGD (Richard et al., 2017), a targeting antibody EGFRvIII (Hadjipanayis et al., 2010), or the targeting toxin chlorotoxin (Stephen et al., 2014) can enhance its tumor-targeting specificity. Finally, manganese, chelated with albumin-binding molecules, has been evaluated as a new contrast agent in both subcutaneous and in situ brain tumor models (Zhou et al., 2019).

2.4.2 Nanomaterials for CT imaging and Surface Enhanced Resonance Raman Scattering

Gold nanoparticles have incredible biosafety properties and are readily synthesized. Thus, a series of gold NPs with Surface Enhanced Resonance Raman Scattering (SERRS) signals were developed to guide brain tumor resection (Saha et al., 2012; Karabeber et al., 2014). These nanoparticles show strong signal intensity in SERRS after 24 h of injection, yielding a clear tumor contour. Besides gold nanoparticles, a new PET/CT imaging reagent named ⁶⁸galliumBNOTA-PRGD2 (68 Ga-PRGD2) showed some GBM diagnosis success (Li et al., 2014). Since GBM cells overexpress avb3, the target integrin of 68 Ga-PRGD2, the nanoparticle accumulates in GBM cells. The sensitivity and specificity of 68 Ga-PRGD2 for GBM grading were respectively 12% and 25% superior to those of the clinical PET/CT agent 18Ffluorodeoxyglucose (FDG). Moreover, Zhao et al. (2016) used the integrin a5b1 to enhance the specificity of 99mTc-HisoDGR SPECT/CT probes. 99mTcHisoDGR yielded precise contours in subcutaneous and in situ models 1.5 and 2 h after injection, respectively.

2.4.3 Nanomaterials for optical imaging

Currently, fluorescence imaging surgery guidance is widely used in clinical practice. Nanoparticles can be used as fluorescent dye carriers or fluorescent dyes to enhance optical imaging, dramatically reducing the failure rate of surgery (Wu et al., 2019). Several nanoparticles, including liposomes and polymer NPs, can effectively deliver live imaging fluorescent dyes to brain tumor sites during the operation. Indocyanine green, a nearinfrared fluorescent probe approved by the FDA, is widely used in tumor tracking and photothermal therapy. Liposome nanoparticles (LP-iDOPE) combined with VEGF-bevacizumab can enhance brain tumor imaging. One day after injection and before surgery, LP-iDOPE enhanced tumor localization (Suganami et al., 2015). Biodegradable polymers, such as polyalkylcyanoacrylate (Vauthier et al., 2003) and poly (lacticco-glycolic acid) (Han et al., 2016; Guan et al., 2017; Orunoğlu et al., 2017), are widely used to prolong the retention period of nanoparticles in vivo. Besides, polymer nanoparticles can load cerebral tumor targeting particles like chlorotoxin or anti-PDGFRD antibody (Monaco et al., 2017). A novel optical contrast agent with good biocompatibility was developed by modifying a polymer nanomatrix with fluorescent dyes and silver nanosheet clusters. The contrast of these nanoparticles was 90% higher than that of the control (Ray et al., 2014).

Being activated by light at a particular wavelength, some nanoparticles can emit light at another (such as quantum dots (Liu M. X. et al., 2017; Tang et al., 2017) and upconversion luminescent materials (Wang et al., 2020)), making them ideal fluorescent dyes for cancer targeting and imaging. Quantum dots are widely used for biological imaging diagnosis due to their excellent water solubility, low fluorescence quenching rate, high fluorescence quantum yield, and stable chemical properties (Wu and Yan, 2013). Thus, quantum dots modified with tumortargeting molecules can achieve specific biological imaging. Moreover, near-infrared technology can increase their fluorescence and Raman signals by several orders of magnitude (Gill and Le Ru, 2011), making them more sensitive for imaging in preclinical and clinical studies. Upconversion nanoparticles have considerable light stability, no fluorescence scintillation, deep tissue penetration, and low light damage effects (Gu et al., 2013; Liu et al., 2021; Mohan and Poddar, 2021), making them promising in vivo imaging probes (Idris et al., 2012; Yang et al., 2012). However, although these nanoparticles have strong tissue penetration ability, they cannot penetrate brain tissue like traditional fluorescent dyes (Li and Wang, 2018). To overcome this problem, research on second near-infrared transparent window (NIR-II, 1,000-1700 nm) fluorescent nanoprobes has gradually developed. NIR-II fluorescent nanoprobes have a stronger ability to penetrate tissues and achieve higher image fidelity. Ag2S quantum dots are one of the typical NIR-II fluorescent probes. Coupled to cyclic RGD peptides, the biocompatible NIR-II Ag2S fluorescent probe can achieve targeted labeling and imaging of U87 cells (Zhang et al., 2012). Considering the enhanced penetration depth of fluorescence signals, Qi et al. (2018) further analyzed the tissue penetration depth of NIR-II imaging probes. Near-infrared aggregation-induced emission under excitation with a 1,300 nm NIR-ii laser allowed the group to observe 5 mm blood vessels at a depth of 1,065 mm in the brain.

2.4.4 Nanomaterials for multimodal imaging

Since each diagnostic method has its own advantages and disadvantages, combining different diagnostic methods can optimize the outcomes. The development of dual-mode imaging nanoparticles has majorly impacted biomedical research. Angiopep-2 (ANG, TFFYGGSRGKRNNFKTEEY) coupled with upconversion dual-mode imaging nanoparticles (ANG/PEG-UCNPs) and Gd were constructed for targeting GBM. Compared with non-ANG and Gd-DTPA imaging, the nanoprobe yielded a significantly enhanced T1-weighted magnetic resonance contrast for glioblastoma. T2-weighted MRI also shows great potential for identifying clear glioblastoma borders (Ni et al., 2014). These results confirmed the advantage of MRI combined with fluorescent nanoprobes in GBM diagnosis. However, although dual-mode imaging improves the accuracy of high-resolution information, it still does not achieve overall tumor visualization. To improve the diagnosis accuracy and sensitivity, various imaging nanoparticles have been explored. For example, gold silicon-based SERS nanoparticles were used for three-mode imaging applications, where gold NPs were encapsulated by Gd3+ ions. MRI, photoacoustic imaging, and SERS showed clear tumors. The three-dimensional rendering of the magnetic resonance and photoacoustic images showed good co-expression signals in the tumor (Kircher et al., 2012; Neuschmelting et al., 2018).

2.4.5 Summary

Traditional GBM diagnosis faces some tough obstacles, such as the BBB and tumor heterogeneity. Although nanotechnologies have solved these problems to a certain extent, single-dimension diagnosis remains extremely limited due to the complexity of GBM. Combining multiple diagnostic methods will become a critical research field. Therefore, it is essential to construct various nanomaterials meeting the requirements for multimode combinations. Table 1 lists existing diagnosis nanomaterials and their related mechanisms.

2.5 Nano-assisted GBM therapy

2.5.1 Ferroptosis

Ferroptosis is an iron-dependent programmed cell death distinct from apoptosis, necrosis, pyroptosis, and autophagy (Bogdan et al., 2016; Zheng et al., 2017; Shen et al., 2018; Zhang et al., 2020). Excessive iron reacts with hydrogen peroxide (H2O2), generating hydroxyl radicals and singlet oxygen in cells (this process is known as the Fenton reaction). High hydroxyl radical levels eventually lead to cytotoxic lipid peroxidation. Since ferroptosis and apoptosis are radically different in mechanism and phenotype, combination therapy targeting these two processes may be a strategy for treating GBM. Based on this idea, Yulin *et al.* proposed an innovative local chemotherapy approach. They constructed iron oxide

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TABLE 1 Nanomaterialbased diagnostic for glioma.

Application	Nanomaterial	Highlight
MR imaging	Self-assembling nanoparticles of amphiphilic gadolinium chelates	High loading capacity of Gd3+ ions, enhanced imaging effect
	conjunction of gadolinium with interleukin (IL)-13, the arginine-	Significantly enhanced tumor targeting ability
	Glycine–aspartic acid (RGD) peptide, the epidermal growth factor receptor (EGFR) deletion mutant (EGFRvIII) antibody or the anti-GD2 antibody	
	Gadolinium metallofullerene nanoparticles	Excellent 1H MR relaxivity
	Tungsten doped iron oxide crystal	Excellent T1-weighted effect
	Superparamagnetic iron oxide nanoparticles	Magnetically controlled target accumulation
	SPIO with brain tumor targeted peptides cRGD, targeted antibody EGFRvIII and targeted toxin chlorotoxin (CTX)	Specific targeting in tumors
Surface Enhanced Resonance Raman Scattering (SERRS) and CT imaging	Au NPs	A stronger SERRS signal intensity 24 h postinjection, and then an accurate outline of the tumor
	68 Ga-PRGD2	Enhanced sensitivity and specificity of glioma grading
	99 mTc-HisoDGR SPECT/CT	Clear visualization in both subcutaneous and orthotopic models respectively 1.5 h and 2 h post injection
Optical imaging	Indocyanine green (ICG) loaded liposomal formulated nanoparticles (LP-iDOPE)	Enhanced imaging effect of brain tumors, excellent tumor- specific localization
	Poly-alkyl-cyano acrylates (PACA) and poly lacticcoglycolic acid (PLGA)	Extending the circulation time of nanoparticles in the body
	Polymer nano matrix loaded with silver nanoplate clusters and a fluorescent dye	Better biocompatibility and contrast
Multimodal imaging	Quantum dots (QDs)	Good solubility in water, high fluorescence quantum yield, low fluorescence quenching rate and stable chemical properties
	Quantum dots modified with tumor targeting molecules	Specific bioimaging
	Upconversion nanoparticles (UCNPs)	Good photostability, no fluorescence scintillation, deep tissue penetration and small photo damage
	Ag ₂ S QDs	Deeper penetration potential through tissues, higher fidelity of images
	Angiopep-2 (ANG) dual-targeting simultaneously Gd-doped upconversion dual-mode imaging nanoparticles (ANG/PEG- UCNPs)	Enhanced T1-weighted MR contrast of glioblastoma, great potential in T2-weighted MRI, ability to show a clear glioblastoma boundary
	Gold-silica-based SERS nanoparticles	Ability to show clear tumor visualization by three modalities in triple-modality imaging

nanoparticles (IONPs) based on gene therapy to treat patients with glioblastoma via ferroptosis and apoptosis after surgery. By modifying the porous structure of carboxyl-linked IONPs, they co-transferred small interfering RNA (siGPX4, targeting glutathione peroxidase 4) and cisplatin with a high drug loading efficiency. During intracellular degradation, IONPs markedly increased iron (Fe2+ and Fe3+) levels and activated reductive nicotinamide adenine dinucleotide phosphate (NADPH) oxidase (NOX), increasing H₂O₂ levels. The Fenton reaction between Fe²⁺, Fe³⁺, and intracellular H₂O₂ produced reactive oxygen species and initiated ferroptosis, while cisplatin destroyed nuclear and mitochondrial DNA, leading to apoptosis. Simultaneously, si-GPX4 was released, inhibiting GPX4 expression and produced a synergistic effect through mechanisms related to ferroptosis. Therefore, this system achieved an excellent therapeutic effect and low systemic toxicity both in vitro and in vivo (Zhang et al., 2020).

2.5.2 Gene therapy

Gene therapy is a potential method for the treatment of GBM. In this context, nanomaterials are mainly used as carriers for genes designed to treat tumors. The designed genes can be suicide genes (that convert nontoxic prodrugs into cytotoxic drugs), immunoregulatory genes (that stimulate the immune system), or tumor suppressor genes (Dixit and Kumthekar, 2017). In addition, Qiang et al. constructed lipid-polymer hybrid nanoparticles (LPHNs-cRGD) to efficiently and specifically deliver a CRISPR/Cas9 plasmid targeting the temozolomide resistance gene O-6-methylguanine-DNA methyltransferase (MGMT). To facilitate the entry of the genes into the GBM in vivo, they non-invasively and locally gained access into the BBB using focused ultrasound microbubbles. The nanocarrier successfully mediated the transfection of pCas9/MGMT, downregulating MGMT expression and increasing the sensitivity of GBM cells to temozolomide (Yang et al., 2021).

2.5.3 Radiotherapy

Radiotherapy is a common treatment method for malignant tumors. However, radiation damages normal tissues and tumor hypoxia can lead to radiation resistance. There are two solutions to overcome these problems: the first is to use more advanced radiotherapy technology, and the second is to develop a new generation of therapeutic agents able to sensitize tumor cells to ionizing radiations to improve their effect. Nanomaterials can be used as radiosensitizers or their carriers. They have achieved good results as radiosensitizers or carriers after photon and particle radiation (Caban-Toktas et al., 2020; Chung et al., 2020; Kazmi et al., 2020; Ruiz-Garcia et al., 2021). Some nanoparticles, such as gadolinium, gold, hafnium, bismuth, and platinum nanoparticles, can also achieve good results as sensitizers (Lux et al., 2019). The principle is that when photons and particles activate the nanoparticles, a photoelectric effect (Kazmi et al., 2020) amplifies the radiation effect (Lacombe et al., 2017; Kuncic and Lacombe, 2018). Gold nanoparticles became an important radiosensitizer due to their biocompatibility, tunable optical properties, and high stability. Kunoh et al. (2019) compared the radiosensitivity of gold nanoparticle-treated and untreated cells. Their experiments showed that pre-treating cells with gold NPs prevented radioresistance development in cancer cells. Interestingly, the gold NPs did not induce apoptosis but increased the number of abnormal nuclei, causing mitotic cell catastrophe.

2.5.4 Photothermal therapy

Photothermal therapy consists in injecting materials with high photothermal conversion efficiency into the body, making them accumulate around the tumor by targeted recognition technology, then irradiating them with an external light source (generally infrared light) to convert light into heat and kill cancer cells. Based on the above ferroptosis treatment, Yulin *et al.* blended gallic acid with Fe²⁺ to form gallic acid/Fe²⁺ nanoparticles with excellent photothermal conversion ability. Near-infrared light irradiation (808 nm) can drastically improve the Fe²⁺ release efficiency of nanoparticles and induce ferroptosis in tumor cells while releasing a large amount of heat to kill tumor cells (Zhang et al., 2021).

2.5.5 Magnetothermal therapy

Magnetothermal therapy-mediated cancer therapy (MHCT) consists in exposing magnetic nanomaterials to an alternating magnetic field to heat tumor tissue and alter cellular mechanisms. A temperature rise from 37°C to 42°C–45°C can induce tumor cell death by activating specific intracellular and extracellular degradation mechanisms (Krawczyk et al., 2011; Zhang and Calderwood, 2011; Gupta and Sharma, 2019). At 42°C, tumor cells undergo irreversible damage leading to apoptosis, while achieving the same effect in normal cells requires at least 55°C.

However, magnetothermal therapy has not yet become one of the main GBM therapy because some challenges remain. First,

the safety, efficacy, and appropriate dose range of MHCT are unclear, and this needs to be taken into account to determine the magnetic nanomaterial dose. Moreover, the choice of magnetic parameters and the appropriate magnetic field strength are also undetermined. Second, injecting effective drugs into targeted GBM cells through clinically feasible methods remains challenging. Most small-molecule drugs cannot penetrate the BBB, which dramatically hinders the delivery of drugs to tumor sites. In addition, some physical limitations affect hyperthermia performance. These include heat distribution, toxicity, magnetic nanosensors efficiency, and the reduction of hyperthermia performance of magnetic nanoparticles (MNPs) in the cellular environment, that is, once they are internalized by the cell (lysosomal aggregation phenomenon) (Di Corato et al., 2014; Soukup et al., 2015). Besides, the lack of methods to accurately measure local body temperature is another obstacle to MNP treatment evaluation (Dewhirst et al., 1997; Arthur et al., 2005). More importantly, achieving the precise targeting of tumor cells by MNPs is also one of the main challenges for GBM treatment.

According to a recent meta-analysis, less than 1% of injected particles accumulate at the tumor site. Thus, the use of targeted strategies to attach specific targeted moieties to the surface of nanomaterials has also become an important unsolved question. However, in some techniques, only 4% of the targeted portion of the used ligand is recognized by its targeted receptor, which may lead to heterogeneity and poor results (Herda et al., 2017). In addition, converting targeted strategies from basic research to clinical research is ineffective, especially MHCT. A study reported that in mouse xenograft models, the accumulation of antibodies usually varies between 0.5% and 50% of the injection dose per Gram of tumor tissue. Meanwhile, we observed that the accumulation of antibodies per Gram of tumor tissue in human tumors was less than 0.01% of the injection dose (Björnmalm et al., 2017). Gupta and Sharma. (2021) proposed magnetic dots coated with carboxymethyl-stevioside as a magnetic hyperthermia agent for GBM treatment. These magnetic dots showed significant water stability, and their specific absorption rate was 209.25 W/g under an alternating magnetic field of 359 kHz and 188 Oe. They also induced notable antimigration and anti-invasive effects on GBM C6 cells by inhibiting the gene expression of matrix metalloproteinases 2 and 9. The key to solving these problems is controlling the amount of magnetic materials that can reach the tumor microenvironment. For this, doping in an appropriate proportion can improve the magnetism of MNPs (Li et al., 2021). Due to the high specific surface area volume ratio, van der Waals force, and strong dipole-dipole interaction, MNPs tend to agglomerate, resulting in increased particle size and reduced magnetism. The high polydispersity of nanoparticles can also reduce the magnetic heating capacity of MNP systems. Therefore, nanoparticle size crucially affects the magnetic and thermal efficiency of nanosystems. In biomedical applications, MNPs with a small diameter (10-100 nm) and narrow size distribution are preferred to prevent their rapid removal from the systemic circulation by the reticuloendothelial system (Cheng et al., 2021). In addition, surface coatings of nanoparticles [such as inorganic materials (alumina or silica), polymers (dextran, chitosan, polyethylene glycol, or stevioside), fatty acids (oleic acid), and liposomes] prevent aggregation and may contribute to colloidal stability through space and electrostatic repulsion (Karimi et al., 2013; Jamari et al., 2020).

Heat shocks can induce the expression of various heat shock proteins (HSPs), which act as molecular chaperones to protect proteins from thermal denaturation, assist protein folding, and induce heat tolerance in cells (Gong et al., 2012). HSP 27, 70, 73, and 90 are considered the key constitutively overexpressed HSPs in GBMs and play an essential role in cancer cell heat resistance against MHCT (Lee Titsworth et al., 2014). Thus, besides improving effective magnetism the at tumor microenvironment, methods targeting these heat shock proteins can also make tumor cells sensitive to magnetic hyperthermia therapy. Possible strategies include specifically inducing an immune response towards these tumor-specific overexpressed HSPs in GBM cells or using HSP gene inhibitors. The heat-induced antitumor immune response is also a new research direction.

2.5.6 Immunotherapy

To date, the standard high-grade GBMs therapy concerns multidisciplinary approaches, containing maximum surgical resection, radiotherapy, and chemotherapy. But the complete resection is nearly impossible because of the invasiveness of GBM, and the recurrence of tumor is practically inevitable even in patients undergoing multimodal therapy. Moreover, these recurrent tumors are often resistant to chemotherapy and radiotherapy (Leiva-Salinas et al., 2017). Thus, it is necessary to develop new therapeutic approaches against GBM. Immunotherapy consists in stimulating the patient's immune system to make it identify and attack malignant tumors through continuous anti-tumor immunity. However, many obstacles hinder the application of immunotherapy in the clinical practice. The first is the complexity of tumors. Tumors have various immune escape mechanisms. Second, the immune environment of brain is distinct from that in other organs, which is unable to produce an immune response against tumors (Zanganeh et al., 2019). Due to the BBB, the transport of immune effectors from the blood to the brain is limited. Moreover, although activated circulating T lymphocytes are present in the central nervous system, there are few naive T cells (Su et al., 2017).

Another challenge limiting immunotherapy against GBM is that various mechanisms promote immunosuppression inside and around the tumor. GBM is classified as a cold immune tumor, and its microenvironment represents an immune desert with little to no immune effector cell infiltration. Key factors of the glioblastoma-mediated immune cold microenvironment contain the abundance of CD4⁺CD25+FOXP3+ regulatory T cells (Tregs) and myeloid cells, as well as immunosuppressive cytokines and secretory factors produced by tumor cells like transforming growth factor- β , interleukin 6 and interleukin 10 (Reardon et al., 2017). GBM grading is related to Treg infiltration into tumors. In malignant GBM, tumor-resident Tregs express high levels of PD-1 (Lowther et al., 2016), an essential inhibitory receptor expressed in activated T cells which is significant in the immune response (Sharpe and Pauken, 2018). Thus, although there is a small amount of immune cell infiltration, immune cells are often in a low response state due to immunosuppressive signals (Woroniecka et al., 2018).

A primary mediator of immunosuppression in GBM patients is tissue hypoxia, which activates signal transducers and activators of transcription 3 (STAT3) and an immunosuppressive signaling pathway which promotes the production of hypoxia-inducible factor-1-alpha (HIF1A). However, it then induces Treg activation and vascular endothelial growth factor (VEGF) synthesis (Almiron Bonnin et al., 2018).

In the face of the many challenges in immunotherapy, nanomaterials can be used as a breakthrough point. Nanomaterials can deliver drugs to tumors and induce cytotoxic anti-tumor T cell responses. Cancer vaccines with designed better efficacy can be by combining immunomodulators and antigens, direct targeting and T cell functionalization, nucleic acid delivery, adjuvants, immune checkpoint inhibitors, and inhibitory tumor microenvironment regulation (Riley et al., 2019). In a current research, poly (lactic-coglycolic acid) nanoparticles modified by Angiopep-2 and IP10-EGFRvIIIscFv fusion proteins crossed the BBB and amassed in brain tissue. After binding to cytotoxic T lymphocytes, the nanoparticles notably increased the immune response and antitumor activity in a GBM model (Wang et al., 2018).

To re-adjust the inhibited GBM immune microenvironment, Tong Fei *et al.* designed a nanocomposite drug based on a polyglycerol functionalized doxorubicin-containing nanodiamond. The nanocomposite regulated the immunosuppressive microenvironment of GBM by activating autophagy, thereby stimulating the immune response. This rebuilding promoted the anti-glioblastoma immune response and strengthened the activation of dendritic cells (Li et al., 2019).

The tumor vascular laminin-411 ($\alpha 4\beta 1\gamma 1$) is associated with the high expression of tumor stem cell markers (Notch, CD133, Nestin, c-Myc) and a shorter survival time for GBM patients. Tao *et al.* designed a nano-bioconjugate which is able to cross the BBB to inhibit laminin-411. This nanobioconjugate targeting the tumor microenvironment prolonged animal survival and inhibited cancer stem cell markers in mice carrying intracranial GBM (Sun et al., 2019).

The low accumulation of antigens in antigen-presenting cells is another obstacle to effective immunotherapy against brain tumors. This is related to the low activation of antigen-presenting cells in GBM. Two approaches could solve this problem: enhancing the antigen-loading capacity of nanovaccines or



adding substances that can sensitize antigen-presenting cells to nanomedicines. A hybrid "cluster bomb" nanovaccine, based on zinc oxide and triblock-copolymer nanoparticles, stimulated cellular and humoral immunity and increased the survival time of tumor-bearing mice (Shen et al., 2019). Another study combined exosomes derived from GBM with α -galactosylceramide (natural killer T-cell activator). Used subcutaneously in glioblastoma-bearing rats, it increased interferon γ and tumor necrosis factor α production and promoted the immune response (Liu H. et al., 2017).

2.5.7 Summary

Because of the heterogeneity, the complexity of the tumor microenvironment, the multiple immune escape mechanisms, and the weak sensitivity of the brain immune system, handling GBM from a single dimension is limited. GBM treatments need to be multimodal. However, combining different methods for killing glioma cells is insufficient. Comprehensive glioblastoma treatments should also target the tumor microenvironment, and stimulate immune cells of the brain (Figure 2).

3 Summary and outlook

This article reviews the definition, classification, diagnosis, and therapeutic applications of nanomaterial-based GBM treatments. Currently, the clinical application of nanoparticles and the complexity of GBM itself still face numerous challenges. Designing nanomaterials more suitable for clinical applications require optimizing nanomaterials by understanding the transport regulation mechanism of the BBB, the composition of the GBM tumor microenvironment and its influence on the BBB. Carefully studying the mechanisms of action of nanomaterials on the brain, discovering new properties of nanomaterials, improving their synthesis, and exploring new and promising drug delivery systems are crucial for developing clinical applications of nanomaterials in the diagnosis and treatment of glioblastoma. Most importantly, since singledimension diagnostic and treatment methods are limited, future studies should focus on multi-dimensional nanomaterials. With further research, the clinical nanodiagnosis and treatment system for GBM is expected to improved.

Author contributions

All authors contributed to the design and writing of the manuscript. XL, XN, and WW proposed the ideas and drafted the outlines. SY, LC, and HX performed the literature search and completed the manuscript. SL and JJ helped revise the manuscript and provided support in need.

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Conflict of interest

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Phytofabrication and characterization of *Alchornea cordifolia* silver nanoparticles and evaluation of antiplasmodial, hemocompatibility and larvicidal potential

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Purpose: The recent emergence of *Plasmodium falciparum (Pf)* parasites resistant to current artemisinin-based combination therapies in Africa justifies the need to develop new strategies for successful malaria control. We synthesized, characterized and evaluated medical applications of optimized silver nanoparticles using *Alchornea cordifolia* (AC-AgNPs), a plant largely used in African and Asian traditional medicine.

Methods: Fresh leaves of *A. cordifolia* were used to prepare aqueous crude extract, which was mixed with silver nitrate for AC-AgNPs synthesis and optimization. The optimized AC-AgNPs were characterized using several techniques including ultraviolet-visible spectrophotometry (UV-Vis), scanning/ transmission electron microscopy (SEM/TEM), powder X-ray diffraction (PXRD), selected area electron diffraction (SAED), energy dispersive X-ray spectroscopy (EDX), Fourier transformed infrared spectroscopy (FTIR), dynamic light scattering (DLS) and Zeta potential. Thereafter, AC-AgNPs were evaluated for their hemocompatibility and antiplasmodial activity against *Pf* malaria strains 3D7 and RKL9. Finally, lethal activity of AC-AgNPs was assessed against mosquito larvae of *Anopheles stephensi, Culex quinquefasciatus* and *Aedes aegypti* which are vectors of neglected diseases such as dengue, filariasis and chikungunya.

Results: The AC-AgNPs were mostly spheroidal, polycrystalline (84.13%), stable and polydispersed with size of 11.77 \pm 5.57 nm. FTIR revealed the presence of

several peaks corresponding to functional chemical groups characteristics of alkanoids, terpenoids, flavonoids, phenols, steroids, anthraquonones and saponins. The AC-AgNPs had a high antiplasmodial activity, with IC₅₀ of 8.05 µg/mL and 10.31 µg/mL against 3D7 and RKL9 *Plasmodium falciparum* strains. Likewise, high larvicidal activity of AC-AgNPs was found after 24 h- and 48 h-exposure: $LC_{50} = 18.41 \mu g/mL$ and 8.97 µg/mL (*Culex quinquefasciatus*), $LC_{50} = 16.71 \mu g/mL$ and 7.52 µg/mL (*Aedes aegypti*) and $LC_{50} = 10.67 \mu g/mL$ and 5.85 µg/mL (*Anopheles stephensi*). The AC-AgNPs were highly hemocompatible (HC₅₀ > 500 µg/mL).

Conclusion: In worrying context of resistance of parasite and mosquitoes, green nanotechnologies using plants could be a cutting-edge alternative for drug/ insecticide discovery and development.

KEYWORDS

Alchornea cordifolia, silver nanoparticles, green synthesis, characterization, hemocompatibility, biocidal activities, Plasmodium falciparum, culicidae mosquitoes

1 Introduction

Nanotechnology has profoundly changed several aspects of human life through technological and health advances made in sectors such as new technologies, energy, cosmetics and health. This term encompasses a set of activities from development research to evaluation of materials sized 1–100 nm—also known as nanomaterials (Subedi, 2013; Bayda et al., 2020). In developed countries such as the United States of America, Japan and China, nanotechnologies are greatly funded, studied and evaluated for their ability to solve diverse problems (Dong et al., 2016; Qiu, 2016). In developing countries, studies are more focused on green nanotechnologies which rely on the development of nanomaterials using living organisms such as plants and microorganisms (Kojom Foko et al., 2019; Kojom Foko et al., 2021).

Green nanotechnologies are very attractive as these are cheaper to implement, safer and eco-friendly as compared to their chemical and physical counterparts (Tran et al., 2013; Gahlawat and Choudhury, 2019). Indeed, chemical and physical methods are time-consuming, costly and request reagents which are harmful to humans and environment (Shakeel et al., 2016). By blending living organisms (e.g., fungi, viruses, bacteria, alga, plants) or derived products with metal source, green metallic nanoparticles (MNPs) are synthesized and then can be tested for different biological and non-biological activities (Honary et al., 2013; Kojom Foko et al., 2019; Bayda et al., 2020; Chugh et al., 2021; Araújo et al., 2022). Also, the utilization of plants is more advantageous than with microorganisms due to increased risk of biohazard and cost to isolate, purify and maintain microbial cultures (Kalishwaralal et al., 2010; Shakeel et al., 2016).

Regarding biological activities, many studies reviewed biocidal potential of green MNPs against non-communicable diseases (e.g., diabetes, cancer), oxidative stress, diverse pathogens (e.g., bacteria, viruses), and disease vectors (e.g., mosquitoes, ticks) (Benelli et al., 2017; Patil and Chandrasekaran, 2020; Araújo et al., 2022). Roughly, plantbased MNPs show a high biocidal potential, and thus were proposed as new avenues for control of infectious diseases, especially mosquito-borne diseases (e.g., malaria, dengue and chikungunya) for which current control methods are jeopardized due to i) their toxicity to humans and environment, and ii) emergence and spread of drug-resistant parasites and insecticide-resistant mosquitoes (Kojom Foko et al., 2019; Kojom Foko et al., 2021). Thus, synthesis of plant-based MNPs could be interesting to develop new drugs and insecticides to control and eliminate mosquito-borne diseases. Malaria is the predominant vector-borne disease globally with an estimated 247 million cases and 619,000 deaths in 2021 (World Health Organization, 2022). Africa bears the bulk of this global malaria burden, with children under 5 years of age and pregnant women being most vulnerable groups (Dongang Nana et al., 2022; World Health Organization, 2022). Resistance of pathogens and mosquito vectors is a great threat to malaria control and elimination efforts (Arya et al., 2021). Recent studies pointed out independent emergence of malaria parasites resistant to current most effective antimalarial drugs (i.e., artemisinin-based combination therapies - ACTs) in two African countries (Rwanda and Uganda) (Uwimana et al., 2021, 2020; Balikagala et al., 2021).

There is paucity of data on biological activities of green MNPs in Cameroon where vector-borne diseases such as malaria are causes of concern (Lehman et al., 2018; Antonio-Nkondjio et al., 2019; Mbohou et al., 2019). In the present study, silver NPs were synthesized using leaves of Alchornea cordifolia (AC-AgNPs), optimized, characterized and evaluated for hemocompatibility and lethal activity against Plasmodium falciparum-Pf (the main and deadliest human malaria species) (Kojom Foko et al., 2021; Kojom Foko et al., 2022b; World Health Organization, 2022), and three mosquito species, i.e., Anopheles stephensi, Culex quinquefasciatus and Aedes aegypti, involved in human transmission of parasites and viruses (dengue, Zika, malaria and lymphatic filariasis) (Benelli et al., 2017; Wang et al., 2017; Patil and Chandrasekaran, 2020). Alchornea cordifolia Schumach. and Thonn.) Müll. Arg. (Euphorbiaceae) is largely distributed in sub-Saharan African countries (e.g., Cameroon, Ghana, Nigeria) where its leaves and root bark are traditionally used by populations for nutritional purposes and treating several infectious and inflammatory ailments such as rheumatism, pain and arthritis (Ngaha Njila et al., 2016; Cesar et al., 2017).



FIGURE 1

Flowchart depicting the study design. AC-AgNPs, *Alchornea cordifolia* silver nanoparticles; GC-MS, Gas chromatography—Mass spectrometry; DLS, Dynamic light scattering; EDX, Energy dispersive X-ray spectroscopy; FTIR, Fourier transformed infrared spectroscopy; PXRD, Powder X-ray diffraction; SAED, Selected area electron diffraction; SEM, Scanning electron microscopy; TEM, Transmission electron microscopy; UV-Vis, Ultraviolet—Visible spectrophotometry.

2 Materials and methods

2.1 Study design

This was an experimental study aimed at determining antiplasmodial, hemocompatibility and larvicidal potential of biosynthesized silver NPs using *A. cordifolia* leaves. The plant was harvested and authenticated taxonomically. Crude extract of *A. cordifolia* leaves (AC-CE) was screened for phytochemical composition and used for AgNPs synthesis. The optimization of AC-AgNPs was made, and the optimized AC-AgNPs were characterized and tested for antiplasmodial, hemocompatibility and larvicidal potential (Figure 1). The study was approved by ethical committee of the National Institute of Malaria Research (NIMR), India (N°PHB/NIMR/EC/2020/55).

2.2 Collection and authentication of plant material

Healthy and fresh leaves of *A. cordifolia* (AC) were collected at Faculty of Sciences (FS), main campus, University of Douala (UD), Littoral Region, Cameroon (Figure 2). Malaria is highly prevalent in Cameroon, and *P. falciparum* is the main malaria species. Other species including *Plasmodium vivax*, *Plasmodium ovale spp* have also been reported across the country (Kojom Foko et al., 2021; Kojom Foko et al., 2022a). The taxonomic authentication was done by Dr Tchiengue Barthelemy at Cameroon National Herbarium, Yaounde, in comparison with voucher specimen number 9657/SRF/ Cam previously deposited.

2.3 Preparation of *A. cordifolia* aqueous extract

About 500 g of fresh A. cordifolia leaves were washed with running tap water and distilled water to remove dust and surface contaminant, and thereafter air-dried for 2 weeks at room temperature. The dried material was introduced in an electric grinder to obtain a fine powder. Ten grams of powder was taken in a conical flask containing 100 mL of distilled water, heated at 80°C for 10 min in a water bath under static conditions (Eva'ane Meva et al., 2016). The mixture was allowed to cool at room temperature, and then filtered using a Whatman paper n°1 to remove particulate matter. The filtrate obtained (crude extract, AC-CE) was used to perform phytochemical screening and AC-AgNPs biosynthesis. The AC-CE was not used more than a week following its preparation in order to avoid gradual loss of viability due to long storages (Eya'ane Meva et al., 2016). The AC-CE was lyophilized and stored for biological assays. The yield of extraction of AC-CE was 41% (w/v).

2.4 Phytochemical screening of *A. cordifolia* aqueous extract

The AC-CE was subjected to gas chromatography-mass spectrometry (GC-MS) analysis to identify the composition and percentage abundance of phytochemical constituents. The GC-MS was carried out on a Perkin Elmer Turbo Mass Spectrophotometer (Norwalk, CTO6859, NY, United States) which includes a Perkin Elmer Auto sampler XLGC. The column used was a Perkin Elmer Elite-5 capillary column measuring 30 m \times 0.25 mm with a film thickness of 0.25 mm composed of 95% dimethyl polysiloxane. The carrier gas used was helium at a flow rate of 1.21 mL/min 1 µL sample injection volume was utilized. The inlet temperature was maintained at 260°C. Oven temperature was programmed initially at 100°C for 2 min, and then programmed to increase to 290°C at a flow rate of 10°C/min (Supplementary Figure S1). The total run time was 39.98 min. The Mass Spectrometry transfer line was maintained at a temperature of 200°C. The source temperature was maintained at 220°C. The GC-MS was analyzed using electron impact ionization at 70 eV. Full scan mode was used to detect analytes. Data were evaluated using total ion count for compound identification and quantification. Measurement of peak areas and data processing were carried out by Turbo-Mass-OCPTVS-Demo SPL software, and spectrums of the components were compared with database of spectrum of known components stored in the GC-MS library.



2.5 Phytofabrication and optimization studies of AC-AgNPs

AC-AgNPs were synthesized by blending AgNO₃ aqueous solution with freshly prepared AC-AE and incubated in dark until color change. The determination of optimal conditions for AC-AgNPs biosynthesis was performed by recording UV-Visible spectra of reaction mixtures after varying four parameters, namely, incubation temperature (35° C– 85° C), incubation time (10 min–5 h), AgNO₃ concentration (0.5-5 mM), and AgNO₃/AC-CE volume ratio ($10-100 \mu$ L) as described earlier (Hawadak et al., 2022). Thereafter, the optimized reaction mixture was centrifuged at 15,000 rpm for 10 min, the pellet was washed twice with distilled water and once with 95% ethanol, filtered using sterile syringe filter (MICRO-POR^{*}, 0.22μ m), and then lyophilized for further AC-AgNPs characterization and biological assays.

2.6 Characterization of AC-AgNPs

The characteristics of green synthesized AC-AgNPs (i.e., surface plasmon resonance-SPR, size, shape, aggregation, functional chemical groups and crystallinity) were determined using several techniques (Figure 1). The formation of AC-AgNPs was monitored by visual inspection of the solution and then followed by UV–Vis spectrum measurement using a double beam spectrophotometer (Model n.o., BRI-2700, BR BIOCHEM Life Sciences Pvt., Ltd., India) operating at 1 nm resolution. Milli Q ultrapure water was used as blank. The selected area electron diffraction (SAED) and powder X-ray diffraction (XRD) were used to determine the physical nature of the AC-AgNPs. The PXRD was made at 45 kv voltage, 40 mA current, 2θ range of 10–80 and speed of 2°/minute (PANanalytical,

Xpert Pro model). The PXRD patterns of optimized AC-AgNPs were compared to Joint Committee on Powder Diffraction Standards files (JCPDS 65-2871 and 31-1238). The size, shape and aggregation patterns of AC-AgNPs were determined using scanning electron microscopy-SEM coupled with EDX (Bruker AXS Microanalysis GmbH Berlin, Germany) and transmission electron microscopy-TEM coupled with SAED (TECNAI TF20, Fei, Electron Optics, Oregon, United States) operating at a potential of 20 kv and 200 kv, respectively. The size of NPs was calculated using the Scherrer equation: $D = \frac{K\lambda}{\beta \cos \theta}$, where D is diameter (nm) of the crystallite (i.e., NPs in this regard), K is the Scherrer constant (range values = 0.68-2.08) depending on shape of nanoparticles (e.g., K = 0.94 for spherical NPs); λ is the X-ray wavelength (in our study PXRD analysis was performed at wavelength for copper, $CuK_{\alpha} = 1.5406 \text{ Å}$), β is the line broadening at full width at half maximum (FWHM) which is expressed in radians, and θ is the Bragg's angle of PXRD-related peaks which is expressed in degrees (Muniz et al., 2016). The atomic composition of the NPs was determined using energy dispersive X-ray (EDX). Fourier transformed infrared spectroscopy (FTIR) was used to determine functional chemical groups capped on the AC-AgNPs surface through potassium bromide method. Sample was grinded with KBr in an infrared path and the spectrum was recorded in the range 400-4000 cm⁻¹ using a FTIR spectrophotometer (Perkin Elmer, Frontier Model). Zeta potential and dynamic light scattering (DLS) were performed to evaluate NPs stability and size distribution using particle size analyzer (Zetasizer nano ZS, Malvern Instruments Ltd., U.K.). In practice, zeta potential of ±30 mV is considered as a good indicator of the stability of colloidal suspensions such as NPs while values outside the range indicate phenomena such as flocculation, aggregation and sedimentation (Kojom Foko et al., 2019).

TABLE 1 Phytochemical screening of the AC-AE using GC-MS analysis.

Peak	Retention time	Area (%)	Name of the compounds
1	6.76	2.58	4-Methylmannitol
2	9.22	0.29	Dodecanoic acid, methyl ester
3	9.52	0.48	2(4H)-Benzofuranone, 5,6,7,7a-tetrahydro-4,4,7a-trimethyl-, (R)-
4	10.04	0.57	1-Hexadecene
5	10.99	0.41	8-Pentadecanone
6	11.59	0.36	1,1,4,7-Tetramethyldecahydro-1H-cyclopropa[e]azulene-4,7-diol
7	12.08	1.42	6-Hydroxy-4,4,7a-trimethyl-5,6,7,7a-tetrahydrobenzofuran-2(4H)-one
8	12.31	0.58	1-Nonadecene
9	12.37	0.51	2(4H)-Benzofuranone, 5,6,7,7a-Tetrahydro-6-hydroxy-4,4,7a-trimethyl-
10	12.51	1.00	(S,E)-4-Hydroxy-3,5,5-trimethyl-4-(3-oxobut-1-en-1-yl)cyclohex-2-enone
11	12.77	1.28	Neophytadiene
12	12.84	1.08	2-Pentadecanone, 6,10,14-trimethyl-
13	13.02	0.16	2-hexadecen-1-ol, 3,7,11,15-tetramethyl-, [R-[R*,R*-(E)]]-
14	13.08	0.32	1,2-Benzenedicarboxylic acid, bis(2-methylpropyl) ester
15	13.58	0.60	1,2-Benzenedicarboxylic acid, bis(2-methylpropyl) ester
16	13.69	1.86	Hexadecanoic acid, methyl ester
17	14.22	2.36	n-Hexadecanoic acid
18	14.36	0.22	1-Octadecene
19	14.84	1.99	Hexadecanoic Acid, trimethylsilyl ester
20	15.33	0.43	9,12-Octadecadienoic acid (Z,Z)-, methyl ester
21	15.39	3.64	9,12,15-Octadecatrienoic acid, methyl ester, (Z,Z,Z)-
22	15.51	25.17	2-Hexadecen-1-Ol, 3,7,11,15-Tetramethyl-, [R-[R*,R*-(E)]]-
23	15.62	0.56	Methyl stearate
24	15.99	1.40	Phytol, TMS derivative
25	16.40	0.54	Phytol, acetate
26	17.63	0.21	4,8,12,16-Tetramethylheptadecan-4-olide
27	19.09	1.41	Bis(2-ethylhexyl) phthalate
28	20.31	0.36	Tetracontane
29	21.14	1.70	Squalene
30	21.57	0.40	.alphaTocospiro B
31	21.78	0.52	Hexatriacontane
32	22.52	1.11	9,12-Octadecadienoic Acid (Z,Z)-, 2,2-Dimethyl-1,3-Dioxolan-4-Ylmethyl Ester
33	24.12	0.81	Vitamin E
34	24.84	3.32	Ethanone, 1-(2,3,4,7,8,8a-hexahydro-3,6,8,8-tetramethyl-1H-3a,7-methanoazulen-5-yl)-
35	25.93	3.65	STIGMASTA-5,22-DIEN-3-OL
36	26.87	4.69	.gammaSitosterol
37	27.32	4.03	9,19-Cyclolanostane-3,7.betadiol, diacetate (20R,14.beta.)
38	28.25	0.99	9,19-Cyclolanostan-3-ol, 24-methylene-, (3.beta.)-

(Continued on following page)

Peak	Retention time	Area (%)	Name of the compounds
39	31.55	14.53	Phytyl tetradecanoate
40	35.11	3.40	Methanesulfonic Acid 2-(3-Hydroxy-4,4,10,13,14-Pentamethyl-2,3,4,5,6,7,10,11,12,13,14,15,16,17-Tetradecahydro-1h-Cyclopenta[A]Phenan
41	37.27	3.92	1-Eicosanol
42	37.77	5.12	9,10,12,13-Tetrabromooctadecanoic acid
	Total	100.00	

TABLE 1 (Continued) Phytochemical screening of the AC-AE using GC-MS analysis.

2.7 Assessment of antiplasmodial potential of AC-AgNPs

Chloroquine-sensitive 3D7 and chloroquine-resistant RKL9 of *Pf* strains were used for antiplasmodial assays for AC-AgNPs, AC-CE, and chloroquine (CQ). The *Pf* culture was maintained using standard protocols (Trager and Jensen, 1976; Schuster, 2002). Briefly, parasite cultures were maintained in fresh AB positive human erythrocytes suspended at 5% hematocrit in RPMI-1640 culture medium supplemented with L-glutamine and HEPES buffer (0.2% sodium bicarbonate, 0.4% albumax, 50 µg/L hypoxanthine, 200 U/mL penicillin and 200 µg/L streptomycin) and incubated at 37°C under a gas mixture of 1% O₂, 5% CO₂ and 94% N₂. Culture of infected erythrocytes were transferred daily into fresh complete culture medium and checked microscopically for parasite growth.

The in vitro evaluation of antiplasmodial activity was performed using culture-adapted Pf strains: i) 3D7, sensitive to CQ, artemisinin and its derivatives and ii) RKL9, resistant to CQ. Antimalarial drug screening was done based on SYBR green I-based fluorescence assay as described previously (Smilkstein et al., 2004). Parasite culture (0.5%-0.8%) was synchronized at ring stage with 5% sorbitol. A volume of 100 µL of complete medium were introduced into each well of 96-well microplate, then dilutions were performed for AC-AgNPs and AC-CE (4, 8, 16, 31.25, 62.5, 125, 250 and 500 µg/mL) and CQ (4, 6.25, 12.5, 25, 50, 100, and 200 µg/mL) were added. Ten microliters (10 µL) of synchronized blood were thereafter added in each well, mixed and kept in an incubator at 37°C for 48 h in 96-well flat bottom tissue culture-grade plates under reduced O2 atmosphere. Each experiment was replicated thrice. CQ was used as standard drug, while complete medium was considered as negative control. After 48 h-incubation, 100 µL of SYBR Green I in lysis buffer (0.2 µL of the fluorochrome/mL of buffer) was added into each well, mixed gently twice, and the plate was then covered with foil and incubated in a dark chamber for 1 h at room temperature. The buffer lysis consisted of Triton X-100 (0.08% v/ v), Tris (20 mM), EDTA (5 mM), and saponin (0.008% wt/v). The fluorescence counts were read using an ELISA reader (Synergy HTX 1708152, Agilent BioTek, Santa Clara, California, United States) with excitation and emission wavelength bands centered at 485 and 530 nm.

2.8 Validation of antiplasmodial assay

The SYBR Green based antiplasmodial assay was validated by inspecting microscopic slides of parasite cultures treated with negative control, standard drug, AC-CE and AC-AgNPs (Kaushik et al., 2015; Hawadak et al., 2022). After 48 h-incubation, thick and thin blood films were made, air-dried and stained with 10% Giemsa stain for 20 min. The number of schizonts with \geq 2 nuclei out of 200 asexual parasites was noted. Also, fluorescence counts of untreated and treated *Pf* cultures were compared to detect any quenching effect-related measurement artefacts which may due to chemical compounds of AC-AgNPs and AC-CE (Kaushik et al., 2015).

2.9 Hemocompatibility investigation

The method described by Wang and others was used to evaluate hemocompatibility of biosynthesized AC-AgNPs (Wang et al., 2010). Human red blood cells (RBCs) were obtained from the ICMR-NIMR malaria parasite bank, washed with incomplete media, and diluted with phosphate-buffered saline (PBS) to obtain a suspension (Hematocrit = 1%). Different concentrations (2, 4, 8, 16, 30, 62.5, 125, 250 and 500 µg/mL) of AC-AgNPs and AC-CE were incubated with RBCs in Eppendorf tubes (20 µL of each concentration in 180 µL blood) at 37°C for 30 min and 24 h at pH of 7.40. The reaction was stopped by placing tubes at 4°C for 15 min. The mixtures were then centrifuged at 3,000 g for 4 min, and 100 μ L of supernatant was loaded into a 96-well plate to measure the released hemoglobin at 540 nm (SPECTROstar^{Nano}, BMG LABTECH GmbH, Ortenberg Germany). Saponin was used as positive control, inducing 100% hemolysis, while PBS was considered as negative control. The experiment was performed in triplicate. RBCs hemolysis at each concentration after 30 min and 24 h was calculated as follows:

% hemolysis =
$$\frac{(As - ANC)}{(APC - ANC)} \times 100$$

where A_S , A_{NC} and A_{PC} are the absorbance of the sample, negative control (PBS) and positive control (saponin).


AC-AgNPs solution and UV-Vis findings. Color of AgNO₃, AC-AgNPs and AC-CE solutions (A). The color change indicates Ag⁺ reduction to elemental nanosilver. UV-visible spectrum of optimized AC-AgNPs for incubation temperature (B), incubation time (C), AgNO₃ concentration (D), and volume of AC-CE (E). AC-AgNPs, *Alchornea cordifolia* silver nanoparticles; AC-CE, *Alchornea cordifolia* crude extract; AgNO₃, Silver nitrate; UV-Vis, Ultraviolet—Visible spectrophotometry.

2.10 Mosquito rearing

The eggs of An. stephensi, Cx. quinquefasciatus and Ae. aegypti were obtained from NIMR Insectarium, New Delhi, India. The characteristics of mosquitoes used are as follows: An. stephensi-laboratory strain collected from Sonepat, Haryana, India (established in 1996; black and brown, malathion-deltamethrin-DDT-resistant susceptible strain). and Cx. quinquefasciatus-laboratory strain collected from Sonepat, Haryana, India (established in 1999; selected for permethrin resistance and is resistant to DDT, malathion and deltamethrin), and Ae. aegypti-laboratory strain collected from Delhi, India (established in 2006; DDT-malathion-deltamethrin strain). Adult Ae. aegypti were derived from batches of 100 eggs in $18 \text{ cm} \times 13 \text{ cm} \times 4 \text{ cm}$ trays containing 500 mL of boiled and cooled water in a laboratory maintained at 25°C-29°C temperature and 65%-70% Relative humidity; 12:12 h Light/Dark photoperiod. Eggs were fed daily with TetraBits fish food (Tetra GmbH, Herrenteich, Germany), and late 3rd and 4th instar larvae were used for larval bioassays.

2.11 Larvicidal bioassays

The protocol described by the World health Organization (WHO) was used for this experiment (WHO, 2016). Late 3^{rd} and 4^{th} instar larvae were exposed to the AC-AgNPs with different concentrations (0–50 µg/ mL). Each concentration was tested in triplicate comprising of 25 larvae placed into plastic bowls (8 cm diameter, 300 mL capacity) containing distilled water. The larval mortality was monitored after 24 h, 48 h and 72 h post-treatment periods, and the lethal concentrations to cause 50%/90% mortality in treated larvae (LC₅₀/LC₉₀) and percentage mortality after post-treatment periods were calculated as described previously in the WHO procedures (WHO, 2016). Distilled water was used as control. All experiments were performed under laboratory conditions as described above.

2.12 Statistical analysis

Data was keyed into an Excel spreadsheet (Microsoft Office, United States) and then exported to statistical package for social sciences v16 (SPSS, IBM, Inc., Chicago, United States), and GraphPad v5.03 (GraphPad PRISM, Inc., San Diego, California, United States) for statistical analysis. Using GraphPad software v8.03 (GraphPad PRISM, Inc., San Diego, CA, United States), fluorescence counts of antiplasmodial assay were used to plot graph of percent inhibition of Pf parasite growth against concentrations of AC-AgNPs, AC-CE, and CQ to determine 50% inhibition concentration (IC₅₀). The dose/time mortality response data of larvicidal assays was analyzed using log-probit regression model to determine LC_{50} and LC_{90} with their confidence interval at 95% (95% CI). The Abbott's formula was used to correct mortality rate if comprised between 5% and 20% in the negative control group (Sun and Shepard, 1947). Experiments were considered invalid when mortality rate in negative control group was >20%.



SEM and TEM analysis of AC-AgNPs. SEM images at 5.00 KX (A) and 50.00 KX (B) of AC-AgNPs. Micrographs of the AC-AgNPs using TEM at 20 nm (C) and 10 nm (D), and size distribution of the nanocrystallites (E). AC-AgNPs, *Alchornea cordifolia* silver nanoparticles; SEM, Scanning electron microscopy; TEM, Transmission electron microscopy.

Regarding hemocompatibility assay, the amount of NPs required to lyse 50% of RBCs (hemolysis concentration, HC_{50}) was determined. Quantitative and qualitative variables were presented as mean \pm standard deviation (SD) and percentages, respectively. One-way analysis (ANOVA), McNemar's and Pearson's independence chi square tests were used to make comparisons. The level of statistical significance was set at *p*-value <0.05.

3 Results

3.1 GC-MS analysis

GC-MS chromatogram of AC-CE revealed several peaks which represent different compounds as shown in Supplementary Figure S1. A total of 42 compounds were identified in AC-CE after comparing the peaks with database of spectrum of known components stored in the GC-MS library (Table 1). Two compounds were predominantly represented, namely, 2hexadecen-1-ol, 3,7,11,15-tetramethyl-, [R-[R*,R*-(E)]]- and phytyl tetradecanoate, with proportions of 25.14% and 14.53%, respectively (Supplementary Figure S2; Table 1).

3.2 UV-Vis spectroscopy and AC-AgNPs optimization

The synthesis of AC-AgNPs was noted after 2 minutes following the incubation of plant extract and AgNO₃ solution as a dark brown color was observed (Figure 3A). The UV-Vis spectrum analysis revealed a SPR at 445 nm wavelength (Figures 3B–E). The SPR did not change with the variation of four parameters used to optimize AC-AgNPs synthesis (AgNO₃ concentration, incubation time, incubation temperature and volume of plant extract). In contrast, the amplitude of UV-Vis curves gradually increased with increasing values of each parameter (Figures 3B–E). Thus, the optimization of AC-AgNPs was achieved for the following parameters: $100 \,\mu$ L of fresh plant extract was mixed with 900 μ L of AgNO₃ (5 mM), and then incubated at 85°C for 5 h under static conditions.

3.3 Electron microscopy analysis of green AC-AgNPs

Analysis of SEM and TEM micrographs of AC-AgNPs is depicted in Figure 4. Based on SEM, agglomeration of AC-AgNPs was observed (Figures 4A, B). TEM images of silver colloidal solution exhibited that AC-AgNPs were polydispersed, predominantly spheroidal with various sizes (Figures 4C, D). The size distribution when a section of these NPs is considered is presented in Figure 4E. Following the digitization phase of various images, size distribution using ImageJ software was found to be within 5–25 nm range. The distribution of AC-AgNPs size was large, with mean size \pm SD of 10.89 \pm 5.67 nm.

3.4 PXRD analysis, SAED patterns, and composition of AC-AgNPs

The PXRD patterns outline that AC-AgNPs are face-centered cubic. The intense and narrow diffraction peaks revealed the formation of pure crystals of silver and silver chloride. The nanosilver crystal peaks obtained at 20 values of 38.07°, 46.20°, 64.33° and 77.40° which correspond to the (111), (200), (220) and (311) planes of the face-centered cubic (fcc) structures, respectively (JCPDS file 65-2871). Additional peaks corresponding to silver chloride nanocrystallites were observed at 20 values of 27.8°, 32.2°, 54.8°, 57.4° and 67.4° indexed to (111), (200), (311), (222) and (400) planes, respectively (JCPDS file 31-1238). SAED suggests that the NPs are polycrystalline with diffraction rings associated due to their stacking each other due to their magnetite phase (Figures 5A, B). The crystallinity percentage of AC-AgNPs was 84.13%. Using data from PXRD, the size of silver nanocrystals and silver chloride nanocrystals based on the Scherrer formula was 13.47 \pm 5.81 nm and 10.42 \pm 3.34 nm, respectively (Table 2). The overall mean size of AC-AgNPs was 11.77 ± 5.57 nm.

The EDX profile of AC-AgNPs showed a strong signal due to silver atom (Ag) which was involved in AC-AgNPs at a percentage of 58.31%. Other signals due to chlorine (Cl), cadmium (Cd), carbon (C) and oxygen (O) were also observed at 5.56%, 0.36%, 9.31% and 26.46%, respectively (Figure 5C). The identity of functional chemical groups at the interface of AC-AgNPs were determined using FTIR which revealed strong signals at $3,416 \text{ cm}^{-1}$, 1630 cm^{-1} , and 1023 cm^{-1} which are characteristics of alcohols (O-H stretch), alkenes (C=O stretch) and alkyl and Aryl Halides (C-F stretch), respectively. Smaller signals corresponding to alkanes (C-H stretch) at 2927 cm⁻¹, alkanes/aldehydes/alkenes (C-H stretch, C-O stretch) at 2857/1739 cm⁻¹, nitriles (C≡N stretch) at 2373/ 2323 cm⁻¹, aromatic compounds (C=C stretch) at 1458 cm⁻¹, and nitro compounds (NO₂ stretch) at 1377 cm⁻¹ were also seen (Figure 5D; Table 3).

3.5 Zeta potential and DLS

The stability of AC-AgNPs was determined using zeta potential, and the analysis revealed a zeta potential value of -18.1 mV which outlines a good stability (Supplementary Figure S3). On analysis of DLS results, the AC-AgNPs had a mean size ±SD of 89.77 ± 16.50 nm, with polydispersity index of 0.242 (Supplementary Figure S4).

3.6 Antiplasmodial assays

High antiplasmodial activity was found for AC-AgNPs against 3D7 (CQ-sensitive) and RKL9 (CQ-resistant) *Pf* strains. Based on IC₅₀ values, AC-AgNPs exhibited higher antiplasmodial activity as compared to that of AC-CE irrespective of plasmodial strain, and differences were statistically significant (p < 0.0001): 8.05 µg/mL vs. 20.27 µg/mL for 3D7, and 10.31 µg/mL vs. 32.55 µg/mL for RKL9. The standard drug CQ exhibited IC₅₀ values of 0.04 µg/

mL and 0.35 μ g/mL against *Pf* strains 3D7 and RKL9 (Figures 6A, B). The SYBR green assay findings were supported by microscopic data. AC-AgNPs and AC-CE elicited no quenching effects as no statistically significant difference was found between fluorescence counts of NPs, standard drug and plant extract (Supplementary Figure S5).

3.7 Hemolysis induced by the green AC-AgNPs

We have noted that hemolysis rates were dependent on substance, dose and time (Figures 7A, B). After 30 min, hemolysis rates elicited by AC-AgNPs and AC-CE were significantly higher than that of CQ at doses \geq 62.5 µg/mL. At these concentrations (125–500 µg/mL), hemolysis rates ranged from 6.25%–13.15% for CQ, 14.55%–48.14% for AC-AgNPs, and 5.50%–40.95% for AC-CE (Figure 7A). To be noted, HC₅₀ was not achieved after 30-min incubation as hemolysis rate was below 50% at 500 µg/mL. After 24 hour-incubation, hemolysis rates increased for all substances tested, with the highest values in AC-AgNPs, treated samples (maximum hemolysis of 98.14% at 500 µg/mL). Statistically significant difference between AC-AgNPs, AC-CE, and CQ were seen at doses \geq 8 µg/mL (Figure 7B).

3.8 Toxic effect of the AC-AgNPs against mosquito species

Mortality of Cx. quinquefasciatus, Ae. aegypti and An. stephensi larval stages was followed 24h, 48h and 72 h after treatment with AC-AgNPs. Mortality rates of the three mosquito species increased as a function of time and concentration (Supplementary Figure S6). After 48 h incubation, larval mortality rates were 100% at doses 23.5 µg/mL for Cx. quinquefasciatus, 20 µg/mL for Ae. aegypti, and 15 µg/mL for An. stephensi (Supplementary Figure S6). AC-AgNPs were more lethal against An. stephensi regardless of exposure time, with LC_{50} values of 10.67 $\mu g/mL$ and 5.85 $\mu g/mL$ at 24 h- and 48 hexposure, respectively. These values were 16.71 µg/mL and 7.52 µg/ mL for Ae. aegypti; 18.41 µg/mL and 8.97 µg/mL for Cx. quinquefasciatus, respectively. Regardless of exposure time, the larvicidal activity of AC-AgNPs was much higher than that of AC-CE for which LC₅₀ of 231.41 µg/mL, 110.33 µg/mL and 53.15 µg/mL against Cx. quinquefasciatus, Ae. aegypti and An. stephensi were found after 24 h exposure, respectively (Tables 4, 5). Interestingly, AC-CE did not cause any larval mortality at LC_{50} and LC₉₀ concentrations found for AC-AgNPs.

3.9 Behavioral and morphological impact of the AC-AgNPs on the larvae

The stereomicroscopic observations of *Ae. aegypti, An. stephensi* and *Cx. quinquefasciatus* larval stages treated with AC-AgNPs are depicted in Figure 8, and revealed the induction of behavioral and morphological changes in mosquito larvae. It was observed that swimming behavior of larvae was reduced,



Patterns of the green synthesized AC-AgNPs using PXRD (A), SAED (B), EDX (C), and FTIR (D). In (A), intensity of peaks is presented as arbitrary units (a.u). In (A), peaks with blue and black round shape indicate silver nanocrystals and silver chloride nanocrystals, respectively. AC-AgNPs, *Alchornea cordifolia* silver nanoparticles; EDX, Energy dispersive X-ray spectroscopy; FTIR, Fourier transformed infrared spectroscopy; PXRD, Powder X-ray diffraction; SAED, Selected area electron diffraction.

with morbid larvae at the bottom of bowls and unable to swim to the surface. Several morphological changes were noted in AC-AgNPs-treated larvae and these included loss of external hairs/ bristles, swelling of the apical cells, pigmentation of the body, shrinkage of the larvae, and necrosis and thickening of the epidermis (Figure 8).

S.No.	Position (2 0)	Peak amplitude (a.u)	FWHM (20)	Cos (θ)	Miller indices (HKL)	Nature	Size (nm)
1	27.83	64.55	1.1219	0.97065	(111)	AgCl	7.62
2	32.31	146.69	1.2155	0.96051	(200)	AgCl	7.11
3	38.15	66.67	0.6446	0.94509	(111)	Ag	13.62
4	46.20	75.30	1.3896	0.91982	(200)	Ag	6.49
5	54.81	19.36	0.9351	0.88778	(311)	AgCl	10.00
6	57.41	16.20	0.7762	0.87710	(222)	AgCl	12.19
7	64.59	20.41	0.4035	0.84531	(220)	Ag	24.33
8	67.41	10.75	0.6573	0.83191	(400)	AgCl	15.17
9	77.40	31.95	1.1259	0.78043	(311)	Ag	9.44

TABLE 2 Principal characteristic values of the powder X-ray diffractogram of AC-AgNPs.

a.u, Arbitrary units; FWHM, full width at half maximum; Cos, Cosinus.

TABLE 3 Functional groups at a given wavenumber for the FTIR spectra of AC-AgNPs.

Absorption (cm ⁻¹)	Appearance	Functional groups	Compound class
3,416	Medium	N-H stretching	Primary amine
2,927	Sharp	C-H stretching	Alkane
2,857	Medium	C-H stretching	Alkane
2,373	Sharp	O=C=O stretching	Carbon dioxide
		C≡N stretching	Nitriles
2,323	Weak	O=C=O stretching	Carbon dioxide
		C≡N stretching	Nitriles
1,739	Sharp	C=O stretching	Ester, Aldehyde, Saturated aliphatic, or $\delta\text{-lactone}$
1,630	Medium	C=C stretching	Conjugated alkene
1,458	Medium	C-H bending	Alkane (methylene or methyl group)
1,377	Medium	C-H bending	Aldehyde or Alkane (gem dimethyl)
1,023	Sharp	C-O stretching	Alcohol, Ether, Carboxylic acids
682	sharp	C=C bending	Alkene or Aromatics
600	sharp	C-I stretching	Halo compound
532	Sharp	C-Br stretching	Alkyl halides

4 Discussion

Vector-borne diseases such as malaria are an important public health problem throughout the world especially in Cameroon. This study demonstrated good hemocompatibility and high biocidal potential of green synthesized AgNPs using *A. cordifolia* leaves (Euphorbiaceae).

The synthesis of AC-AgNPs through green route was rapid as color change was noted a few minutes after mixing AC-CE and AgNO₃ aqueous solutions, thereby outlining the onset of AC-AgNPs synthesis through reduction of Ag^+ ions into Ag^0 . This observation was further confirmed upon analysis of UV-Vis spectra with a peak at 445 nm wavelength. Karthik and others showed a close value

(434 nm) for *Acalypha indica*, another Euphorbiaceae plant (Karthik et al., 2017). The UV-Vis peak corresponds to SPR phenomenon during which electron on NPs surface enter into resonance with the wavelength of incident light (Kojom Foko et al., 2021). The SPR band was increasing with parameters used for optimizing AC-AgNPs synthesis (i.e., AgNO₃ concentration, AC-CE volume, incubation time and incubation temperature), and such findings were seen previously with plants growing in Cameroon, especially *Megaphrynium macrostachyum* (Eya'ane Meva et al., 2016), and *Selaginella myosurus* (Belle Ebanda Kedi et al., 2018).

The biofabricated AC-AgNPs were small and mostly spherical which is consistent with earlier reports using



laboratory (A) P. falciparum 3D7 and (B) RKL9 strains. AC-AgNPs, Alchornea cordifolia silver nanoparticles: AC-CE. Alchornea cordifolia crude extract; CQ, Chloroquine; $\mathrm{IC}_{\mathrm{50}},\,\mathrm{50\%}$ Inhibition concentration. CQ was used as standard drug. Reference P. falciparum strains 3D7 (CQ-sensitive) and RKL9 (CQ-resistant) were used. The experiments were triplicated.

Morinda citrifolia and Adiantum raddianum (Suman et al., 2015; Govindarajan et al., 2017a). Using a systematic review, we previously reported that the bulk of NPs tested against Plasmodium parasites and mosquito vectors were spherical with a large range of size (Kojom Foko et al., 2019). Also, the nucleation theory of NPs synthesis suggests that slow rate of seed formation is expected to lead to broad size distribution of NPs (Liu et al., 2020). This result suggests that AC-AgNPs nucleation process was heterogeneous, and this can be influenced by several factors such as mixing time and solvation dynamics (Thanh et al., 2014; Deshpande et al., 2021). Size and shape of green NPs are modulated by complex interactions of plant- and experiment condition-related factors, and are crucial parameters that determine their physico-chemical and biological activities (Pal et al., 2007; Adams et al., 2014). Based on TEM analysis, AC-AgNPs were polydispersed with varied size. Such variation is commonly seen in AgNPs fabricated with plant extracts (Kojom Foko et al., 2019; Kojom Foko et al., 2021).

The analysis of PXRD and SAED patterns outlined that AC-AgNPs were polycrystalline with a crystallinity percentage of 84.13% and presence of additional peaks on diffractogram. This finding outlines that biosynthesized AC-AgNPs were not totally pure. At



Alchornea cordifolia crude extract; CQ, Chloroquine; IC₅₀, 50% Inhibition concentration. The experiment was performed in triplicate. Saponin was used as positive control and PBS as negative control. Statistically significant at *p < 0.05, **p < 0.01 and ***p < 0.01

nanoscale level, a large number of metals present as face-centered cubic structures and tend to agglomerate due to high tension surface of ultrafine NPs (Belle Ebanda Kedi et al., 2018), thereby explaining the crystalline nature of AC-AgNPs. Also, with increasing nucleation and growing over time, NPs form twinned structures that then multiply with their surfaces bounded to cubic facets with the lowest binding energy (Annamalai and Nallamuthu, 2016). The SAED pattern clearly confirmed the crystalline nature of AC-AgNPs.

Silver atom was mainly involved in AC-AgNPs synthesis while other atoms such as oxygen and chlorine were also found, and these could be due to phytochemical compounds in AC-CE. FTIR spectrum revealed the presence of several peaks corresponding to functional chemical groups (e.g., O-H, C≡N, C=C) attributable to alkanoids, terpenoids, flavonoids, phenols, steroids, anthraquonones or saponins, and confirmed results from GC-MS-based phytochemical analysis done here and reported elsewhere (Osadebe et al., 2012). These compounds are likely involved in reducing silver ions during NPs synthesis along with their capping

Time	LC ₅₀	95% CI	LC ₉₀	95% CI	Regression equation ^a	χ^2 (<i>p</i> -value)		
Culex quinquefa	Culex quinquefasciatus							
24 h	231.41	200.01-308.77	524.35	450.81-703.41	y = -1.16 + 0.004x	3.31 (0.85)		
48 h	188.71	161.69-214.03	431.49	362.87-539.18	y = -0.92 + 0.005x	2.84 (0.78)		
72 h	147.50	123.61-170.73	391.20	334.12-507.03	y = -0.76 + 0.005x	3.91 (0.69)		
Aedes aegypti	Aedes aegypti							
24 h	110.33	85.44-214.11	160.14	141.00-348.11	y = -3.25 + 0.036x	8.44 (0.01)		
48 h	90.30	71.80-109.37	141.42	123.02-241.88	y = -2.41 + 0.020x	5.15 (0.97)		
72 h	71.52	66.27-100.11	113.11	91.76-199.44	y = -2.30 + 0.044x	4.01 (0.17)		
Anopheles steph	Anopheles stephensi							
24 h	53.15	47.33-60.01	121.88	87.14-199.01	y = -2.88 + 0.050x	5.86 (0.001)		
48 h	41.57	37.51-50.43	102.11	75.55-111.77	y = -3.36 + 0.081x	8.30 (0.32)		
72 h	37.23	28.83-52.68	57.01	41.15-63.52	y = -2.52 + 0.070x	0.71 (0.15)		

TABLE 4 Larval toxicity of AC-AE against larval stages of Cx. quinquefasciatus, Ae. aegypti and An. stephensi after 24h, 48h and 72 h exposure.

Control no larval mortality recorded; LC₅₀, LC₉₀ Lethal concentration of the substance that kills 50%, 90% of the exposed larvae, respectively; LC₅₀ and LC₉₀ are expressed in µg/mL; 95% CI, Confidence interval at 95%; γ^2 Chi square; Statistical significance was set at *p*-value <0.05.

^aDetermined using the probit model.

TABLE 5 Larval toxicity of AC-AgNPs against larval stages of Cx. quinquefasciatus, Ae. aegypti and An. stephensi after 24h, 48h and 72h exposure.

Time	LC ₅₀	95% CI	LC ₉₀	95% CI	Regression equation ^a	χ^2 (<i>p</i> -value)		
Culex quinquefa	Culex quinquefasciatus							
24 h	18.41	11.75-21.02	24.35	19.11-38.96	y = -9.16 + 0.59x	62.31 (<0.0001)		
48 h	8.97	6.27-10.60	17.22	11.44–19.52	y = -7.32 + 0.52x	57.14 (<0.0001)		
72 h ^b	_	_	_	_	_	_		
Aedes aegypti	Aedes aegypti							
24 h	16.71	15.86-17.53	24.16	20.98-27.59	y = -2.27 + 1.36x	7.15 (0.52)		
48 h	7.52	5.81-9.42	16.63	15.54-17.97	y = -1.35 + 1.50x	10.3 (0.24)		
72 h ^b	_	_	_	_	_	_		
Anopheles stephensi								
24 h	10.67	7.59-13.75	21.62	12.49-28.76	y = -3.58 + 1.48x	5.35 (0.48)		
48 h	5.85	3.75-8.94	12.06	10.55-19.80	y = -5.35 + 2.35x	8.30 (0.32)		
72 h ^b	—	_	_	_	_	_		

Control no larval mortality recorded; LC₅₀ LC₅₀ Lethal concentration of the substance that kills 50%, 90% of the exposed larvae, respectively; LC₅₀ and LC₉₀ are expressed in µg/mL; 95% CI, Confidence interval at 95%; χ^2 Chi square; Statistical significance was set at p-value <0.05.

^aDetermined using the probit model.

^bNo data were computed as all larvae were dead after 48 h

and stabilization (Hawadak et al., 2022). We found two predominant compounds in plant extract (2-hexadecen-1-ol, 3,7,11,15tetramethyl-, [R-[R*,R*-(E)]]- (acyclic diterpene alcohol) and phytyl tetradecanoate (fatty acid phytyl ester) using GC-MS. Although mechanism of action of NPs reduction is uncertain, it is likely these compounds, alone or in combination with other compounds in plant extract, were involved in reduction, capping and stabilization of AC-AgNPs.

Zeta potential defines the stability of colloidal suspensions such NPs, and is a common parameter used to surface charge on a particle. In this study, zeta potential of AC-AgNPs was -18.1 mV. This value indicates a good stability of AC-AgNPs in dispersion medium. Indeed, negative surface charge is due to the binding affinity of AC-CE compounds with the NPs, conferring stability of AC-AgNPs and preventing several phenomena such as aggregation, sedimentation or flocculation which are known impair stability of particles (Faisal et al., 2021).

High lethal activity of green AC-AgNPs against Pf strains 3D7 and RKL9 was observed, with $IC_{50} < 10 \mu g/mL$ for 3D7 and $IC_{50} < 20 \,\mu\text{g/mL}$ for RKL9. This is consistent with value reported by Hawadak et al. and Rajkumar et al. using green NPs mediated by Eclipta prostrata and Azadirachta indica, respectively (Rajakumar et al., 2015; Hawadak et al., 2022). In contrast, our values are lower than those found previously with different



Morphological deformities induced by the exposure to AC-AgNPs (LC₅₀ dose) on larval stages of *Ae. aegypti*, *Cx. quinquefasciatus*, and *An. stephensi*. **(A)** *Ae. aegypti* larvae (Control), **(B–D)** *A. aegypti* larvae (AC-AgNPs-treated), **(E)** *Cx. quinquefasciatus* larvae (Control), **(F–H)** *Cx. quinquefasciatus* larvae (AC-AgNPs-treated), **(I)** *An. stephensi* larvae (Control), **(J)** *An. stephensi* larvae (AC-AgNPs-treated). Arrows indicate the difference morphological abnormalities seen in AC-AgNPs-treated larvae: swelling of the apical cells (blue arrows), pigmentation of body (yellow arrows), shrinkage of the larvae (red arrows), loss of external anal and head hairs/bristles (green arrows), necrosis and thickening of the epidermis (black arrows).

Plasmodium strains (Kojom Foko et al., 2019). This antiplasmodial activity exhibited by the AC-AgNPs is due to above mentioned phytochemical compounds which served as bioreactors for NPs reduction and capping. Several studies suggested potential mechanisms of action of NPs against Plasmodium parasites (Coxgeorgian et al., 2019; Abubakar et al., 2020). The NPs could induce parasite death by acting on several targets including cell membrane, enzymes and internal organelles (Shakeel et al., 2016; Kamaraj et al., 2017; Varela-Aramburu et al., 2020). Using in vivo model, Karthik and others showed that antiplasmodial activity of marine actinobacterialmediated gold NPs was associated with increased production of tumor growth factor but reduction in tumor necrosis factor, thereby emphasizing an immunomodulatory role of NPs (Karthik et al., 2013). Pf is highly prevalent in Cameroon (Kojom Foko et al., 2018; Antonio-Nkondjio et al., 2019; Kojom Foko et al., 2021), and our findings suggest that AgNPs could be interesting as antimalarial drug. A large number of NPs-related chemical and/or physical factors could explain discrepancies obtained between our findings

and those from previous studies. These included mainly size distribution, shape, capping/reducing agents, aggregation and surface charge. Even though AC-AgNPs synthesized in this study showed broad size distribution (range 6–28 nm), these are still interesting for future antimalarial drug development. Optimal NPs size for integration into human drugs varies depending on the specific drug and its intended application (Mitchell et al., 2021). This size distribution found here is consistent with previous studies on potential of MNPs as either drug delivery agent (i.e., passive targeting to enhance the accumulation of drugs in tumors) or antimalarial drug (i.e., active targeting to specific cells/tissues) (Santos-Magalhães and Mosqueira, 2010; Rahman et al., 2019). It should be interesting to conduct more studies to define consistent NPs size cut-offs for antimalarial therapy purposes.

It is known that antimalarial drugs such as ACTs, the current medicines used for treating uncomplicated malaria in most of endemic countries, can induce hemolysis in patients (Rehman et al., 2014). Therefore, new antimalarial drug candidates should be screened for hemocompatibility profile. The hemolysis rate was

below at 50% after 30 minute-incubation, thereby underlining a $HC_{50} > 500 \,\mu$ g/mL for the AC-AgNPs. The biofabricated AC-AgNPs were therefore highly hemocompatible, consistent with findings of Hossain and coworkers, who reported HC50 of 700 and 800 µg/mL for green aqueous and ethanolic NPs mediated by Andrographis paniculata stem (Hossain et al., 2019). Hemolysis increased as a function of time for AC-CE and AC-AgNPs which is in line with previous studies (Laloy et al., 2014; Avitabile et al., 2020). Hemolysis activity of NPs is strongly dependent on their size with higher hemolytic activity seen in smaller NPs (Chen et al., 2015). Thus, the small size of AC-AgNPs could likely explain their hemolytic activity (De La Harpe et al., 2019). Also, the anti-hemolytic activity of AC-AgNPs can be partially attributed to biomolecules coated on their surface. In fact, polyphenols are known to delay solubilization and inhibit oxidation of lipid frame; terpenes and flavonoids prevent interactions with hydrophobic parts of proteins and lipids, resulting in protecting and stabilizing cells membrane (Hoshyar et al., 2016; De La Harpe et al., 2019).

The phytofabricated AC-AgNPs exhibited a high toxicity against larval stages of Ae. aegypti, Cx. quinquefasciatus and An. stephensi, with LC_{50} below 20 µg/mL. Consisting with previous reports on diverse families of plants such as A. raddianum (Pteridaceae), Hugonia mystax (Linaceae), Psidium guajava (Myrtaceae), Holostemma ada-kodien (Apocynaceae) and Aganosma cymosa (Apocynaceae) (Govindarajan et al., 2017a, 2017b; Benelli and Govindarajan, 2017; Alyahya et al., 2018; Ntoumba et al., 2020). In contrast, some authors reported $LC_{50} > 20 \,\mu g/mL$ for AgNPs fabricated with Ventilago maderaspatana (Rhamnaceae), Naregamia alata (Meliaceae), Hedychium coronarium (Zingiberaceae) and Sargassum wightii (Sargassaceae) (Azarudeen et al., 2017a, 2017b; Kalimuthu et al., 2017; Murugan et al., 2017). The discrepancy observed between studies is likely due to a cocktail of factors including the phytochemical composition of plant used for NPs synthesis, size/shape of NPs and mosquito strains. The mechanisms through which NPs induce larval mortality are still elusive, but it is thought that nanosized materials such as NPs can easily pass through insect exoskeleton and cell membrane, bind to sulphur-containing proteins and/or DNA which then lead to interference with homeostatic and physiological processes essential for larvae (e.g., copper homeostasis, osmoregulatory and spiracle-related respiratory systems) (Armstrong et al., 2013; Kojom Foko et al., 2021; Araújo et al., 2022). Other authors reported NP-induced physical and molecular degradation of insect gut as additional death cause (Kalimuthu et al., 2017, 2016; Banumathi et al., 2017; Ishwarya et al., 2017; Suganya et al., 2019). Also, these putative mechanisms could also explain behavioral and morphological modifications in AC-AgNPstreated larvae seen in this study and by several earlier studies on extracts and NPs (Banumathi et al., 2017; Ishwarya et al., 2017; Suganya et al., 2019).

5 Conclusion

In this study, we synthesized, optimized, characterized and evaluated some medical applications of green AC-AgNPs

including antiplasmodial, hemocompatibility and larvicidal potential. The synthesis was rapid and the optimized AC-AgNPs were mostly spheroidal, small-sized, dispersed, stable and polycrystalline in nature. Several phytochemicals including alkanoids, terpenoids, flavonoids, phenols and steroids were responsible for reduction, capping and stabilization of AC-AgNPs. The AC-AgNPs exhibited higher antiplasmodial and mosquito larvicidal activities compared to plant extract. The AC-AgNPs induced several mortality-associated behavioral and morphological changes in larval stages of Ae. aegypti, An. stephensi and Cx. quinquefasciatus. Finally, the AC-AgNPs exhibited good hemocompatibility with $HC_{50} > 500 \,\mu\text{g/mL}$. In worrying context of resistance of malaria parasites to current drugs and mosquitoes to different classes of insecticides, green nanotechnology could be a valuable and cutting-edge alternative for advanced drug/insecticide development and research.

Data availability statement

The original contributions presented in the study are included in the article/Supplementary Material, further inquiries can be directed to the corresponding author/s.

Author contributions

LPKF and VS conceptualized the study. LPKF performed laboratory experiments and drafted the first version of the manuscript. JH helped in laboratory experiments. PBEK collected the plant and brought it to the National Herbarium for taxonomical authentication. JH, VV, PBEK, and CEEM helped in data interpretation. JH, VV, PBEK, CEEM, KR, VP, and VS revised the manuscript for important intellectual content. VV and KR supervised larvicidal assays and validated data. VS supervised the work at all stages. All authors read and approved the final version of the manuscript before submission.

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Supplementary material

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Glossary

AC Alchornea cordifolia ACT Artemisinin based combination therapy AgNPs Silver nanoparticles ANOVA Analysis of variance a.u Arbitrary units CE Crude extract CI Confidence interval Cos Cosinus CQ Chloroquine DDT Dichlorodiphenyltrichloroethane DLS Dynamic light scattering DNA Deoxyribonucleic acid EDX Energy dispersive X-ray ELISA Enzyme-linked immunosorbent assay FCC Face centered cubic FTIR Fourier transformed infrared spectroscopy FWHM Full width at half maximum GC-MS Gas chromatography coupled with mass spectrometry HC50 50% hemolysis concentration

IC50 50% inhibition concentration ICMR Indian Council of Medical Research JCPDS Joint Committee on Powder Diffraction Standards LC Lethal concentration MNPs Metallic nanoparticles NIMR National Institute of Malaria Research PBS Phosphate-buffered saline Pf Plasmodium falciparum PI Polydispersity index PXRD Powder X-ray diffraction RBC Red blood cell SAED Selected area electron diffraction SD Standard deviation SEM Scanning electron microscopy SPR Surface plasmon resonance TEM Transmission electron microscopy UD The University of Douala UV-Vis Ultraviolet-Visible WHO World Health Organization

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Bacterial-mediated synthesis and characterization of copper oxide nanoparticles with antibacterial, antioxidant, and anticancer potentials

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The application of novel bacterial strains for effective biosynthesis of nanoparticles minimizes negative environmental impact and eliminates challenges of available approaches. In the present study, cell-free extract of Stenotrophomonas sp. BS95. was used for synthesis of copper oxide nanoparticles (CuONPs). Characterization of crude and calcined CuONPs was carried out by UV-vis spectroscopy, X-ray diffraction (XRD), fourier transform infrared (FTIR) spectroscopy, zeta potential, dynamic light scattering, field emission scanning electron microscopy, transmission electron microscopy, and atomic force microscopy. Afterward, biogenic CuONPs were evaluated for antibacterial, antioxidant, and cytotoxic effects using broth microdilution method, DPPH assay and alamarBlue assay, respectively. Finally, molecular mechanisms behind anticancer effects of CuONPs was ascertained by real time PCR. UV-vis absorbance spectra registered surface plasmon resonance peaks at 286 nm and 420 nm for crude and calcined CuONPs, respectively. FTIR spectra exhibited bands associated with organic functional groups of bacterial proteins, confirming capping and functionalization of CuONPs. The average crystallite size of crude and calcined CuONPs was determined as 18.24 and 21.3 nm by XRD, respectively. The average zeta potentials of crude and calcined CuONPs were as -28.57 \pm 5.13 and $-29.47 \pm 4.78 \,\text{mV}$, respectively, indicating their high stability. Electron microscopy revealed that crude and calcined CuONPs were roughly spherical particles with an average size of 35.24 ± 4.64 and 43.68 ± 2.31 nm, respectively. Biogenic CuONPs induced antibacterial effects with minimal inhibitory concentrations ranging from 62.5 to 1,000 µg/ml against Gram-negative and Gram-positive strains. The antioxidant activity of crude and calcined CuONPs was found to be 83% + 2.64% and 78% + 1.73%, respectively. More intriguingly, CuONPs exerted considerable cytotoxic effects on human colon and gastric adenocarcinoma cells, while induced low toxicity on normal cells. Anticancer effects of biogenic CuONPs were confirmed by significant changes induced in the expression of apoptosis-related genes, including P53, BAX, BCL2 and CCND1. Hence, biosynthesized CuONPs could be considered as potential antimicrobial, antioxidant and anticancer agents.

KEYWORDS

stenotrophomonas sp. BS95, copper oxide nanoparticles, antimicrobial activity, antioxidant effects, anticancer properties

Introduction

Nanotechnology is as an evolving interdisciplinary research field that focuses on nanoparticles (NPs) with improved size-dependent features, such as robustness of the colloidal moiety, great surface area and high bioavailability to name a few (Pan et al., 2021). Conventionally, synthesis of NPs can be performed by physical, chemical and mechanical approaches, however, high cost and toxicity of these methods have shifted fabrication of NPs toward biological systems (Chandrasekaran et al., 2020; Lahiri et al., 2021). Among all microorganisms, bacteria are attractive candidates to synthesize NPs because of remarkable benefits like high stability, short generation time, mild experimental condition, easy culture, resistance to most toxic heavy metals and their ability to produce sustainable NPs at a large scale (Yusof et al., 2019; Waris et al., 2021).

Unique characteristics of metal oxide NPs made them suitable candidates for various commercial and domestic applications, such as energy harvesting, food processing and environmental protection (Tsuzuki, 2021). In addition, there is a growing interest to develop nano-scale pharmaceuticals for management of global health concerns. For instance, antibiotic resistance crisis, which has placed a substantial clinical and financial burden on healthcare systems, can be controlled by antibacterial potential of NPs. Furthermore, increasing incidence and mortality rate of cancer, which is due to poor diagnosis and low specificity of common chemotherapeutics, can be reversed by novel nano-based agents (Pearce et al., 2017; Fu et al., 2018; Zhang et al., 2018).

Copper oxide nanoparticles (CuONPs) have attracted much attention because of appropriate redox potential, high specific surface area, and excellent stability in different solutions (Nagajyothi et al., 2017; Verma and Kumar, 2019). The aim of present study was to synthesize CuONPs by a toxic-free, rapid, and eco-friendly approach and evaluate its biological activities. To do so, CuONPs were first synthesized by Stenotrophomonas sp. BS95, and then characterization was carried out by well-established techniques; Optical properties were defined by ultraviolet-visible (UV-vis) spectroscopy and fourier transform infrared (FTIR) defined functional groups. X-ray diffraction (XRD) was used to determin the crystallite size, while physical stability and surface charge were measured by zeta potential. Analysis of hydrodynamic particle size was carried out by dynamic light scattering (DLS) and electrone microscopy was used to study various surface phenomena such as morphology and roughness. Then after, crude and calcined CuONPs were assessed for antibacterial, antioxidant, and cytotoxic effects using broth micro-dilution method, DPPH assay and alamarBlue assay, respectively. Finally, molecular mechanisms behind anticancer effects of biogenic CuONPs was unraveled by real time polymerase chain reaction (PCR).

Materials and methods

Biosynthesis of CuONPs by hydrothermal cell lysate supernatant

Synthesis of CuONPs in the present study was carried out using bacterial cell lysate supernatant (CLS), as previously described (Nakhaeepour et al., 2019). A cold-tolerant bacterium, namely, *Stenotrophomonas* sp. BS95 was isolated from alpine soil samples collected in western Iran, and identified using 16s rRNA gene sequencing analysis. The data presented in this study are deposited in the GenBank repository, accession number OQ253458. To synthesize CuONPs, this strain was cultured in tryptic soy broth (TSB) medium (Merck) and incubated in a shaking incubator at 28°C and 150 rpm for 72 h. Afterward, the culture medium was centrifuged at 7,500 rpm for 15 min and the pellet was washed with 1 mM NaCl solution (Merck). Then, the cell pellet was resuspended in distilled H₂O and after 24 h incubation, it was placed in an ultrasonic bath sonicator for 20 min to obtain the cell lysate. Upon centrifugation at 5,000 rpm for 20 min, the supernatant was added to 0.01 M copper (II) sulfate pentahydrate (CuSO₄,5H₂O) and heated at 121°C for 20 min. Followed by centrifugation at 10,000 rpm for 10 min, biogenic CuONPs were obtained and washed with distilled H2O and ethanol. Then, the purified NPs were dried in a vacuum oven at 80°C for 4 h to achieve crude CuONPs, and calcined CuONPs were obtained after incubation in a muffle furnace (470°C) for 4 h.

Characterization of synthesized CuONPs

UV-vis spectroscopy

To define optical properties of crude and calcined CuONPs, UVvis spectroscopy was used (Ssekatawa et al., 2022). In this regard, 1 mg of each sample was dispersed in distilled H_2O and dispensed into different cuvettes. Optical properties of samples were then obtained by UV-vis spectrophotometer (Shimadzu UV-1700, Japan), scanning at a resolution of 1 nm between 200 and 800 nm ranges.

FTIR spectroscopy

The organic functional groups of crude and calcined CuONPs were identified by FTIR spectroscopy (Sarkar et al., 2020). In this regard, 2 mg of biogenic CuONPs and 2 g of potassium bromide (KBr) were mixed and compressed to obtain translucent circular pellets. Then, samples were scanned through 4,000 to 400 cm⁻¹ wavenumber range and a resolution of 4 cm⁻¹ for at least 32 scans per sample using Thermo Nicolet 6700 FTIR spectrometer (Nicolet Avatar, Madison, WI, United States of America). To note, KBr pellet was used as control.

XRD analysis

To confirm crystallinity of crude and calcined CuONPs, XRD analysis was performed (Nakhaeepour et al., 2019). To do so, GNR Explorer X-ray diffractometer (Italy) fitted with Cu-Ka radiation ($\lambda = 1.5418 \text{ A}^\circ$) and scanning from $2\theta = 20^\circ - 80^\circ$, with a voltage of 40 kV, current of 30 mA and integration time of 0.2 s/step was used. Obtained data was visualized in OriginPro 2019b software, and validated by standard CuONPs 2 θ values from the International Center for Diffraction Data (ICDD) database. The average crystallite size was calculated using the following formula:

 $d = 0.9 \lambda / \beta \cos \theta$

in which d is the average crystallite size, β is full peak width at half maximum, λ is the wavelength of X-ray (1.5418 Å) and θ is the 2 θ angle in peak.

DLS and zeta potential

The average size and stability of crude and calcined CuONPs were determined at neutral pH and room temperature as previously described (Sarkar et al., 2020). Briefly, to evaluate the average size distribution, 100 μ g/ml of each sample was dispersed in ethanol and sonicated for 5 min. Afterward, samples were examined by DLS analyzer (vasco3. Cordouan, France) for three times, and zeta potential was determined by an electrophoretic light scattering instrument (Zeta Compact, CAD, France).

FESEM, TEM and AFM analysis

To determine the particle size distribution and nanostructure of crude and calcined CuONPs, FESEM was used (Ali et al., 2020). In summary, each sample was spread onto an aluminum tape and coated with gold to become a conductor. Micrographs were taken at different magnifications using FESEM (Mira 3-FEG TESCAN, Czech Republic), operating at around 30 kV accelerating voltage. Morphological and topographical characteristics of biogenic CuONPs were also studied by TEM (Ssekatawa et al., 2022). In this regard, 1 mg/ml of each sample was dispersed in ethanol, sonicated and finally loaded on copper grid thin films. Micrographs were obtained by TEM (912AB, LEO, Germany) operating at around 100 kV accelerating voltage. The particle size distribution (PSD) plots were obtained by determining the size of 50 particles for each sample using ImageJ software.

The surface morphology of crude and calcined CuONPs was investigated by AFM (Barani et al., 2021). To do so, NSC15-type silicon probes with the radius of tip curvature less than 10 nm were used and samples were analyzed by AFM device (Brisk model, Ara Research, Iran). The height of each sample was finally estimated in scanning areas of $1 \times 1 \mu m$.

Antibacterial activity

The antibacterial activity of biogenic CuONPs against pathogenic bacteria including *Bacillus subtilis* PTCC 1023, *Staphylococcus aureus* ATCC 25923, *Pseudomonas putida* KT2440 and *Escherichia coli* PTCC 1860 was evaluated using broth micro-dilution method (Binesh et al., 2021) The suspension of 1.5×10^8 CFU/mL bacteria (according to the 0.5 McFarland standard) was prepared in nutrient broth (NB, Merck). To assess minimum inhibitory concentration (MIC), 100 µl of CuONPs with serial dilutions (500–1.9 µg/ml) and 10 µl of pathogenic bacteria were transferred to each well of 96-well plates containing 100 µL Muller-Hinton broth medium (MHB, Merck) and incubated at 37°C in a shaker incubator for 18 h. The absorbance was then recorded at 630 nm using a spectrophotometer (Stat Fax 2100, England). To evaluate minimum bactericidal concentration (MBC), $5 \mu l$ from each dilution was spread on Muller-Hinton agar (MHA, Merck) plates and incubated at 37° C for another 24 h.

Analysis of antioxidant activity

The antioxidant activity of crude and calcined CuONPs was assessed by measuring their capability to scavenge synthetic stable radicals of 2,2-diphenyl-1-picrylhydrazyl (DPPH), as previously reported (Barani et al., 2021). To do so, 0.14 mM DPPH in methanol was added to each well of 96-well plates containing different concentrations of CuONPs (31.2, 62.5, 125, 250, 500, and 1,000 μ g/ml) and incubated at 37°C for 30 min, while ascorbic acid was used as a standard solution. Finally, the absorbance (A) was recorded at 517 nm using spectrophotometer (Awarness), and free radical scavenging activity of CuONPs was calculated using the following formula:

DPPH radical scavenging activity (%) =
$$\left(\frac{AS - (AT - AB)}{AS}\right) \times 100$$

in which AT is the absorbance of test wells, AB is the absorbance of blank wells, and AS is the absorbance of standard solution.

Cell culture, treatment and viability assay

Human colon and gastric adenocarcinoma cells (LoVo and MKN-45 cell lines, respectively) along with human dermal fibroblasts (HDF cell line) were purchased from Pasteur Institute (Tehran, Iran). MKN-45 and HDF cells were cultured in Dulbecco's modified Eagle's medium (DMEM, Biowest), whereas LoVo cells were grown in Roswell Park Memorial Institute-1640 (RPMI-1640, Biowest). All media were supplemented with 10% fetal bovine serum (FBS) (Biowest) and 1% penicillin-streptomycin (Biowest). Cells were maintained at 37° C in normoxic (95% and 5% CO₂ in air) and hypoxic (93% N₂, 5% CO₂ and 2% O₂) conditions.

To evaluate cytotoxic effects of CuONPs and determine the half maximal inhibitory concentration (IC_{50}) values, alamarBlue assay was performed (Movaffagh et al., 2021). To do so, LoVo and MKN-45 cells were seeded at a density of 14,000 cell/well, in each well of 96-well plates, while HDF cells were seeded at a density of 10,000 cell/well. After 24 h incubation, cells were treated with 50, 100, and 200 µg/ml crude and calcined CuONPs, while untreated cells were considered as control. At the end of treatments (24 h), alamarBlue solution (Sigma-Aldrich) was added to each well (20 µl/ well) followed by 2 h incubation at 37°C. Then, the absorbance (A) of each well was recorded at 600 nm using spectrophotometer (BioTek), and the viability of cells was calculated based on the following equation:

Cell viability (%) =
$$\left[100 - \left(\frac{AT - AU}{AB - AU}\right)\right] \times 100$$

in which AT is the absorbance of treated cells, AU is the absorbance of untreated cells, and AB is the absorbance of blank control.

Gene name	Forward (5' to 3')	Reverse (5' to 3')	Product size (bp)
TBP	ACAACAGCCTGCCACCTTA	GAATAGGCTGTGGGGGTCAGT	120
P53	GTTCCGAGAGCTGAATGAGG	TTATGGCGGGAGGTAGACTG	123
BAX	GGACGAACTGGACAGTAACATGG	GCAAAGTAGAAAAGGGCGACAAC	150
BCL2	GATGACTGAGTACCTGAACCG	CAGAGACAGCCAGGAGAAATC	124
CCND1	TGAAGGAGACCATCCCCCTG	TGTTCAATGAAATCGTGCGG	151

TABLE 1 List of primers, their sequence, and product length used in the present study.

Gene expression analysis

To assess the effects of biogenic CuONPs on the expression of apoptosis-related genes, real time PCR was applied (Mirzaei et al., 2022). In summary, total cellular RNA was extracted from LoVo cells treated with 100 µg/ml crude and calcined CuONPs, as well as untreated cells, using a total RNA isolation kit (DENAzist Asia). RNA purity was then evaluated by spectrophotometer at 260 and 280 nm (Nanodrop 2000 Thermo). For synthesis of cDNAs, M-MuLV reverse transcriptase (Parstous) was used according to the manufacturer's instruction. The validity of amplified cDNAs was then confirmed by PCR using TBP primers and final products were loaded on 1.5% agarose gel for electrophoresis. Real time PCR was conducted in an iQ5 real-time PCR detection system (Bio-Rad) using SYBR green master mix (BioFact) and specific primers listed in Table 1. To compare the level of gene expression, TBP transcripts were used as internal control and normalized values were plotted as relative fold change over untreated cells. PCR cycling conditions were as follows: 95°C for 5 min [95°C for 20 s, 58°C for 30 s, 72°C for 30 s] (35 cycles) for P53, BAX, BCL2 and CCND1 primers.

Statistical analysis

The data were analyzed by one-way ANOVA and Dunnett's multiple comparison tests using GraphPad Prism version 8.4.3 software. Values were expressed as mean \pm SD, and p values less than 0.05, 0.01, 0.001 and 0.0001 were considered to be statistically significant.

Results

Biosynthesis of CuONPs

In the present study, synthesis of CuONPs was carried out using CLS of a psychrotolerant *Stenotrophomonas* species. In this approach, complicated downstream processes were not required and thus, the risk of microbial contamination was low. The biosynthesis of CuONPs was validated by monitoring four flasks containing *Stenotrophomonas* sp. BS95 after 48 h incubation, bacterial CLS, CuSO₄ solution and the reaction mixture of bacterial CLS with CuSO₄ (Figure 1). The instant precipitation of green aggregates, which did not change over 24 h incubation, indicated the formation of CuONPs.

Optical properties of biogenic CuONPs

UV-vis spectroscopy was carried out to obtain optical properties of crude and calcined CuONPs. As presented in Figure 2A, surface plasmon resonance (SPR) peaks were recorded at 286 and 420 nm, which were assigned to efficient bio-reduction of CuSO₄ to CuONPs. Since the shift of UV-vis absorbance toward short wavelengths is mainly attributed to decrease in the NP size, the left shift observed in UV spectrum was associated with the small size of our biogenic crude CuONPs.

FTIR analysis

The FTIR spectra of crude and calcined CuONPs revealed similar absorbance bands within the wavenumber range of 3,200-3,600, 1,652 and 1,235-1,360 cm⁻¹ (Figure 2B). As shown, three distinct bands were registered for crude CuONPs in the range of 2,958, 1,080 and 547 cm⁻¹, while two unique absorbance bands were recorded for calcined CuONPs in the range of 521 and 620 cm⁻¹.

Analysis of XRD pattern

Completely similar to the standard pattern of CuO nanocrystals (JCPDS File No: 01-080-1917), analysis of XRD patterns revealed that all peaks representing CuONPs were present in our biogenic crude and calcined NPs (Figure 3). Bragg peaks positioned at 20 values of 32.5°, 35.7°, 38.7°, 46.2°, 53.5°, 65.8°, 66.2°, 68.08° and 72.3° were registered for crude CuONPs, corresponded to the planes of (100), (002), (101), (102), (110), (103), (200), (112) and (004), respectively. However, diffraction peaks observed at 20 values of 32.18°, 35.19°, 38.6°, 52.9°, 65.3°, 68.2° and 71.8° were registered for calcined CuONPs, assigned to crystal planes of (100), (002), (101), (110), (103), (112) and (004), respectively. The average crystallite size, measured using the Debye–Scherrer equation, was as 18.24 nm and 21.3 nm for crude and calcined CuONPs, respectively.

Measurement of stability and particle size

The average zeta potentials of crude and calcined CuONPs were as -28.57 ± 5.13 mV and -29.47 ± 4.78 mV, respectively, indicating their high stability (Figures 4A, B). The size distribution histograms obtained from DLS analysis showed that the size of the crude



Visual detection of biosynthesized CuONPs. Four conical flasks containing *Stenotrophomonas* sp. BS95 after 48 h incubation (A), bacterial CLS (B), CuSO₄ solution (C), and biogenic CuONPs (D).



The UV-vis spectrum of crude and calcined CuONPs (A); maximum absorbance bands were observed at 286 nm and 420 nm, respectively, FTIR spectrum of crude and calcined CuONPs (B).

CuONPs ranged from 25 to 110 nm with a mean distribution diameter of 55.92 nm, and from 38 to 128 nm with an average of 68.35 nm for calcined CuONPs. In addition, the calculated polydispersity index (PDI) were as 0.26 nm and 0.13 nm for crude and calcined CuONPs, respectively (Figures 4C, D).

Nanostructure analysis by FESEM, TEM and AFM

FESEM was used to determine morphology and size details of biogenic CuONPs. As presented in Figures 5A, B, crude and calcined CuONPs were spherical particles with the average PSD of 37.73 ± 3.27 and 48.37 ± 5.17 nm, respectively. Likewise, TEM micrographs revealed that crude and calcined CuONPs were roughly spherical particles with PSD values of 35.24 ± 4.64 and 43.68 ± 2.31 nm, respectively (Figures 5C, D).

To provide further insights into topological appearance and size of CuONPs, AFM images were prepared. As presented in Figure 6, crude and calcined CuONPs were detected as individual conical grains extending upwards with the average height as 5.155 and 6.547 nm, respectively.

Antibacterial effects of biogenic CuONPs

The antibacterial activity of CuONPs was evaluated against Gram-negative and Gram-positive bacteria using the broth microdilution method (Table 2). Calculating MIC values revealed that crude CuONPs inhibited the growth of *B. subtilis*, *S. aureus*, *P. putida* and *E. coli* at 62.5, 125, 250, and 500 µg/ml, respectively. Furthermore, the MIC values of calcined CuONPs were found to be 250 µg/ml for *B. subtilis* and *S. aureus*, and 500 and 1,000 µg/ml for *P. putida* and *E. coli*, respectively. This observation demonstrated





that calcination reduced the bacterial inhibitory activity of CuONPs. As also indicated in Table 2, MBC values for crude CuONPs in the present study were 250 μ g/ml for *B. subtilis, S. aureus* and *P. putida*

and 500 μ g/ml for *E. coli*. Likewise, the MBC values of calcined CuONPs were determined as 250 μ g/ml for *B. subtilis* and *S. aureus*, and 500 μ g/ml for *P. putida* and *E. coli*.



Antioxidant activity of biogenic CuONPs

Evaluating the antioxidant activity of biogenic CuONPs in the present study indicated that they scavenged DPPH radicals in a dose-dependent manner. As presented in Figure 7, upon administration of 2000 µg/ml crude CuONPs, $83\% \pm 2.64\%$ antioxidant activity was detected, while calcined CuONPs exhibited lower activity (78% ± 1.73%) in the same concentration. To note, in all concentrations, significant difference (p < 0.0001) was detected in the scavenging activity between ascorbic acid and biogenic CuONPs.

Anticancer properties of biogenic CuONPs

Assessment of cell viability revealed that CuONPs induced cytotoxic effects in a dose-dependent manner. As shown in Figure 8, treatment of LoVo, MKN-45 and HDF cells with 50, 100 and 200 µg/ml crude and calcined CuONPs significantly reduced cell viability in comparison with untreated cells. To note, toxic effects of CuONPs were also cell type-dependent, as viability of LoVo cells reduced to lower amounts in comparison with MKN-45 cells, and more interestingly, our biogenic CuONPs induced lowest toxic effects on non-cancerous HDF cells. Similarly, morphological alterations, in the form of dispersed cells with cytoplasmic granulation, were apparent upon administration of

crude and calcined CuONPs when compared with untreated cells (Figure 9). Calculated IC_{50} values of crude and calcined CuONPs on LoVo, MKN-45 and HDF cells are presented in Table 3. Due to high toxicity of CuONPs in LoVo cells, we also assessed their anticancer potential in hypoxic condition. Our results demonstrated that in comparison with untreated cells, biogenic CuONPs induced considerable toxicity in hypoxic condition as well (Table 3; Figure 8).

To unravel mechanisms underlying anticancer effects of crude and calcined CuONPs, alterations induced in the expression of apoptosisrelated genes was investigated by real time PCR. As shown in Figure 10, upon 24 h treatment of LoVo cells with 100 µg/ml biogenic CuONPs, significant (p < 0.0001) over expression of *P53* and *BAX* was detected in comparison with untreated cells. On the other hand, crude and calcined CuONPs significantly (p < 0.001) downregulated the expression of *BCL2* and *CCND1* when compared with untreated cells.

Discussion

Biosynthesis of CuONPs was carried out in the present study by an efficient non-toxic approach using *Stenotrophomonas* sp. BS95 CLS. It has been shown that cellular biomolecules such as enzymes and proteins presented in the bacterial CLS could reduce copper ions into copper atoms leading to CuONPs formation (Nakhaeepour et al., 2019; Bandeira et al., 2020). This



TABLE 2 The MIC and MBC of biogenic CuONPs against different Gram-positive and Gram-negative bacteria.

MIC (µg/ml)					MBC (µg/ml)			
	B. subtilis	S. aureus	P. putida	E. coli	B. subtilis	S. aureus	P. putida	E. coli
Crude CuONPs	62.5	125	250	500	250	250	250	500
Calcined CuONPs	250	250	500	1,000	250	250	500	500

mechanism might be involved in biosynthesis of CuONPs in our study as well. The reduction of $CuSO_4$ was subjected to spectral analysis by the UV-vis spectroscopy. SPR peaks of crude and calcined CuONPs were recorded between 286 and 420 nm, respectively. In line with these findings, it has been reported that UV-vis absorbance of biogenic CuONPs fall between 285 and 570 nm (Cheirmadurai et al., 2014; Duman et al., 2016). Furthermore, changes in the SPR peak position after calcination could be explained by elimination of capping agents, increased crystallite size and/or agglomeration of metal oxide NPs, as previously described (Tang et al., 2012; Kayani et al., 2015; Gharibshahi et al., 2017).

FTIR spectrophotometry was carried out to analyze bacterial biomolecules involved in reducing copper ions to CuONPs and subsequent capping. The organic functional groups entrapping CuONPs were determined as previously reported (Ali et al., 2020; Chandrasekaran et al., 2020; Kouhkan et al., 2020). Strong peaks at

3,200–3,600 cm⁻¹ demonstrated the N-H stretching vibrations of amine group or amide linkages in the protein contents of bacterial plasma membrane. The peak observed at 2,958 cm⁻¹ was associated with C-H stretching vibration of the aldehyde compound. The band at 1,652 cm⁻¹ was corresponded to the stretching vibration of C=O, usually found in proteins. The peaks seen at 1,080, 1,235 and 1,360 cm⁻¹ were further associated with the stretching vibration of C–N of aliphatic and aromatic amines. In this study, the bands at 547 and 521 cm⁻¹ for Cu–O confirmed the synthesis of CuONPs. To note, deletion of absorption peaks at 2,958 and 1,080 cm⁻¹ in calcined CuONPs might be due to the removal of peaks corresponding to functional groups, including aldehyde and amine.

The presence of sharp structural peaks in XRD patterns and crystallite size <100 nm revealed the nanocrystalline nature of crude and calcined CuONPs, which are in consistence with previous studies (Ahamed et al., 2014; Shankar and Rhim, 2014; Duman



The antioxidant activity of biogenic CuONPs at different concentrations. Ascorbic acid was used as a standard. ****p < 0.0001 indicate significant difference with ascorbic acid.



FIGURE 8

Dose-response curves representing the effects of crude and calcined CuONPs on viability of LoVo cells in normoxic (A) and hypoxic conditions (B), MKN-45 cells (C), and HDF cells (D). Results are shown as mean \pm SD. *p < 0.05, **p < 0.01***p < 0.001 and ****p < 0.0001 indicate significant difference with untreated cells.



FIGURE 9

Morphological alterations of cells after administration of crude and calcined CuONPs. Phase contrast photomicrographs of LoVo (A–C) and HDF (D–F) cells; untreated (A, D), treated with 200 µg/mL crude (B, E) and calcined (C, F) CuONPs.

TABLE 3 Calculated IC_{50} (µg/ml) values of crude and calcined CuONPs on different cell lines.

	LoVo	MKN	HDF	LoVo-hypoxia
Crude CuONPs	48.36	90.23	158.2	50.84
Calcined CuONPs	44.96	117.5	222.8	92.43

et al., 2016). Metal oxide NPs with zeta potential values higher than +30 mV or lower than -30 mV typically have high degree of stability, which is of utmost importance to avoid their agglomeration in colloidal solution (Joseph and Singhvi, 2019). In the present study, zeta potential measurement indicated higher stability of crude and calcined CuONPs compared to previous reports (Tiwari et al., 2016; Chandrasekaran et al., 2020).

Defining the morphology and PSD of biogenic CuONPs by FESEM, TEM and AFM revealed their spherical shape, while crude CuONPs presented smaller size in comparison with calcined CuONPs. These observations confirmed results obtained from XRD and DLS analysis, and are in consistence with previous reports on biogenic CuONPs (Nasrollahzadeh et al., 2015; Gu et al., 2018; Kouhkan et al., 2020; Sarkar et al., 2020).

Evaluating antibacterial activity of CuONPs implied on their inhibitory effects on both Gram-negative and Gram-positive pathogens, although antibacterial potential of crude CuONPs was higher than calcined CuONPs. In consistence with current findings, previous reports indicated that metal oxide NPs such as CuONPs induced remarkable antimicrobial activity due to their small size and extremely large surface area that provide better contact with microorganisms (Azam et al., 2012; Laha et al., 2014). Various studies have also demonstrated bactericide effects of CuONPs against same pathogenic bacteria, for instance, MBC values of crude CuONPs on *E. coli* and *S. aureus* have been reported as 250 and 2,500 µg/ml, respectively (Ren et al., 2009). In addition, MBC of mechanochemically synthesized CuONPs against E. coli and S. aureus were as 750 and 5,000 µg/ml, respectively (Moniri et al., 2019). In another research, the MBC of phytofabricated CuONPs was reported as 10,000 µg/ml for both E. coli and S. aureus (Alavi et al., 2021). Based on our findings, biogenic CuONPs induced higher growth inhibitory and toxic effects on Gram-positive bacteria compared with Gram-negative ones. In this regard, it has been shown that lipopolysaccharide layer on the outer membrane of Gram-negative bacteria acts as an effective protection against NPs (Franco et al., 2022) that could explain, to some extent, observed effects in the present study. CuONPs induce antibacterial effects through binding to the bacterial cell membrane, production of reactive oxygen species and release of Cu2+ ions that demolish DNA and cellular proteins, affect the membrane permeability, and finally induce cell death (Obeizi et al., 2020). Thus, considerable activity of biogenic CuONPs against B. subtilis, S. aureus, P. putida, and E. coli in our study might be mediated through the same mechanisms.

Current results also revealed high antioxidant activity of crude and calcined CuONPs, which has been attributed to the binding of transition metal ion catalysts to free radicals (Omran and Baek, 2021). Based on a recent report, the free radical scavenging activity of CuONPs may be enhanced by various bio-reductive groups (capping agents) of the bacterial proteins (Ssekatawa et al., 2022). As explained above, FTIR analysis confirmed the capping of crude CuONPs with aldehyde and amine groups, unlike calcined CuONPs. Therefore, higher antioxidant activity of crude CuONPs in our study was presumably due to bacterial-derived functional groups.

Malignancies of the gastrointestinal tract, including colorectal and gastric carcinomas, account for 36.2% of cancer mortality (Hong et al., 2022). Although use of chemical drugs is a systemic treatment for cancer patients, low specificity of common



Analysis of gene expression by real-time PCR. The expression of *P53, BAX* (**A**), *BCL2* and *CCND1* (**B**) were normalized and plotted as relative fold change compared with the untreated cells. Data are represented as mean \pm SD. **p < 0.01, ***p < 0.001, ****p < 0.0001 indicate significant difference with untreated cells.

chemotherapeutics causes many side effects that are mostly intolerable to patients and lead to reduced survival rates (Pearce et al., 2017; Fu et al., 2018; Zhang et al., 2018). To introduce novel and more effective therapeutics, we investigated anticancer potential of CuONPs, and obtained findings revealed that cytotoxicity of our biogenic CuONPs was dose- and cell type-dependent. Intriguingly, crude and calcined CuONPs induced more toxic effects on human colon and gastric adenocarcinoma cells than normal fibroblasts. Current results are in consistence with previous reports. For instance, it has been demonstrated that CuONPs synthesized by marine endophytic actinomycete and Vibrio sp. VLC. induced toxic effects on human lung and esophageal carcinoma cells in a dose-dependent manner, with IC₅₀ values of 500 and 37.52 µg/ml, respectively (Nakhaeepour et al., 2019; Zhao et al., 2022). In addition, low toxicity of CuONPs on normal HDF cells in the present study is in line with another research, which reported minimal toxic effects of CuONPs on human dermal fibroblasts (Sulaiman et al., 2018).

Hypoxia, a biological phenomenon in which oxygen level is below the tissue demand, is a feature of solid tumors, and an indicator of poor prognosis in many cancers including colon and gastric adenocarcinomas (Qi et al., 2020; Pei et al., 2021). Hypoxia causes a range of genetic, transcriptional, and metabolic adaptations in advanced tumors that ultimately promote survival and metastasis of cancer cells (Li et al., 2020). Thus, it has been recommended to assess anticancer effects of potent agents in hypoxic condition for accurate evaluation of their therapeutic potential (Nobre et al., 2018). Current findings that revealed considerable cytotoxicity of biogenic CuONPs in hypoxic condition suggest that these NPs have the potential to induce anticancer effects *in vivo*, although more research on animal models is required.

Carrying gene expression analysis indicated significant induction in P53 expression after administration of biogenic CuONPs. P53 is a tumor suppressor gene with critical roles in the cell cycle regulation and apoptosis (Vousden and Lu, 2002; Lacroix et al., 2006). Similar to our results, it has been reported that CuONPs induced apoptosis in human lung carcinoma cells through upregulation of P53 (Kalaiarasi et al., 2018). In the present study, CuONPs effectively downregulated the expression of BCL2, while induced the expression of BAX. Current findings are in agreement with previous studies, which demonstrated that CuONPs inhibited cell growth and induced apoptosis in acute myeloid leukemia, breast, gastric, colon and lung cancer cells via significant induction of BAX and downregulation of BCL2 (Shafagh et al., 2015; Gopinath et al., 2016; Khan et al., 2017; Kalaiarasi et al., 2018). Present results also revealed that CuONPs significantly decreased the expression of CCND1, a core cell cycle regulator that promotes cell proliferation and plays a major role in oncogenesis (Qie and Diehl, 2016). Similar to our findings, it has been shown that inhibited proliferation and induced apoptosis of oral carcinoma cells upon administration of metal oxide NPs were mediated by downregulation of CCND1 (Li et al., 2020).

Conclusion

Emergence of antibiotic-resistant strains is an intractable challenge to public health worldwide. In addition, acquired chemoresistance of cancer cells has vastly limited the clinical outcome of current pharmaceutical drugs. Although inorganic NPs could act as potent antibiotics and anticancer agents, disadvantages of conventional methods for NP synthesis have enforced to develop alternative approaches. In the present attempt, we successfully used *Stenotrophomonas* sp. BS95 to

synthesize functional CuONPs. UV vis spectroscopy, XRD and DLS analyses and electron microscopy revealed small size, crystallin nature and spherical shape of CuONPs. In addition, good dispersion and high stability of biogenic CuONPs were confirmed by zeta potential analysis, and functional groups were determined by FTIR spectroscopy. Evaluating biological effects of CuONPs exhibited their antibacterial activity against B. subtilis, S. aureus, P. putida, and E. coli. Furthermore, biogenic CuONPs possessed remarkable antioxidant potential and induced considerable anticancer effects on human colon and gastric adenocarcinoma cells via modulation of apoptosis-related genes. Interestingly, biogenic CuONPs induced low toxicity on normal cells, and had the potential to exert cytotoxic effects in hypoxic condition. According to the current findings, our biogenic CuONPs could be considered as effective agents with potential medical applications. Nevertheless, complementary studies on other pathogenic bacteria, more cell lines and animal models are required to better evaluate the efficacy and safety of biogenic CuONPs. In addition, conjugation of biogenic CuONPs with antibiotics and anticancer drugs might improve the clinical outcome of current therapeutic modalities.

Data availability statement

The original contributions presented in the study are included in the article/supplementary material, further inquiries can be directed to the corresponding author.

Author contributions

All authors participated in the design, interpretation and analysis of the data and review of the manuscript; ST, MN, and

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YA carried out the experiments and analysis of the results, BS and FBR designed and supervised the project, supplied all reagents, and edited the manuscript.

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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A peroxidase-like magneto-gold nanozyme AuNC@Fe₃O₄ with photothermal effect for induced cell apoptosis of hepatocellular carcinoma cells *in vitro*

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Hepatocellular carcinoma (HCC) is one of the most commonly diagnosed and malignant cancers worldwide. Conventional therapy strategies may not completely eradicate the tumor and may cause side effects during treatment. Nano-catalytic therapy, as a novel strategy, has attracted a great deal of attention. This study aimed to synthesize a multifunctional magneto-gold nanozyme AuNC@Fe₃O₄ and evaluate its anti-cancer potential in HepG2 cells in vitro. The characteristics of AuNC@Fe₃O₄ were assessed using a transmission electron microscope, dynamic light scattering, and energy-dispersive X-ray. The photothermal performance and peroxidase (POD)-like activity of AuNC@Fe₃O₄ were detected, using thermal camera and ultraviolet-visible spectrophotometer, respectively. The anti-cancer potential of AuNC@Fe₃O₄ was examined using cell counting kit-8, live/dead cell staining, and apoptosis analysis. Further research on HepG2 cells included the detection of intracellular reactive oxygen species (ROS) and lysosomal impairment. We observed that the AuNC@Fe₃O₄ had a small size, good photothermal conversion efficiency and high POD-like activity, and also inhibited cell proliferation and enhanced cell apoptotic ability in HepG2 cells. Furthermore, the AuNC@Fe₃O₄ enhanced ROS production and lysosomal impairment via the synergistic effect of photothermal and nano-catalytic therapies, which induced cell death or apoptosis. Thus, the magneto-gold nanozyme AuNC@Fe₃O₄ may offer a potential anti-cancer strategy for HCC.

KEYWORDS

POD-like, magneto-gold, nanozyme, photothermal effect, cell apoptosis, hepatocellular carcinoma

1 Introduction

Data from Global Cancer Incidence, Mortality and Prevalence 2020 revealed that liver cancer was the sixth commonly diagnosed and the third lethal cancer worldwide (Sung et al., 2021). China accounted for 23.7% and 30% of the global morbidity and mortality from liver cancer, respectively (Ferlay et al., 2021). It was predicted that between 2020 and 2040, there would be a 55% increase in the number of new cases of liver cancer per year, and the percentage of people who would die from the disease in 2040 would be more than 56.4% of those in 2020 (Rumgay et al., 2022). Primary liver cancer can be classified into three types: cholangiocarcinoma, hepatocellular carcinoma (HCC), and a combination of the two, with HCC accounting for approximately 90% of all cases (Llovet et al., 2016). HCC progression is

influenced by several risk factors, such as alcohol abuse, smoking, toxic chemicals, and hepatitis virus (especially for HBV) infections (Yang et al., 2019). Owing to the absolute number of HBV-infected populations (Liu et al., 2016), the mortality rate of HBV-related liver cancer was consistently higher than the global level (Liu et al., 2019), which increased the burden of HCC in China.

In most cases, conventional treatments, such as surgery, radiation, and chemotherapy, do not completely eradicate the tumor and may cause side effects during treatment, such as cancer palindromia and drug resistance (Zhu et al., 2016; Xu et al., 2019; Raoul and Edeline, 2020; Liu and Song, 2021). For example, the surgery was initially considered to be used for patients with early-stage HCC; However, over 50% of patients experienced a recurrence within a year following surgery (Gil et al., 2015; Weber et al., 2015). Sorafenib was an option for patients with advancedstage cancer, however, it was only effective in less than a third of them and caused drug tolerance or cytotoxicity (Llovet et al., 2008; Cheng et al., 2009; Bruix et al., 2012; Anwanwan et al., 2020). Besides surgery and chemotherapy, radiation therapy is a non-invasive and local ablative treatment approach to kill cancer cells. However, the efficiency of radiation therapy is easily limited by radioresistance, due to the DNA damage response and cell cycle checkpoints activation (Yoon and Seong, 2014; Wahl et al., 2016; Sun et al., 2020). Although the traditional strategies of HCC control the growth of HCC and prolong the survival time of patients, it still cannot satisfy their needs. Thus, it is necessary to discover a more efficient treatment approach to improve the quality of life for patients.

In recent years, nano-catalytic therapy, as a new tumor treatment strategy, has attracted the attention of an increasing number of researchers. Nanozymes are nanomaterials that catalyze chemical reactions of substrates under physiological states, obeying the patterns of enzyme kinetics (Wei et al., 2021). In 2007, Yan's team was the first to report that magnetic nanoparticles Fe₃O₄ possessed peroxidase (POD)-like activity, and proposed the concept of nano-catalysis (Gao et al., 2007). Furthermore, Shi et al. innovatively paved the way for further applications of nanoparticles in tumor nano-catalytic therapy, by disrupting the Fenton reaction that induced H₂O₂ disproportionation for •OH generation (Zhang et al., 2016). Currently, nano-catalytic therapy and photothermal therapy (PTT) are frequently employed in the treatment of tumors. The integration of PTT and nano-catalytic therapy has contributed to improving their cancer therapy efficiency. For instance, hyperthermia promoted the enzymatic activity of Fe₃O₄ nanozyme to generate more •OH, and simultaneously, •OH heightened the therapeutic impact of PTT (Wu et al., 2019; Zuo et al., 2022). It has also been reported that the Fe₃O₄@ZIF-8/GOx@ MnO₂ hybrid nanozyme can enhance the efficiency of nanoparticles in anti-tumor therapy by combining multiple therapeutics (Zhang et al., 2021b).

 Fe_3O_4 and Au nanoparticles, as is well known, demonstrated the unique characteristics of a high photothermal effect and POD-like activity (Zeng et al., 2013; Vallabani et al., 2017; Ghosh et al., 2022; Huang et al., 2022). Encouraged by the aforementioned description, we wonder if AuNC@Fe₃O₄ which has been employed as magnetic resonance imaging/com-puterized tomography multimodal imaging contrast agents of cancer owing to their high relaxivity value and excellent contrast enhancement (Wang et al., 2016b), also retains the photothermal and catalytic ability, or is beneficial to cancer therapy.

In this study, we synthesized multifunctional magneto-gold nanozyme AuNC@Fe₃O₄ and evaluated their anti-cancer ability in HCC cells *in vitro*. The AuNC@Fe₃O₄ exhibited high photothermal effect and POD-like activity. The results also reflected the influence of AuNC@Fe₃O₄ on engendering cell death and apoptosis. Furthermore, the synergistic effect of PTT and nano-catalytic therapy on reactive oxygen species (ROS) and lysosomal impairment in HepG2 cells were also studied.

2 Materials and methods

2.1 Materials and reagents

Ferric slat, gold (III) chloride (HAuCl4), other reagents related to AuNC@Fe₃O₄ synthesis and 3,3',5,5'-Tetramethylbenzidine (TMB) were purchased from Sigma, Inc. (St. Louis, United States). H₂O₂ solution and different pH buffer solutions (pH = 2, 3, 4, 5, 6, 7, 8, and 9) were bought from Aladdin (Shanghai, China). Human umbilical vein endothelial cells (HUVEC), human HCC cell lines (HepG2 cells) and the specific culture mediums for the two cell lines were purchased from Procell (Wuhan, China). Cell Counting Kit-8 (CCK-8) was obtained from Sangon Biotech (Shanghai, China). 2',7'-Dichlorodihydrofluorescein diacetate (DCFH-DA) was obtained from MedChemExpress (New Jersey, United States). Calcein-AM/propidium iodide (PI) kit, Annexin V-FITC apoptosis detection kit, Lyso-Tracker Red kit and Hoechst 33342 staining solution were purchased from Beyotime. Inc. (Shanghai, China).

2.2 AuNC@Fe₃O₄ synthesis

The AuNC@Fe₃O₄ was synthesized according to previous methods (Wang et al., 2016b). AuNC was initially synthesized and coated with poly (vinyl pyrrolidone) (PVP). Subsequently, PVP was replaced with 2-aminoethanethiol, and AuNC was transformed into AuNC-NH2 for interacting with carboxyl group functionalized Fe₃O₄ nanoparticles. The ultra-small Fe₃O₄ particles were prepared. To produce Fe₃O₄-COOH, ferric slats were vigorously stirred in pre-prepared polymer poly (acrylic acid) (PAA) solution. N-(3-Dimethylaminopropyl)-Nethylcarbodiimide and N-hydroxysuccinimide activated the Fe₃O₄-COOH, which then reacted with AuNC-NH₂ to generate AuNC@Fe₃O₄. The AuNC@Fe₃O₄ was centrifugated, washed with ethanol and water, and then dispersed in ddH₂O with different concentrations for further experiments.

2.3 AuNC@Fe₃O₄ characterization

The size of AuNC or AuNC@Fe₃O₄ nanoparticles was analyzed using a transmission electron microscope (TEM). Dynamic light scattering (DLS) was applied to detect hydrodynamic particle diameter and intensity of AuNC@Fe₃O₄ nanoparticles on a Malvern Zetasizer NANO ZS. Energy-dispersive X-ray (EDX) was utilized to analysis the element of AuNC@Fe₃O₄ nanoparticles on a FEI TECNAI G20 high-resolution TEM.

2.4 AuNC@Fe₃O₄ photothermal performance *in vitro*

To investigate the photothermal effect of the magneto-gold nanoparticles, First, 200 μ l of AuNC@Fe₃O₄ solution with distinct concentrations (0, 50, 100, 200, 300, 400, and 500 μ g/ml) was exposed to 808 nm laser at 1.0 W for 720 s; Second, 200 μ l of AuNC@Fe₃O₄ solution with concentration of 50 μ g/ml was exposed to 808 nm laser at different powers (1.0, 1.2, and 1.4 W) for 720 s. The thermal image and temperature change were recorded at different times by an infrared (IR) thermal camera (Fotric 220). As a control, ddH₂O was irradiated under the same conditions.

To investigate the photothermal stability of the magneto-gold nanoparticles, AuNC@Fe₃O₄ aqueous solution (500 µg/ml) was irradiated under 808 nm laser at 1.0 W for 420 s, then the irradiation was turned off. After that, the temperature was further measured for another 360 s. The experiment was then repeated four more times. The thermal image and temperature change were recorded at different times by the IR thermal camera (Fotric 220). As a control, ddH₂O was irradiated under the same operation.

To evaluate the photothermal conversion efficiency of AuNC@ Fe_3O_4 , the data from the cooling periods were calculated, according to previous report (Ren et al., 2015). Briefly, when the system reached energy balance, the equation was:

$$\sum_{i} m_i C_{p,i} \frac{dT}{dt} = Q_{AuNC@Fe_3O_4} + Q_s - Q_{loss}$$
(1)

where Cp and m were the heat capacity and mass of AuNC@Fe₃O₄ solution, respectively. T was the temperature of AuNC@Fe₃O₄ solution. $Q_{AuNC@Fe_3O_4}$ represented energy absorbed by AuNC@Fe₃O₄ nanoparticles. Q_s represented the energy absorbed by ddH₂O. Q_{loss} was the heat lost to the surroundings.

The equation for $Q_{AuNC@Fe_3O_4}$ was:

$$Q_{AuNC@Fe_{3}O_{4}} = I(1 - 10^{-A_{\lambda}})\eta$$
(2)

where *I* represented the laser power density, A_{λ} denoted the absorbance of AuNC@Fe₃O₄ solution under 808 nm in a 96-well plate, and η represented its photothermal conversion efficiency.

The equation for Q_{loss} was

$$Q_{loss} = hA\Delta T \tag{3}$$

where A was the surface area of the container, h denoted the heat transfer coefficient; ΔT represented the temperature changes, expressed as T-T_{surr} (where T and T_{surr} represent the solution and surrounding air temperature, respectively).

When heating ddH₂O, the heat input and output reached energy balance at the maximum steady-state temperature, therefore the equation for Q_s was:

$$Q_s = Q_{loss} = hA\Delta T_{max,H_2O} \tag{4}$$

where $\Delta T_{max,H_2O}$ was the temperature changes of ddH₂O.

When the system reached its maximum balanced temperature, the energy input (the heat absorbed by $AuNC@Fe_3O_4$ and ddH_2O) was equal to the heat lost into the surrounding, and the equation could be:

$$Q_{AuNC@Fe_{3}O_{4}} + Q_{s} = Q_{loss} = hA\Delta T_{max,mix}$$
(5)

where $\Delta T_{max,mix}$ was the changed temperature of the AuNC@Fe₃O₄ solution.

According to Eqs 2, 4, 5, η was:

$$\eta = \frac{hA\Delta T_{max,mix} - hA\Delta T_{max,H_2O}}{I(1 - 10^{-A_{\lambda}})} = \frac{hA(\Delta T_{max,mix} - \Delta T_{max,H_2O})}{I(1 - 10^{-A_{\lambda}})}$$
(6)

To calculate the unknown hA, θ was introduced, and could be expressed as following:

$$\theta = \frac{\Delta T}{\Delta T_{max}} \tag{7}$$

Adding Eq. 7 into Eq. 1, the new equation could be:

$$\frac{d\theta}{dt} = \frac{hA}{\sum_{i} m_{i}C_{p,i}} \left(\frac{Q_{AuNC@Fe_{3}O_{4}} + Q_{s}}{hA\Delta T_{max}} - \theta \right)$$
(8)

During the cooling period, the $Q_{AuNC@Fe_3O_4} + Q_s = 0$ in Eq. 8 was:

$$dt = -\frac{\sum_{i} m_{i} C_{p,i}}{hA} \frac{d\theta}{\theta}$$
(9)

which could be changed as following:

$$t = -\frac{\sum_{i} m_{i} C_{p,i}}{hA} ln \theta \tag{10}$$

where $\frac{\sum_{i} m_i C_{p_i}}{hA}$ was calculated by time versus $-\ln(\theta)$ plot. Since the mass of AuNC@Fe₃O₄ (1 × 10⁻⁷ kg) was relatively small when compared to that of ddH₂O (m = 2 × 10⁻⁴ kg), its m and Cp were neglected. The value of *hA* was then calculated using m_{H_2O} of 2 × 10⁻³ kg; C_{p,H_2O} of 4.2 × 10³ J/(Kg.°C). Furthermore, the η of AuNC@Fe₃O₄ was determined by substituting the value of *hA* and other parameters into Eq. 6. The values of other parameters were as follows: I = 2.3 W/cm², $A_{\lambda} = 0.105$, $\Delta T_{max,mix} = 25.3$, and $\Delta T_{max,H_2O} = 0.1$.

2.5 POD-like activity assay

To evaluate the catalytic properties of AuNC@Fe₃O₄, AuNC@ Fe₃O₄ (final concentration: 0, 5, 10, 20, 50, and 100 µg/ml), TMB (final concentration: 0.4 mM), and H₂O₂ (final concentration: 50 µM) was added into a final volume of 500 µl of phosphatebuffered saline (PBS) solution. The absorbance of the buffer was measured using an ultraviolet-visible (UV-vis) spectrophotometer at 500–800 nm. The POD-like activity assay of AuNC@Fe₃O₄ at varying pH levels (pH = 2, 3, 4, 5, 6, 7, 8, and 9) was performed in the presence of H₂O₂ and TMB in PBS solution, and the absorbance at 652 nm was detected by a microplate reader.

2.6 POD-like catalytic kinetic determination

When TMB was used as a substrate, the AuNC@Fe₃O₄ (final concentration: 50 μ g/mL), TMB (final concentration: 0.0, 0.2, 0.4, 0.6, and 0.8 mM), and H₂O₂ (final concentration: 50 μ M) was added into a final volume of 100 μ l of PBS solution. The absorbance at 652 nm was detected by a microplate reader.

When H_2O_2 was used as a substrate, the AuNC@Fe₃O₄ (final concentration: 50 µg/ml), H_2O_2 (final concentration: 0, 10, 20, 30, 40, 50, 60, 70, and 80 µM) and TMB (final concentration: 0.4 mM) was added into a final volume of 100 µl of PBS solution. The absorbance at 652 nm was detected by a microplate reader.

Based on Michaelis-Menten Eq. 11 and saturation curve, the $V_{\rm max}$ and Michaelis-Menten constant could be calculated,

$$\frac{1}{V} = \frac{K_m}{V_{max}} \frac{1}{[s]} + \frac{1}{V_{max}}$$
(11)

and the V was calculated using Eq. 12:

$$V = \frac{A}{(b \times \mathbf{\mathcal{E}}_{652\,nm} \times t)} \tag{12}$$

where *A* was the absorbance of the reaction system at 652 nm. t = 600 s, which was the reaction time. b = 0.3125 cm, which was the light path in the reaction solution, and $\mathcal{E}_{652\,nm}$ = 39,000 M⁻¹ cm⁻¹ (Dashtestani et al., 2019).

2.7 Cell viability assay

The HepG2 and the HUVEC cells were cultured to assess the cytotoxicity of AuNC@Fe₃O₄ through CCK-8 assay. 4000 of cells were cultured at 96-well plate well overnight at 37°C in a humidified incubator with 5% CO₂. Subsequently, 100 μ l of fresh medium with distinct concentrations of AuNC@Fe₃O₄ (0, 10, 20, 30, 40, and 50 μ g/ml) was changed and cultured for 24 h. The CCK-8 solution (final volume: 10 μ l) was added into 100 μ l of medium, and incubated for 2 h. Then, the absorbance of medium was detected at 450 nm.

2.8 Live/dead cell staining assay

HepG2 cells were cultured overnight in a 12-well plate with 500 μ l of culture medium. The cells were then treated with PBS or AuNC@Fe₃O₄ (50 μ g/ml) for 12 h. Then, the cells were cultured for an additional 12 h after either being irradiated by an 808 nm laser for 5 min at 1.4 W or not. The culture medium was then removed, and cells were washed once with PBS and incubated with 500 μ l stain solution for 15 min. Finally, the cells were washed thrice with PBS and photographed by an inverted fluorescence microscope.

2.9 Apoptosis analysis

To investigate the ability of AuNC@Fe₃O₄ for inducing cell apoptosis, HepG2 cells were quantitatively detected by a flow cytometer. The cells were initially seeded into a 6-well plate and treated under different conditions for 24 h. They were collected with 0.25% trypsin and washed thrice with ice-cold PBS. Subsequently, these cells were resuspended in 195 μ l of binding buffer. Ten microliters of PI and 5 μ l of Annexin V-FITC were added, and the mixture was incubated for 20 min at room temperature, and cells were detected by flow cytometer.

2.10 Intracellular ROS detection

The intracellular POD-like catalytic ability of AuNC@Fe₃O₄ was detected using DCFH-DA. Except for an additional 4 h of culture, the method used for the laser-irradiated groups was similar to the treatment described above. Furthermore, 1 ml of PBS with DCFH-DA (5 μ M) was added, and the mixture was incubated for another 30 min at 37°C in a humidified incubator with 5% CO₂. The wells were then washed thrice with PBS to remove the excess dye and photographed by an inverted fluorescence microscope.

2.11 Lysosomal impairment assay

After treatment, lysosomes and cell nuclei were stained with Lyso-Tracker Red and Hoechst 33342, respectively, according to the manufacturer's instructions. Subsequently, an inverted fluorescence microscope was used to capture images of cells.

2.12 Statistical analysis

Statistical analysis was achieved by GraphPad Prism version 8 (GraphPad Software, United States). Results were represented as mean \pm standard deviation. The student t-test was used to compare the means of multiple groups. The statistical significances were as follows: * 0.01 < p < 0.05, ** 0.001 < p < 0.01, and ***p < 0.001.

3 Results and discussion

3.1 Synthesis and characterization of AuNC@ Fe $_3O_4$

The structure and characteristics of AuNC and AuNC@Fe₃O₄ were determined by TEM. The results demonstrated that the diameter of AuNC and AuNC@Fe3O4 were 25-40 and 50-100 nm, respectively, with high uniformity and no agglomeration (Figures 1A, B). DLS was used to confirm the size of AuNC@Fe₃O₄, and the average hydrodynamic size distribution of these nanoparticles was approximately 55 nm (Figure 1C). The increase in the hydrodynamic size might be owing to the attachment of Fe₃O₄ to the surface of the AuNC. Elemental mapping analysis revealed the presence of the atoms Au, Fe and O, proving that AuNC@Fe₃O₄ was successfully formed (Figure 1D; Table 1). The "-CO-NH-", that came from the reaction of Fe_3O_4 -COOH and AuNC-NH₂ and the carbon-coated brace used during sample preparation or analysis might have contributed to the existence of C element that was also present (Phongtongpasuk et al., 2016).

3.2 Photothermal performance of AuNC@ Fe_3O_4

The thermal camera was used to investigate the photothermal conversion capabilities of AuNC@Fe₃O₄. The temperature changes of AuNC@Fe₃O₄ solution with different concentrations under



TABLE 1 The statistics of elements analysis for AuNC ${}_{\odot}\text{Fe}_{3}O_{4}$ by energy-dispersive X-ray (EDX).

Compound	Element	Weight (%)
AuNC@Fe ₃ O ₄	Au	23.1
	Fe	29.4
	0	42.4
	С	5.1

808 nm laser irradiation at 1.0 W for 360 s were recorded. As depicted in Figure 2A, the temperature of the solution increased in a concentration- and time-dependent pattern. For example, the temperature of different concentrations of AuNC@Fe₃O₄ solution reached steady statue at 8 min. The temperature of AuNC@Fe₃O₄ solution (500 µg/ml) was changed significantly from 25.9°C to 52.3°C compared with the neglected increase in that of ddH₂O (from 26.0°C to 26.5°C), indicating the good photothermal response of AuNC@Fe₃O₄. For further investigation, the AuNC@Fe₃O₄ solution (50 µg/ml) was irradiation at different powers (1.0, 1.2, and 1.4 W). The laser power was increased from 1.0 to 1.4 W, which resulted in a significant increase in the temperature of the AuNC@ Fe₃O₄ solution. A temperature of 45.3°C was achieved after 10 min of 808 nm laser irradiation at 1.4 W (Figure 2B). PTT, a promising cancer treatment strategy, converts light energy into heat to generate an area of hyperthermia, where tissues can be exposed to high temperatures (from 42°C to 45°C), which can damage or kill tumor cells (Tchouagué et al., 2019; Qu et al., 2022). The results of Figures 2A, B suggested a potential application of AuNC@Fe₃O₄ in antitumor.

Additionally, five cycles of the "On and Off" model were used to measure the temperature curve of the AuNC@Fe₃O₄ solution to assess its photothermal stability. The AuNC@Fe₃O₄ showed excellent photothermal stability since the temperature was raised to 52.9 °C and there was no reduction in the temperature rise following laser irradiation during the five cycles (Figure 2D).

Moreover, the average of the data from the five cooling periods was used to get the photothermal conversion efficiency (η) of AuNC@Fe₃O₄. The plot of the time value and $-\ln$ (θ) was displayed in Figure 2E, and its slope was 99.526. Using Eqs 6, 10, the η of AuNC@Fe₃O₄ was calculated to be 39.58%, which was similar with or higher than the PPT reagents previously reported, such as, EA-Fe@BSA NPs (31.2%) (Tian et al., 2020), Fe₃O₄@ Carbon@Platinum-Chlorin e6 (28.28%) (Xu et al., 2022b), Au nanorods (22%) (Zeng et al., 2013), Au nanoshells (13%) (Hessel et al., 2011), PANi@Au (40.4%) and Au nanoparticles (21.7%) (Zhang et al., 2021a).

Collectively, these findings suggested that $AuNC@Fe_3O_4$ exhibited good photothermal conversion and photothermal stability, which implied a promising application in PTT for tumors.

3.3 POD-like activity of AuNC@Fe₃O₄

It was reported that Au and Fe₃O₄ nanoparticles demonstrated POD-like enzyme activity (Zandieh and Liu, 2021), therefore it was necessary to investigate whether the AuNC@Fe₃O₄ possessed similar characteristics. The peroxidase mimicking activity of AuNC@Fe₃O₄ was validated by TMB. TMB could be oxidized to blue oxTMB by \bullet OH and detected at 652 nm, using UV-vis spectrophotometer (Zhu et al., 2022). As presented in Figure 3A,



Photothermal performance analysis of AuNC@Fe₃O₄. (A) Temperature change curves of water and AuNC@Fe₃O₄ aqueous solution after different treatments (B) Temperature change curves of AuNC@Fe₃O₄ after different treatments (C) Photostability of AuNC@Fe₃O₄ solution under irradiation for five cycles. (D) Time versus $-ln(\theta)$ plot of the AuNC@Fe₃O₄ solution.



POD-like activity assay of AuNC@Fe₃O₄. (A) Ultraviolet-visible (UV-vis) absorption spectra of the reaction system with different concentrations. (B) Michaelis-Menten curve of AuNC@Fe₃O₄ for H₂O₂. (C) Lineweaver-Burk plotting of AuNC@Fe₃O₄ for H₂O₂. (D) Michaelis-Menten curve of AuNC@Fe₃O₄ for TMB. (E) Lineweaver-Burk plotting of AuNC@Fe₃O₄ for TMB. (F) The absorbance of the reaction system at 652 nm under different pH values.

the groups with different concentrations of AuNC@Fe₃O₄ had varying absorbance intensities at 652 nm. The group of AuNC@ Fe₃O₄ with 100 μ g/ml showed the strongest signal at 652 nm,

followed by the group with 50 $\mu g/ml$. The intensity of absorbance tested with H₂O displayed no peak at 652 nm. These findings, which indicated that the AuNC@Fe₃O₄ possessed POD-like enzyme

activity, were further verified by the inset digital photos (Figure 3A). To further confirm the POD-like enzyme specificity of AuNC@ Fe₃O₄, the UV-vis absorption spectra of the reaction system with varying conditions was collected. It was observed from Supplementary Figure S1 that absorbance peak of the AuNC@ Fe₃O₄+TMB or AuNC@Fe₃O₄+H₂O₂ group was negligible. In the absence of AuNC@Fe₃O₄, the TMB + H₂O₂, TMB or H₂O₂ group showed no significant absorbance peak at 652 nm, which was consistent with the AuNC@Fe₃O₄, only group. The results suggested that, except AuNC@Fe₃O₄, other components in the reaction system could hardly catalyzed the conversion of H₂O₂ to •OH and oxidized TMB, which indicated AuNC@Fe₃O₄ exhibited a specific activity of POD-like enzyme.

Kinetic parameters were analyzed to quantitate the POD-like activity of AuNC@Fe₃O₄ using the initial rate method (Gao et al., 2017). First, the absorbance of the system was measured, while varying the concentrations of H₂O₂ concentrations from 0 to 80 mM and maintaining a TMB concentration of 0.4 mM (Supplementary Figure S2). Second, the velocity of reaction was calculated according Eq. 12 and the plot was consistent with traditional Michaelis-Menten curve (Figure 3B), which demonstrates that the catalytic reaction rate increased with the growth of substrate concentration and achieved steady state at high concentrations (Huang et al., 2022). Third, after Lineweaver-Burk fitting, the enzyme kinetic parameters, such as Michaelis-Menten constants (Km) was calculated to be 47.65 mM and the maximum reaction velocity (V_{max}) was 3.18 \times 10⁻⁷ M s⁻¹ (Figure 3C). Forth, the absorbance of the solution was measured at 652 nm while varying TMB concentrations and maintaining H2O2 concentration as a constant (Supplementary Figure S3). Last, the K_m and V_{max} were 0.25 mM and 9.03 \times $10^{^{-8}}\,M\,s^{^{-1}}$ respectively, and the results were presented in Figures 3D, E.

When the H₂O₂ was used as substrate, the velocity of AuNC@ Fe₃O₄ was faster than that of Fe₃O₄ (Vallabani et al., 2017), and K_m value that was lower than that of Fe₃O₄ (Vallabani et al., 2017). Similarly, when the TMB was used as substrate, AuNC@Fe₃O₄ had a velocity that was faster than that of Au NRT, Au NC, Au NS, and horseradish peroxidase (Ghosh et al., 2022), and its value of K_m was also lower than those of them. In the catalytic reaction system, the K_m represents the affinity between the enzyme and substrates, and the lower the K_m, the higher enzyme affinity (Jiang et al., 2018). Therefore, the results suggested that the catalytic ability and the affinity between AuNC@Fe₃O₄ nanozyme and substrates (such as TMB and H_2O_2) was stronger than that of Fe_3O_4 and Au nanoparticles. The following factors may contribute to the significant increase in POD-like activity of AuNC@Fe3O4 nanoparticles: the electronic structure of the interfaces between the Fe₃O₄ and Au, the synergistic effect, and polarization effects from Au to Fe₃O₄ (Lee et al., 2010; Sun et al., 2013; Wang et al., 2016a).

Considering the complex tumor microenvironment, such as hypoxia and weak acidity (Li et al., 2020; Zhao et al., 2021), it was unclear whether AuNC@Fe₃O₄ exhibits POD-like enzyme activity even at low pH. At low pH values ranging from 2 to 6, the AuNC@Fe₃O₄ exhibited higher POD-like enzyme activity, and the optional pH was 4. When the pH was higher than 7, the PODlike enzyme activity was reduced dramatically (Figure 3F). The results hinted that AuNC@Fe₃O₄ might have significantly varied POD-like enzyme activity between distinct parts of normal (pH = 7.4) and cancer tissues (pH = 6.5), especially for lysosomes (pH = 4.5-5.5) and endosomes (pH = 5.5-6.8) (Kuppusamy et al., 2002; Wojtkowiak et al., 2011).

Overall, these findings provided evidence for the high POD-like catalytic activity of AuNC@Fe₃O₄ nanozyme and implied potential catalytic ability in tumor.

3.4 In vitro anti-tumor effect of AuNC@ Fe_3O_4

It is important to examine the biocompatibility of AuNC@Fe₃O₄ before performing further clinical applications. Therefore, HepG2 and HUVEC cells were incubated with AuNC@Fe3O4 at varying concentrations for 24 h to estimate the cytotoxicity using CCK-8 assay. Low concentrations of AuNC@Fe₃O₄ did not affect the survival rate of HepG2 cells; however, at 40 and 50 μ g/ml, the viability of cells decreased to 77% and 60%, respectively (Figure 4A). In contrast, the viability of HUVEC cells was not drastically affected by AuNC@Fe₃O₄ after 24 h incubation at the varying treatments (Figure 4B). The findings indicated that AuNC@Fe₃O₄ was not toxic to normal cells at the concentration ranging from 0 µg/ml to 50 µg/ ml and demonstrated good biocompatibility. The reason why AuNC@Fe₃O₄ showed more sensitive to HepG2 could be attributed to the fact that the pH of the tumor was lower than that of normal tissues (Kuppusamy et al., 2002; Wojtkowiak et al., 2011) and that the AuNC@Fe₃O₄ had higher POD-like enzyme activity in a lower pH reaction system, which meant it produced more •OH, which could be lethal to cells (Cui et al., 2018; Malfanti et al., 2022).

To explore the anti-tumor effect of AuNC@Fe₃O₄, the live/dead cell staining assay was utilized. There were nearly no dead cells in the PBS and PBS + NIR (near-infrared) groups; However, when the cells were treated with AuNC@Fe₃O₄ or AuNC@Fe₃O₄+NIR, the number of dead cells increased significantly, with the last group having the most cell death (Figure 4C). To further verify this result, the flow apoptosis assays of HepG2 cells with different conditions was conducted. As depicted in Figures 4D, E; Supplementary Table S1, the results indicated that approximately 51% apoptotic cells (Q2+Q3) were observed in the AuNC@Fe₃O₄+NIR group, which was greater than other groups.

The results revealed that AuNC@Fe $_3O_4$ displayed good biocompatibility, and the laser irradiation augmented the anti-tumor ability of AuNC@Fe $_3O_4$.

3.5 ROS and lysosomal impairment induced by AuNC@Fe $_3O_4$

To confirm the synergistic effect of PTT and POD-like enzyme catalytic activity of AuNC@Fe₃O₄, the production of the ROS in HepG2 cells was validated using the DCFH-DA probe. As reported, DCFH-DA crossed the cell membrane and was subsequently oxidized to DCF with green fluorescence (Afri et al., 2004). It was evident from Figure 5A that the HepG2 cells incubated with AuNC@Fe₃O₄ exhibited a higher green fluorescence signal than PBS and PBS + NIR, indicating the ability of AuNC@Fe₃O₄ to effectively



Evaluation for anti-tumor effect of AuNC@Fe₃O₄ *in vitro*. (A) Cell viability of HepG2 cells treated with AuNC@Fe₃O₄ for 24 h. (B) Cell viability of human umbilical vein endothelial cells (HUVEC) cells treated with AuNC@Fe₃O₄ for 24 h. (C) Calcein-Am/propidium iodide (PI) staining of HepG2 cells under different conditions. Scale bar: 100 μ m. (D) Apoptosis analysis of HepG2 cells with different treatments. (E) The histogram results of apoptotic HepG2 cells derived from (D). * 0.01 < p < 0.05, ** 0.001 < p < 0.01, and ***p < 0.001.



FIGURE 5

Analysis for synergistic effect of photothermal therapy (PTT) and catalytic activity of AuNC@Fe₃O₄. (A) Reactive oxygen species (ROS) detection of HepG2 cells with varying treatments. (B) Lysosomal impairment detection of HepG2 cells with varying treatments. Scale bar: $50 \mu m$.

catalyze the conversion of intracellular H_2O_2 into •OH in cancer cells. Compared with the AuNC@Fe₃O₄ group, the signal of the AuNC@Fe₃O₄+NIR group was stronger. The similar result was collected by the detection of the absorbance intensities at 652 nm of the reaction system with or without NIR irradiation, using UV-vis spectrophotometer. We found the signal of reaction system with NIR irradiation was higher than that of group without irradiation (Supplementary Figure S4). The results confirmed that photothermal effect enhanced the POD-like enzyme catalytic activity of AuNC@Fe₃O₄.

The phenomenon could be attributed to the localized surface plasmon resonance (LSPR), which was the collective oscillation of surface free electrons in metal nanoparticles under light irradiation, leading to local heating (also called photothermal effect) and hot carriers (such as hot electrons and hot holes). One hand, energy of hot electrons may transfer to local heating by electron-phonon interactions, causing a rise in temperature (Brongersma et al., 2015). Similar with natural enzymes, the catalytic ability of nanozymes could be enhanced by elevated temperature (Wang et al., 2021; Zhu et al., 2022). Another hand, hot electrons could be transferred from AuNPs to empty orbits of H_2O_2 , and activated the H_2O_2 to generate •OH under NIR light irradiation (Wang et al., 2017; Xu et al., 2022a).

It was reported that increased ROS could disrupt normal structure of the lysosomes (Shyam et al., 2021); however, whether the AuNC@Fe₃O₄ could induce lysosomal impairment remained unknown. The fluorescence images (Figure 5B) demonstrate that the PBS alone and PBS + NIR groups had negligible effects on the lysosomal impairment and that there were more HepG2 cells with lysosomal impairment following incubation with AuNC@Fe₃O₄. As expected, the lysosomal signal was the weakest in the AuNC@Fe₃O₄ under laser irradiation group. The results confirmed the synergistic effect of PTT and POD-like enzyme catalytic activity of AuNC@ Fe₃O₄ on lysosomal impairment. Additionally, lysosomal impairment may contribute to an increase in lysosomal membrane permeability, a decrease in lysosomal quantity, a disruption in lysosomal enzyme activities, an increase in ROS levels, and most importantly, the induction of cell apoptosis (Abulikemu et al., 2022).

This at least partly, explained why $AuNC@Fe_3O_4$ with or without laser irradiation could cause cell death or apoptosis.

4 Conclusion

In summary, this study aimed to synthesize magneto-gold nanozyme AuNC@Fe₃O₄ and evaluate its anti-cancer effects for HCC *in vitro*. The AuNC@Fe₃O₄ showed the typical small size of about 55 nm. Additionally, it demonstrated a high photothermal conversion efficiency and POD-like activity. The CCK-8 results demonstrated that AuNC@Fe₃O₄ had good biocompatibility and HCC cell-killing ability. Moreover, AuNC@Fe₃O₄ could synergistically stimulate cell death or apoptosis. Finally, it was observed that magneto-gold nanocomposites could facilitate 808 nm laser irradiation to increase their catalytic ability to produce ROS. This might promote lysosomal impairment, causing cell death or apoptosis. These results suggested that the AuNC@Fe₃O₄ may offer a promising anti-cancer strategy for HCC *via* the synergistic effect of PTT and nano-catalytic therapy. Further research is required to investigate the therapeutic efficacy of AuNC@ Fe₃O₄ for HCC *in vivo*.

Data availability statement

The original contributions presented in the study are included in the article/Supplementary Material, further inquiries can be directed to the corresponding author.

Author contributions

XS and GW proposed the conception and design of the study. XS and JL performed the experiments and interpretation of the data. XS and JL drafted the manuscript. GW reviewed and revised the manuscript. All authors read and approve the final manuscript.

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Supplementary material

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fbioe.2023.1168750/ full#supplementary-material

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Fullerenol inhibits tendinopathy by alleviating inflammation

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Tendinopathy is a common disease in orthopaedics, seriously affecting tendon functions. However, the effects of non-surgical treatment on tendinopathy are not satisfactory and surgical treatments possibly impair the function of tendons. Biomaterial fullerenol has been proved to show good anti-inflammatory effects on various inflammatory diseases. For in vitro experiments, primary rat tendon cells (TCs) were treated by interleukin-1 beta (IL-1 β) combined with aqueous fullerenol (5, 1, 0.3 µg/mL). Then inflammatory factors, tendon-related markers, migration and signaling pathways were detected. For in vivo experiments, rat tendinopathy model was constructed by local injection of collagenase into Achilles tendons of rats and fullerenol (0.5, 1 mg/mL) was locally injected 7 days after collagenase injection. Inflammatory factors and tendon-related markers were also investigated. Fullerenol with good water-solubility showed excellent biocompatibility with TCs. Fullerenol could increase expression of tendonrelated factors (Collagen I and tenascin C) and decrease expression of inflammatory factors (matrix metalloproteinases-3, MMP-3, and MMP-13) and reactive oxygen species (ROS) level. Simultaneously, fullerenol slowed the migration of TCs and inhibited activation of Mitogen-activated protein kinase (MAPK) signaling pathway. Fullerenol also attenuated tendinopathy in vivo, including reduction of fiber disorders, decrease of inflammatory factors and increase of tendon markers. In summary, fullerenol is a promising biomaterial that can be used to treat tendinopathy.

KEYWORDS

fullerenol, inflammation, ROS, MAPK, tendinopathy

1 Introduction

Tendinopathy is chronic disorders of tendons usually caused by overuse, with an incidence of 0.2%–0.3% of adult patients (van der Vlist et al., 2020). Among them, athletes are the riskiest ones to suffer from tendinopathy with a morbidity of approximate 52% (Kujala et al., 2005) (Lagas et al., 2020). Tendinopathy causes pain, diffuse or localized swelling, loss of tissue integrity and impaired performance (Millar et al., 2021). The pathological mechanisms of tendinopathy are multiple, including apoptosis disorder, mechanical overload, imbalance of matrix metalloproteinases (MMPs) and tissue inhibitors of metalloproteinases (TIMPs), genetic factors, inflammation (Yuan et al., 2002) (Arnoczky et al., 2004) (Mokone et al., 2006). Current managements of tendinopathy consist of drug treatments, physical therapy, and surgery. However, curative effects of drug treatments and physical therapies are short-term. Surgeries possibly lead to secondary injury and tendon function postoperatively cannot recover to preoperative level. Thus, it is necessary to develop a new treatment method with minor injury.

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Pathology of tendinopathy still remains controversial. Inflammation plays a key role in the appearance of tendinopathy, especially in the early phase (Legerlotz et al., 2012). From the perspective of risk factors, injuries, repetitive mechanical overloading and hypoxia all elevate inflammatory cytokines, such as tumor necrosis factor alpha (TNF- α), interleukin-1 beta (IL-1 β), prostaglandin E2 (PGE2) (D'Addona et al., 2017). Moreover, hypoxic damage or increased oxygen demand of tendon cells caused by mechanical stresses also tends to raise oxygen free radicals like reactive oxygen species (ROS), leading to secondary damage of tendon tissues. Simultaneously, Dakin et al. found that both tendinopathic and ruptured Achilles tendons of human expressed many CD14⁺ and CD68⁺ cells and showed a complex inflammation signature, involving interferon, nuclear factor-kappa B (NF-κB) and signal transducer and activator of transcription 6 (STAT-6) activation pathways, which also proved that inflammation was a vital pathological process of tendinopathy (Dakin et al., 2018). Therefore, inhibiting inflammation is possibly an effective method to attenuate tendinopathy.

Fullerenol is a fullerene derivative with good water solubility, which expands its use in biological and medical fields. Structurally, there are numerous carbon-carbon double bonds in fullerenol, contributing to its antioxidative activity of scavenging reactive oxygen species (ROS) (Markelić et al., 2022). A lot of studies have reported good protective effects of fullerenol on cells under oxidative stress and DNA damage. It was found that fullerenol showed excellent curative or preventive effects on bleomycin-induced pulmonary fibrosis (Zhou et al., 2018), intervertebral disk degeneration (Yang et al., 2014a), myocardial ischemia-reperfusion injury (Ding and Li, 2020), osteoarthritis (Pei et al., 2019). In the meantime, fullerenol has the ability to rescue HaCaT human skin keratinocytes and corneal epithelial cells from ultraviolet B (wavelength between 280 and 320 nm) (Saitoh et al., 2011) (Chen et al., 2022). Apart from these, fullerenol presents promising results of osteogenic differentiation induction to repair bone defect. Despite of the favorable therapeutic benefits of fullerenol on multiple diseases, there is no study on the effects of fullerenol on tendinopathy.

Since inflammation was an important feature of tendinopathy, and fullerenol showed brilliant anti-inflammatory and antioxidant effects in various diseases, we hypothesized that fullerenol could mitigate tendinopathy by inhibiting inflammation. Therefore, this study aims to explore the effects of fullerenol on tendinopathy and investigate the potential mechanisms, in order to provide a new treatment method of tendinopathy.

2 Materials and methods

2.1 Characterization of fullerenol

Fullerenol powder was purchased from Chengdu Zhongke Times Nano Energy Tech Co., Ltd. The fullerenol powder was tested by Transmission Electron Microscope (TEM, TF20) for size and morphology and by Fourier transform infrared spectrometer (FTIR, Thermo Scientific Nicolet iS20, United States). At room temperature, fullerenol was suspended in distilled water to make aqueous fullerenol with the concentration of 50 mg/mL. Then, size distribution and zeta potential were investigated by Nano Sizer and Zeta potential Tester (Omni, United States). 50 mg/mL aqueous fullerenol was stored at room temperature shielded from light for further use. For cell treatment, aqueous fullerenol was diluted with Dulbecco's modified Eagle's medium (DMEM, Gibco, United States) to the concentration of 10, 5, 3, 1, 0.5, 0.3, 0.1 μ g/mL and was sterilized with 0.22 μ m filter membranes (Millipore, United States).

2.2 Tendon cells isolation and culture

Tendon cells (TCs) were isolated from the Achilles tendons of rats, as described previously (Jiao et al., 2022b). In brief, Achilles tendons of one-week-old rats were cut after disinfection. And tendons were immersed in 0.06% collagenase type I (Worthington, United States) solution at 37°C overnight. Then, the solution was centrifuged and the supernatant was discarded. The sediment was suspended and incubated in DMEM supplemented with 10% fetal bovine serum (FBS, Gibco, United States) and 1% antibiotics (penicillin and streptomycin, Gibco, United States). Cells were subcultured when they reached 80%–90% confluence.

2.3 Cell Count Kit-8 assay

Cell Count Kit-8 (CCK-8) assay was performed using the kit (Dojindo, CK04-05, Japan) according to the instruction. TCs were seeded into 96-well plates with a density of 3×10^3 per well. Then, TCs were incubated using DMEM containing fullerenol with different concentrations (10, 5, 3, 1, 0.5, 0.3, 0.1 µg/mL). At 1 and 3 days, TCs were cultured in DMEM medium with 10% CCK-8 reagent at 37°C for 2 h. The absorbance of the supernatant at 450 nm was measured using a microplate reader (Infinite M200 Pro, Tecan, Switzerland).

2.4 Live/dead cell staining

Live/dead cell staining was performed using the kit (KeyGEN, Nanjing, China) according to the manufacturer's instruction. The samples were observed using a confocal microscope (Leica, Germany). Live (green) cells stained by Calcein AM were detected with excitation at 488 nm, and dead (red) cells stained by PI were observed with excitation at 555 nm.

2.5 RNA extraction and qRT-PCR

For inflammation induction and fullerenol treatment, TCs were treated by 50 ng/mL IL-1 β combined with aqueous fullerenol. Then, total RNA was extracted using TRIzol reagent (Thermo Scientific, United States). A NanoDrop 1,000 spectrophotometer (Thermo Scientific, United States) was used to evaluate RNA purity and quantification. 1,000 ng of the extracted RNA was reverse transcribed to cDNA using

Primer	Forward primer (5' to $3'$)	Reverse primer (5' to 3')
GAPDH	GGCAAGTTCAACGGCACAGT	GCCAGTAGACTCCACGACAT
COL1A1	TGACTGGAAGAGCGGAGAGTA	GGGGTTTGGGCTGATGTACC
TNC	TGCCATAGCAACAACAGCCAT	AACTCTCCACCTGAGCAGTC
MMP-3	TGCTCATGAACTTGGCCACT	GTGGGAGGTCCATAGAGGGAT
MMP-13	GGGAACCACGTGTGGAGTTAT	GACAGCATCTACTTTGTCGCC

TABLE 1 Sequences of primers for gRT-PCR.

Abbreviations: GAPDH, glyceraldehyde-3-phosphate dehydrogenase; COL1A1, collagen 1A1; TNC, tenascin C; MMP-3, matrix metalloproteinases-3; MMP-13, matrix metalloproteinases-13.

PrimeScript Master Mix (Takara, RR036A, Japan). The qRT-PCR reaction was performed with 2× SYBR Green qPCR Master Mix (Low ROX) (Bimake, B21703, China) and Applied Biosystems 7,500 Real-Time PCR System (Applied Biosystems, Foster City, CA, United States). The relative mRNA levels were calculated with $2^{-\Delta\Delta CT}$ method. Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) was used as internal control. The primers used in this study are listed in Table 1.

2.6 Scratch assay

For scratch assay, TCs were seeded into 6-well plates and cultured to reach 80% confluence. Then, a straight scratch was scraped with a 200- μ L pipette tip. And TCs were incubated with IL-1 β and aqueous fullerenol. Cell migration was determined by measuring the distance at 0, 12 and 24 h.

2.7 Transwell assay

For the Transwell assay, Transwell chambers (BD Science, United States of America) were used. In 24-well plate, TCs (5 \times 10⁴) in 150 μ L of serum-free basal medium were seeded into the upper chamber, and 650 μ L of DMEM supplemented with 10% FBS, IL-1 β and aqueous fullerenol was added into the lower chamber. The Transwell system was placed in a 5% CO₂ incubator at 37 C for 24 h. Then, the cells were fixed and stained with crystal violet solution.

2.8 Protein extraction and Western blotting

For inflammation induction and fullerenol treatment, TCs were treated by 50 ng/mL IL-1 β combined with aqueous fullerenol. For protein extraction, TCs were lysed using RIPA lysis buffer (Beyotime, China) supplemented with 1% protease and phosphatase inhibitor cocktail (100X) (Thermo Scientific, United States). Then, the mixture was centrifugated at a speed of 14,000 RCF for 15 min. The supernatant was separated and mixed with SDS-PAGE sample loading buffer (Beyotime, China) and boiled at 99°C for 5 min. Protein samples were electrophoresed on SDS gels and transferred onto polyvinylidene fluoride membranes (Millipore, United States). The membrane was then blocked in Tris-buffered saline Tween

20 (Solarbio, China) containing 5% non-fat milk (Sangon Biotech, China) or 5% bovine serum albumin (MPbio, United States) for 1 h at room temperature. After that, the membrane was incubated with primary antibodies at 4°C overnight and secondary antibodies for 1 h at room temperature. Protein immunoreactivity was detected with LI-COR Odyssey Fluorescence Imaging System (LI-COR Biosciences, United States), and ImageJ was used to measure the protein expression. The anti-bodies used were as follows: p44/42 MAPK (Erk1/2) (Cell Signaling Technology, United States), Phospho-p44/42 MAPK (Erk1/2) (Cell Signaling Technology, United States), SAPK/JNK (Cell Signaling Technology, United States), Phospho-SAPK/JNK (Cell Signaling Technology, United States), p38 MAPK (Cell Signaling Technology, United States), Phospho-p38 MAPK (Cell Signaling Technology, United States), Anti-rabbit IgG (H + L) (800 4X PEG Conjugate) (Cell Signaling Technology, United States), Anti-mouse IgG (H + L) (800 4X PEG Conjugate) (Cell Signaling Technology, United States).

2.9 Animal experiments

All animal experiments were approved by the Ethics Committee of Shanghai Ninth People's Hospital, Shanghai Jiaotong University School of Medicine (Approval number: SH9H-2021-A895-1). To establish tendinopathy models, Sprague-Dawley (SD) rats (male, 8 weeks old), purchased from Shanghai JieSiJie Laboratory Animals Co., LTD., were anesthetized. Then, 50 mg/mL collagenase type I (Worthington, United States) solution was injected into the Achilles tendons to trigger inflammation. At 7 days after injection, 50 µL aqueous fullerenol with the concentration of 0.5 mg/mL and 1 mg/mL were injected. At the 21st day after collagenase injection, tendons were collected and used for histological observation.

2.10 Histological observation

The histological observation methods were similar to those previously reported (Jiao et al., 2022a). Briefly, after fixation, embedding and section, Hematoxylin-eosin (HE) and Masson trichrome staining were performed. The method of evaluating fiber alignment was described in the previous studies (Adeoye et al.,



2022) (Ozlu et al., 2019) (Erisken et al., 2013). For immunohistochemical staining, we incubated sections overnight with different antibodies (COL I, COX-2, IL-6; Servicebio; China). On the next day, the sections were incubated with the secondary antibody (HRP-anti-rabbit IgG, Servicebio, China). After that, they were observed and captured.

2.11 Statistical analysis

All results are shown as the mean \pm standard deviation. Student's t-test was used for comparisons between two groups, and one-way analysis of variance followed by Tukey's *post hoc* analysis was used for comparisons between three or more groups. Statistical significance was set at p < 0.05.

3 Results

3.1 Characterization of fullerenol

To detect the characterization of fullerenol, we investigated the size and morphology of fullerenol powder by TEM. Shown in Figure 1A, the diameter of fullerenol powder was over 1 μ m. Furthermore, we detected characteristic absorption peaks of fullerenol powder by FTIR spectra. In Figure 1B, four characteristic absorption peaks existed. In detail, broad O–H

Repeat	Zeta potential (mV)	Mobility (µ/s)/(V/cm)	Conductance (µS)	Count rate (kcps)
1	-16.61	-1.30	25	586
2	-13.21	-1.03	26	601
3	-16.51	-1.29	26	601

TABLE 2 Zeta potential of aqueous fullerenol.



Biocompatibility of fullerenol with different concentrations. (A) Optical density (OD) value of TCs treated with different-concentration (0, 0.1, 0.3, 0.5, 1, 3, 5, 10 µg/mL) fullerenol at 1 and 3 days tested by CCK-8. (B) Live/dead cell staining of TCs treated with 0, 0.3, 1, 5 µg/mL fullerenol at 3 days. Scale bar = 25 µm. (Data are presented as the mean \pm standard deviation. *p < 0.05, **p < 0.01).

stretching vibration (vO-H) presented at 3,412.80 cm⁻¹, C=C stretching vibration (vC = C) was shown at 1,596.71 cm⁻¹, O-H in-plane deformation vibration (δ sC-OH) existed at 1,354.07 cm⁻¹, and C-O stretching vibration (vC-O) was at 1,082.82 cm⁻¹. Then, we dissolved fullerenol by water into the concentration of 50 mg/mL, which became brown to black liquid (Figure 1C). Although the diameter of fullerenol powder was over 1µm, according to size distribution, particles in aqueous fullerenol were mostly from 100 to 1000 nm (Figure 1D). Notably, there was another peak from 3,000 to 6000 nm, which was probably caused by agglomeration due to high concentration. At the same time, the surface zeta potential was -15.44 ± 1.93 mV (Table 2).

3.2 Fullerenol shows low cytotoxicity on rat TCs

To detect the cytotoxicity of fullerenol, we performed CCK-8 assay. Shown in Figure 2A, at 1 day after fullerenol (10, 5, 3, 1, 0.5, 0.3, 0.1 μ g/mL) treatment, no significant difference existed between TCs treated with fullerenol and without fullerenol, indicating no cytotoxicity at 1 day. However, at 3 days, optical density (OD) value in the 10 μ g/mL group was obviously lower than control group, suggesting that 10 μ g/mL fullerenol influenced cell viability of TCs at 3 days. Based on this, we chose three concentrations (5, 1, 0.3 μ g/mL) to conduct further experiments. Furthermore, we verified the cytotoxicity of fullerenol with the three concentrations at 3 days *via* live/dead cell staining (Figure 2B). It was found that almost no dead TCs existed at the three concentrations. All the results showed that low-concentration fullerenol had good cytocompatibility with TCs.

3.3 Fullerenol inhibits inflammation of TCs caused by IL-1 β and rescues the impairments of TCs

Next, we investigated the effects of fullerenol on the inflammation of TCs and the expression of tendon-related markers. Collagen 1A1 (COL1A1) is the most important component of tendon tissues and expresses lower in tendinopathy (Cho et al., 2021; López De Padilla et al., 2021). Tenascin C (TNC) is a glycoprotein abundantly expressed in tendons subjected to high tensile and compressive stress (September et al., 2007). TNC has been proved in the regulation of cell-matrix interaction (September et al., 2007). Shown in Figure 3A, after adding IL-1β, expression of COL1A1 and TNC decreased, although there was no significant difference. Fullerenol enhanced the RNA level of COL1A1 and TNC remarkably, especially 5 µg/mL. Contrary to COL1A1 and TNC, IL-1β augmented matrix metalloproteinases-3 (MMP-3) and matrix metalloproteinases-13 (MMP-13) expression, which were closely related to inflammation. As an anti-inflammatory material, fullerenol lowered MMP-3 and MMP-13, suggesting that fullerenol alleviated inflammation. Consistent with RNA, the tendency of TNC, COL I and MMP-13 were increased by IL-1β and decreased by fullerenol (Figure 3B). In view of anti-oxidant effects of fullerenol, we also verified the anti-oxidant effect of fullerenol in tendinopathy. In Figure 3C, IL-1B induced ROS upregulation, showing that IL-1ß exacerbated oxidant stress in TCs. However, after fullerenol treatment, ROS level of TCs diminished in a concentration-dependent manner and nearly disappeared in the concentration of 5 µg/mL. All the above



results implied that fullerenol could attenuate inflammation and ROS level in TCs induced by IL-1 β .

3.4 Fullerenol inhibits migration of TCs

It was reported that migration of TCs increased in an inflammatory environment (Jiao et al., 2022b) (Wang et al., 2019). Next, we investigated the influences of fullerenol on TCs migration through scratch assay and transwell assay. Shown in Figures 4A,C, TCs in all the five groups migrated gradually at 12 and 24 h. The addition of IL-1 β accelerated migration of TCs to around 50% at 12 h and approximate 70% at 24 h. However, fullerenol was able to inhibit the migration of TCs effectively. Notably, the inhibitory effects were concentration dependent. Extremely low concentration like 0.3 µg/mL did not depress the migration of TCs, while 1 µg/mL and 5 µg/mL could suppress the migration. Similarly, in transwell assay, the number of TCs in IL-1 β group increased obviously compared with control group (Ctrl). But the number declined after fullerenol treatment in a concentration.

dependent manner (Figures 4B,D). The above results implied that fullerenol could effectively inhibit migration of TCs.

3.5 Fullerenol inhibits tendinopathy *via* MAPK pathway

Mitogen-activated protein kinase (MAPK) signaling pathway was reported to play a key role in inflammation (Jiao et al., 2022b) (Zhu et al., 2021). So, we explored the activation of MAPK signaling pathway. P38 MAPK pathway was strongly activated in stress, immune response and regulation of cell survival and differentiation (Cuadrado and Nebreda, 2010). Apparently, in our study, inflammation induced by IL-1 β increased the phosphorylation level of p38. Nevertheless, addition of fullerenol availably hindered the activation of p38 (Figures 5A,B). Interestingly, phosphorylation level of p38 decreased in a concentration-dependent manner. Identically, Erk1/2 and JNK was activated by IL-1 β and the activation was inhibited by fullerenol (Figures 5C–F). The results of Western blot suggested



Migration of TCs after IL-1 β (50 ng/mL) and fullerenol treatment. (A) Migration of TCs in control (Ctrl), IL-1 β , IL-1 β +0.3 µg/mL, IL-1 β +1 µg/mL, IL-1 β +5 µg/mL groups tested by scratch assay. Scale bar = 250 µm. (B) Migration of TCs in Ctrl, IL-1 β , IL-1 β +0.3 µg/mL, IL-1 β +1 µg/mL, IL-1 β +5 µg/mL groups tested by transwell assay. Scale bar = 100 µm. (C) Quantitative results of scratch assay. (D) Quantitative results of transwell assay. (Data are presented as the mean \pm standard deviation. *p < 0.05, **p < 0.01).

that fullerenol could restrain the activation of MAPK pathway induced by IL-1 $\beta.$

3.6 Fullerenol inhibits tendinopathy in vivo

Next, we furtherly tested anti-inflammatory effects of fullerenol on tendinopathy *in vivo*. Shown in Figure 6A, after collagenase I injection, tendinous fibers were fractured and arranged disorderly compared with Ctrl group in HE and Masson staining. But fullerenol alleviated impairment of tendinous fibers. Meanwhile, we detected expression of Collagen I (COL I), Cyclooxygenase 2 (COX-2) and IL-6 by immunohistochemical staining (Figure 6B). In Collagenase group, COL I decreased and inflammatory factors (COX-2 and IL-6) increased in comparison with Ctrl group, showing that collagenase induced inflammation of tendon tissues. Fullerenol could alleviate severity of inflammation and promote expression of COL I. All these data hinted that fullerenol reduced inflammation in tendinopathy *in vivo*.

4 Discussion

Tendinopathy is a common overload injury, with an incidence of two to three per 1,000 patients in general medicine practice (van der Vlist et al., 2021). It is challenging to manage tendinopathy. Current treatments have more or less limitations. For example, conservative treatments like eccentric exercises and shockwave therapy are not suitable for all kinds of tendinopathy (Figueroa et al., 2016). Pharmacological management, especially injection, is



+0.3 μ g/mL, IL-1 β +1 μ g/mL, IL-1 β +5 μ g/mL groups were examined by Western Blotting. (F) Quantitative results of phosphorylation levels of JNK. (Data are presented as the mean \pm standard deviation. *p < 0.05, **p < 0.01)

another important way to treat tendinopathy. Unfortunately, no standard procedure of treat tendinopathy pharmacologically is established because there is a lack of comparative studies on effects of various drug injections (Aicale et al., 2020). Furthermore, surgical treatment impaired the function of tendons and there is a need of high-quality evidence on the effects of surgeries on different tendinopathy, such as chronic patellar tendinopathy (Khan and Smart, 2016). Therefore, it is necessary to develop a new method of tendinopathy management with low side effects.

Fullerenol is a hydroxylated derivative of fullerene. Identical to previous studies (Wang et al., 2016) (Zha et al., 2022), the diameter of our aqueous fullerenol ranged from 100 to 1,000 nm. Notably, fullerenol was also manufactured to be nanomaterial with diameter lower than 100 nm (Chen et al., 2022). In terms of biological function, fullerenol show good biocompatibility and low side effects in numerous studies (Yang et al., 2021) (Yang et al., 2014b) (Zhu et al., 2007). However, it was also reported that fullerenol was cytotoxic toward human retinal pigment epithelial

(hRPE) cells at concentrations of $10-50 \mu$ M, and increased phototoxicity on hRPE cells in particular (Wielgus et al., 2010). So, in spite of good biocompatibility, fullerenol is not absolutely safe to all the normal tissue cells. In our study, we detected the cytotoxicity of fullerenol toward TCs. It was found that after short-term (1 day) treatment, fullerenol showed no cytotoxicity at concentrations of 0.1–10 µg/mL. Nonetheless, with extension of treatment time to 3 days, fullerenol at high concentration (10 µg/mL) became deleterious for cell viability of TCs. Overall, fullerenol, consisting of carbon, hydrogen and oxygen elements, exhibited good biocompatibility to TCs. But the extremely high concentration also caused impairment on TCs. All the results of cell viability are almost identical to previous studies.

Inflammation is one of the most features of tendinopathy and suppressing inflammation has been a vital treatment for tendinopathy. Mounting studies have employed antiinflammatory drugs and biomaterials to treat tendinopathy. Chen et al. reported that ibuprofen-loaded hyaluronic acid nanofibrous membranes could reduce inflammation to prevent postoperative





tendon adhesion (Chen et al., 2019). In the study of Choi et al., they synthesized lactoferrin-immobilized, heparin-anchored, poly (lactic-co-glycolic acid) nano-particles (LF/Hep-PLGA NPs) and also found that LF/Hep-PLGA NPs enhanced tendon restoration via inhibiting inflammation (Choi et al., 2020). In terms of drugs, plenty of drugs like aspirin were found to be conducive to tendinopathy treatment through many signaling pathways, such as JNK/ STAT3 pathway (Wang et al., 2019). In our study, we investigated whether fullerenol helped mitigate tendinopathy. From RNA to protein level, under inflammatory environment, fullerenol increased expression of COL I and TNC and decreased expression of MMP-3 and MMP-13. It might imply that fullerenol played a key role in both reducing inflammation and protecting tendon tissues. Furthermore, production of ROS is another factor of damage to tendon. The results of our study showed that fullerenol also had the ability to remove ROS, which was possibly due to many carbon-carbon double bonds in fullerenol.

Stimulation of inflammation tends to change the behaviors of localized cells in different tissues. Cell migration is one of the most important behaviors affected by inflammation. In previous studies, it was found that the migration of human bronchial epithelial cells increased after being treated with TNF- α , which was also a way to induce inflammation (Ren et al., 2021). Similarly, migration of fibroblasts is also influenced by inflammation. Fibroblast-like synoviocytes (MH7A cell) in rheumatoid arthritis migrated faster than synoviocytes in control group (Cai et al., 2021) (Cai et al., 2022). Zhang et al. also reported that macrophage migration inhibitory factor, a proinflammatory cytokine, promoted migration of joint capsule fibroblasts (Zhang et al., 2021). Here, in our study, we investigated the influence of fullerenol on TCs migration under the inflammatory environment by scratch assay and transwell assay. Obviously, IL-1 β treatment induced inflammation of TCs successfully and made migration of TCs faster, which was consistent with our previous study (Jiao et al., 2022b). In view of good anti-inflammatory and anti-oxidant effects of fullerenol on TCs from mRNA level to protein level, it could inhibit migration of TCs under inflammatory environment, unsurprisingly. After tendons are injured, migration of TCs may lead to formation of a fibrotic scar, causing loss of mechanical strength of the original tendon (Wang et al., 2019) (Nichols et al., 2019). Based on our results, fullerenol slowed the migration of TCs which was likely to reduce the impairment of tendon tissues.

A series of signaling pathways is of great importance in tendinopathy inhibition. Various inflammation-related pathways such as NF-KB, c-Jun N-terminal kinase (JNK)/ STAT-3 signaling pathways (Wang et al., 2019) (Vinhas et al., 2020). Besides, MAPK signaling pathways play a part in tendinopathy (Wu et al., 2022) (Moqbel et al., 2020). MAPK cascade consists of three protein kinases, including a MAPK and two upstream components, MAPK kinase (MAPKK) and MAPKK kinase (MAPKKK). So far, three MAPK pathways are found in mammalian cells, namely, the extracellular signal-regulated kinases (ERKs) pathway, the c-Jun amino terminal kinase (JNK) pathway and the p38 MAPK pathway (Kumar et al., 2003). Considering the importance of MAPK signaling pathway in inflammation, we explored the activation of MAPK pathway after fullerenol treatment. Excitingly, fullerenol curbed the phosphorylation level of all p38, ERK, and JNK, suggesting fullerenol could effectively inhibit tendinopathy via MAPK pathway.

We furtherly detected the effects of fullerenol on tendinopathy in vivo. Collagenase injection has been a common method of constructing tendinopathy model (Liu et al., 2021) (Wu et al., 2019). Collagenase injection caused disorders and swelling of fibers in tendons. At the same time, collagenase leads to increase of inflammatory factors and decrease of Collagen I. Since fullerenol had good water-solubility, we decided to inject fullerenol locally into Achilles tendons, which effects for tendinopathy compared had better with intraperitoneal injection. In our study, identical to results of cell experiments, aqueous fullerenol alleviated disorders of tendon fibers and decreased expression of inflammatory factors like COX-2 and IL-6. The animal experiments showed fullerenol was an excellent and convenient therapeutic approach to tendinopathy.

Since tendinopathy is a localized inflammatory disease, traditional drug administration is the most common treatment. Here, we firstly injected aqueous fullerenol to treat tendinopathy. In view of good biocompatibility and antiinflammatory effects, fullerenol shows good prospects in treating localized inflammatory diseases like tendinopathy in the future.

5 Conclusion

In conclusion, fullerenol is a promising biomaterial which has brilliant biocompatibility and anti-inflammatory effects and can be used to treat tendinopathy. Utilization of fullerenol helps reduce the

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side effects caused by drug administration and lower the economic burden.

Data availability statement

The raw data supporting the conclusion of this article will be made available by the authors, without undue reservation.

Ethics statement

The animal study was reviewed and approved by the Ethics Committee of Shanghai Ninth People's Hospital, Shanghai Jiaotong University School of Medicine.

Author contributions

Conceptualization, XZ and YG; methodology, XJ and ZW; software, XJ; validation, ZW, YL, and TW; formal analysis, XJ and ZW; resources, CX; writing—original draft preparation, XJ, ZW, and YL; writing—review and editing, XJ, ZW, and XZ; supervision, YG; funding acquisition, YG. All authors have read and agreed to the published version of the manuscript.

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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A fresh pH-responsive imipenem-loaded nanocarrier against *Acinetobacter baumannii* with a synergetic effect

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In recent years, the treatment of Acinetobacter baumannii infections has become a pressing clinical challenge due to its increasing incidence and its serious pathogenic risk. The research and development of new antibacterial agents for A. baumannii have attracted the attention of the scientific community. Therefore, we have constructed a new pH-responsive antibacterial nano-delivery system (Imi@ZIF-8) for the antibacterial treatment of A. baumannii. Due to its pH-sensitive characteristics, the nano-delivery system offers an improved release of the loaded imipenem antibiotic at the acidic infection site. Based on the high loading capacity and positive charge of the modified ZIF-8 nanoparticles, they are excellent carriers and are suitable for imipenem loading. The Imi@ZIF-8 nanosystem features synergistic antibacterial effects, combining ZIF-8 and imipenem to eliminate A. baumannii through different antibacterial mechanisms. When the loaded imipenem concentration reaches 20 µg/mL, Imi@ZIF-8 is highly effective against A. baumannii in vitro. Imi@ZIF-8 not only inhibits the biofilm formation of A. baumannii but also has a potent killing effect. Furthermore, in mice with celiac disease, the Imi@ZIF-8 nanosystem demonstrates excellent therapeutic efficacy against A. baumannii at imipenem concentrations of 10 mg/kg, and it can inhibit inflammatory reaction and local leukocyte infiltration. Due to its biocompatibility and biosafety, this nano-delivery system is a promising therapeutic strategy in the clinical treatment of A. baumannii infections, providing a new direction for the treatment of antibacterial infections.

KEYWORDS

Acinetobacter baumannii, Imi@ZIF-8, antibacterial infection, synergistic effect, antibiofilm

Introduction

Acinetobacter baumannii is an aerobic, gram-negative coccobacillus and is a conditional pathogen that can cause community-acquired and hospital-acquired infection (Whiteway et al., 2022). In clinical settings, it is responsible for ventilator-associated pneumonia, catheter-associated blood and urinary tract infections, sepsis, endocarditis, skin and wound infections and meningitis (Chen, 2020). *A. baumannii* is ubiquitous in nature and persistent in the hospital environment, where it often infects immunocompromised patients, particularly those in intensive care units (Shan et al., 2022). According to worldwide

data, the detection rate of *A. baumannii* in intensive care units rose from 4% to 7% between 1986 and 2003. Moreover, ICU mortality increased from 53.3% to 84.3% in patients with respiratoryassociated pneumonia due to extensively drug-resistant *A. baumannii* (Saipriya et al., 2020). *A. baumannii* was regarded as ESKAPE pathogens. In order to study and develop effective antibacterial drugs against *A. baumannii* with acquired resistance to antibiotics, the WHO has placed it on priority list (Tacconelli et al., 2018). So, there is a need for research into combatting this pathogen.

Genomic and phenotypic identification analyses were performed, demonstrating the association between A. baumannii infection and multiple virulence factors, including outer membrane lipopolysaccharides, capsular polysaccharides, proteins, phospholipases, protein secretion systems, quantum sensing and biofilm production (Harding et al., 2018; Dehbanipour and Ghalavand, 2022). These virulence factors contribute to the colonization of pathogenic bacteria and exacerbate antibiotic resistance. At present, antibiotics for the treatment of A. baumannii infection are mainly selected based on the sensitivity of pathogens to antibiotics, the severity of the disease and the infection site. Based on its pathogenicity, the clinical treatments of Α. baumannii include sulbactams, carbapenems, aminoglycosides, polymyxin, tegacycline and combined antibiotic therapy (Fishbain and Peleg, 2010; Isler et al., 2019). Carbapenems are highly effective broad-spectrum antibacterial drugs that exhibit strong antibacterial activity against gram-positive bacteria, gramnegative bacteria and anaerobes and are regarded as the "last line of defense" of antibiotics. Their antibacterial mechanism is similar to β-like lactam antibiotics, as they inhibit the formation of the bacterial cell wall by binding with penicillin-binding proteins (PBPs), leading to inactivation (Papp-Wallace et al., 2011; Skariyachan et al., 2019). Imipenem is the most widely-used carbapenem antibiotic in clinical practice and has high bactericidal activity, fast bactericidal rate and excellent bacterial inhibition against A. baumannii, especially for patients with moderate to severe infections and multi-drug resistant bacterial infections. Moreover, imipenem is a time-dependent antibacterial drug with a post-antibiotic effect on bacteria. Imipenem preferentially combines with PBP2, followed by PBP1a and PBP1b, and has a weak affinity for PBP3. This mechanism of action can reduce the release of lipopolysaccharide during bacteriolysis, providing good biological safety (Rodloff et al., 2006; Zhanel et al., 2007; Nowak and Paluchowska, 2016). However, with the wide application of carbapenem antibiotics in recent years, the drug resistance rate of A. baumannii to carbapenems has increased year by year (Jiang et al., 2022). Polymyxins are effective for treating A. baumannii infection, with low rates of resistance, but are associated with a higher risk of nephrotoxicity (Liu et al., 2021). The overall economic benefit of high-dose sulbactam or combination therapy is poor, and clinical data on the efficacy and resistance of the other antimicrobials mentioned are limited (Betrosian et al., 2008; Chu et al., 2013). Hence, finding new treatment schemes is an urgent problem.

The researcher has extensively studied and formulated treatment plans against bacterial infection and drug resistance, including the development of new antibiotic resistance inhibitors. However, the research of new antibiotics is time-consuming and costly, and the development of antibacterial resistance is significantly faster than the research and development of new antibiotics. Furthermore, drug-resistant inhibitors mainly inhibit the severe evolution of bacteria into drug-resistant bacteria, but the prevalence of drug resistance within a patient or in the population remains a challenge (Baym et al., 2016; Chang et al., 2022). In recent years, nanomaterials have attracted attention due to their physical and chemical properties, such as structural stability, large specific surface area, high porosity, easy surface modification, structural diversity and biocompatibility. They have been applied to biomedical engineering projects, such as biosensors (Singh et al., 2022), drug carriers (Vangijzegem et al., 2019; Huang et al., 2021; Liu et al., 2022), medical implants (Akgöl et al., 2021) and medical imaging (Kalva et al., 2022). The rise of nanomedicine has enabled advances in antimicrobial therapy.

Nanocarriers can selectively transport antibiotics to the site of infection due to their targeting properties, thus improving drug distribution, increasing the effectiveness of antibiotics, reducing drug side effects and overcoming bacterial resistance (Nazli et al., 2022). Many nanomaterials have been used in antimicrobial therapy, such as inorganic metal nanomaterials (gold, silver, copper, zinc, titanium), metal oxide nanoparticles (copper oxide, zinc oxide, titanium oxide, iron oxide), carbon-based nanomaterials (graphene and its derivatives, graphene quantum dots, carbon quantum dots), and organic nanostructures (chitosan, dendrimers, liposomes, micelles, vesicles), etc. (Modi et al., 2022). The antibacterial mechanisms of nanomaterials involve 1) the use of size, surface properties and other unique physicochemical properties to damage important intracellular components and interfere with the normal physiological metabolic processes of bacteria, ultimately leading to bacterial death (Cheng et al., 2022); 2) the use of the enzyme-like activity of nanomaterials, regulating the level of reactive oxygen species (ROS) to exert a strong bactericidal effect by disrupting bacterial biofilms (Godoy-Gallardo et al., 2021); 3) smart response platforms based on nanomaterials, such as pH, enzymes and temperature to enhance antimicrobial activity (Jiang et al., 2020); 4) the use of external stimulus-response properties of nanomaterials, such as light and microwaves, or synergistic antimicrobial drugs to achieve an antimicrobial activity in a single or combined treatment (Díez-Pascual, 2020; Nazli et al., 2022). Zeolite imidazole ester skeleton-8 (ZIF-8) is a porous crystalline material formed by the coordination self-assembly of zinc ion and 2-methylimidazole. ZIF-8 has not only high loading capacity but also has antibacterial activity, acid sensitivity, low cytotoxicity, and good biocompatibility, providing broad applications in biomedicine (Lian et al., 2022). The pH of the microenvironment of infected tissues is slightly lower than that of normal tissues due to acid production at the site of infection, so ZIF-8 can be used as an acid-responsive drug carrier for antimicrobial therapy with superior loading capacity, controlled drug release and enhanced targeting (Abdelhamid, 2021; Tan et al., 2022).

In summary, a novel antimicrobial nanodelivery system with pH acid response function was constructed by synthesizing positively-charged ZIF-8 (PEI@ZIF-8) and loading negatively-charged imipenem onto ZIF-8 nanoparticles (Imi@ZIF-8) using positive and negative charge adsorption forces under ultrasonic stirring. The loaded imipenem is effectively released at the site of

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bacterial infection due to the acidic microenvironment, displaying the antimicrobial synergy between imipenem and ZIF-8. In addition, subsequent research and experiments found that the Imi@ZIF-8 nanodrug system exerts a strong killing effect on *A. baumannii*. The development of an antibacterial nanosystem with good biocompatibility and biosafety provides a new strategy for the clinical treatment of *A. baumannii*.

Materials and methods

Materials

Imipenem was purchased from Aladdin Technology (China). Fetal bovine serum (FBS) and high glucose (Dulbecco's modified Eagle's medium DMEM) were obtained from Life Technologies. Dialysis membranes (2000 D), hematoxylin and eosin (HE), crystal violet (CV), and phenazine methosulfate were provided by Solarbio Technology (China). Mueller-Hinton (MH) broth was obtained from Solarbio. Polycarbonate porous membrane syringe filters (200 nm) were obtained from Whatman. The bacterial stock solution and reactive oxygen species (ROS) assay kits were purchased from Beyotime Technology (China). The LIVE/DEAD backlight bacterial viability kit was purchased from Yeasen Biotechnology (China). Anti-IL-6 and anti-Ly6G antibodies were manufactured by BOSTER Biological Technology (China). The HRP-conjugated goat anti-mouse IgG was purchased from BOSTER Biological Technology.

Experimental cells and experimental animals

The mouse lung epithelial (MLE-12) cell line was provided by the Department of Laboratory Medicine, Xiangya Medicine School, Central South University. MLE-12 cells were cultured in a sterile environment at 37°C and 5% CO₂. We prepared 5 mL of 10% fetal bovine serum, 45 mL of DMEM (high sugar type) and 1 mL of penicillin-streptomycin (1%) to form the complete medium for the cells used. The 6-week-old female BALB/c mice used in our experiments were purchased from Shrek Laboratory Animals Ltd. in Hunan, China. All mice were kept in specific pathogen-free conditions at the Animal Resource Center of Xinxiang Medical University. All animal experiments were approved by the Xinxiang Medical University Experimental Animal Ethics Committee. The ethical approval code is XYLL-20230027.

Bacterial strains and bacterial cultures

The *A. baumannii* strains were obtained from the Department of Laboratory Science, The Third Affiliated Hospital of Xinxiang Medical University (Xinxiang, Henan, China). The bacteria were stored in bacterial lyophilization solution at -80° C. A disposable sterile inoculating loop was used to collect the lyophilized solution, and the bacteria were allowed to grow on blood agar plates by triple zoning, followed by incubation at 5% CO₂ and 37°C for 24 h. Finally, individual colonies from the plates were lysed in a sterile LB broth liquid medium with a disposable sterile inoculating loop and incubated overnight at 200 rpm at 37°C to allow bacterial to growth to their logarithmic phase. The overnight culture was added to a fresh, sterilized LB broth medium, and the OD_{600} value was measured using an enzyme marker at 0.6. All experiments were repeated three times, and the average of the three values was taken.

Preparation of PEI@ZIF-8

2-methylimidazole was dissolved in methanol to form solution A, and zinc nitrate was dissolved in methanol to form solution B. Solutions A and B were mixed and allowed to react for a certain period of time, and ZIF-8 was obtained by centrifugation (Wang et al., 2020). The precipitate was washed then three times with water. ZIF-8 and PEI were mixed and stirred for a certain period of time, and the PEI-modified ZIF-8 was collected by centrifugation. Finally, the precipitate was washed three times with water.

Determination of encapsulation and loading rates of imipenem in Imi@ZIF-8

1 mg of PEI@ZIF-8 was dissolved in 1 mL of deionized water (ddH_2O) , and 1.2 mg of imipenem was added and stirred for 4–5 h at room temperature with a magnetic stirrer. The solution was left for 12 h in darkness, then the precipitate was centrifuged in darkness to obtain Imi@ZIF-8. The maximum absorbance of imipenem in the supernatant was determined by UV-visible spectrophotometry (Yu et al., 2014; Wang et al., 2016). The standard concentration gradient of imipenem was set, and the standard curve of concentration and absorbance was established. Then, the content of unbound imipenem in the supernatant was calculated.

The calculation formulas of encapsulation efficiency (EE) and loading efficiency (LE) are as follows:



Characterization of Imi@ZIF-8

Transmission electron microscopy (TEM) was performed to observe the particle size and morphology of ZIF-8, while the surface morphology of ZIF-8 was assessed using Scanning electron microscopy (SEM). A zeta potential-particle size analyzer was used to evaluate imipenem, ZIF-8, and Imi@ZIF-8, whereas dynamic light scattering instrument (DLS) was performed to analyze the hydrated particle size of ZIF-8 and Imi@ZIF-8. The characteristic absorption peak positions of ZIF-8, imipenem and Imi@ZIF-8 were detected by a ultraviolet-visible (UV-vis) spectrophotometer to further validate the successful construction of the nano-delivery system.



Characterization of ZIF-8 and Imi@ZIF-8. (A) TEM image of ZIF-8. (B) TEM image of Imi@ZIF-8. Scale bar is 150 nm. (C) SEM image of ZIF-8. (D) SEM image of Imi@ZIF-8. Scale bar is 200 nm. (E) Particle sizes of Imi@ZIF-8 detected by TEM. (F) Dynamic light scattering (DLS) analyze the hydrated particle size of ZIF-8 and Imi@ZIF-8. (G) UV-vis spectrometries of imipenem, ZIF-8 and Imi@ZIF-8. (H) Zeta Potential values of imipenem, ZIF-8 and Imi@ZIF-8. Data are presented as means \pm SD (n = 3).

Release characteristics of Imi@ZIF-8

Imipenem release from Imi@ZIF-8 was monitored at pH = 7.4 and pH = 6.5 to investigate the acidic pH response properties of Imi@ZIF-8. Specifically, 5 mL of Imi@ZIF-8 was packed into dialysis bags, which were then immersed in 50 mL of PBS solution at pH = 7.4 and pH = 6.5, respectively (Zhou et al., 2022). The dialysate was collected at 6, 12, 18, and 24 h. The absorbance value of imipenem in the dialysate at 299 nm was measured using a UV-Vis spectrophotometer to determine the cumulative release of imipenem from the solution, and the percentage release—time curves at pH = 7.4 and pH = 6.5 were plotted. All experiments were repeated three times, and the average of the experimental values was taken.

In vitro antibacterial and anti-biofilm effects of Imi@ZIF-8

Minimum inhibitory concentration (MIC)

The minimum inhibitory concentration (MIC) of imipenem and ZIF-8 against *A. baumannii* was determined using the broth dilution method. A suspension of *A. baumannii* at a concentration of 1.5×10^6 CFU/mL was added to a 96-well plate. First, the imipenem solution was prepared by serial 2fold dilution to a final concentration of 0, 0.5, 1, 2, 4, and 8 µg/mL of imipenem in each well and a volume of 100 µL. A suspension of 1.5×10^6 CFU/mL of *A. baumannii* was then prepared, and 100 µL of the suspension was added to each well using a pipette. The suspensions were incubated for 16–20 h at 37°C and 5% CO₂. Similarly, the minimum inhibitory concentration of ZIF-8 against *A. baumannii* was tested.

Disc diffusion method

After obtaining individual colonies, the concentration of *A. baumannii* solution was adjusted to 0.5 MCF with physiological saline, and the solution was applied to MH agar with a sterile cotton swab in a flat layer. After placing a filter paper disc on the staining medium, different contents of the imipenem, ZIF-8 and Imi@ZIF-8 were injected into the paper. The MH agar plates were incubated at 5% CO₂, 37°C for 24 h, and the size of the inhibition circles was observed and measured.

Live/dead backlight bacteria assay

A. baumannii was incubated with the corresponding treatment group materials for 24 h, then centrifuged and washed with sterile PBS 1–2 times, the precipitation was the treated A. baumannii. The DMAO/EthD—III mixed fluorescent dye was prepared according to the manufacturer's instructions, and the bacteria were resuspended with the prepared fluorescent dye mixture. The two mixtures were mixed and incubated at room temperature for 15–20 min in darkness. Subsequently, 15 μ L of the stained bacterial suspension was dropped onto a sterile slide and covered with an 18 mm square sterile slide. The survival of bacteria was observed under a laser confocal microscope (CLSM).

Crystal violet staining

100 μ L of drug culture solution was added to each well of a 96well cell culture plate, and 100 μ L of an overnight culture of *A. baumannii* was inoculated at 37°C for 24 h to allow the cells to adhere to the wall. PBS was used to carefully wash three times to remove planktonic bacteria, and the bacterial biofilm was fixed with a formaldehyde solution. The samples were left to dry naturally, and 200 μ L of 1% crystal violet dye was added to



each well and left for 15 min at room temperature. The stain was then washed off 2–3 times with PBS, left to dry, and lysed using 95% ethanol. Then, the solution was incubated in an incubator at 37° C for 25 min. The absorbance value of the lysate was measured using an enzyme marker (570 nm) to determine the biofilm biomass.

The percentage of biofilm eradication and inhibition was calculated as follows:

$$Bio film \ eradication / inhibition \ rate \ (\%) = \left(1 - \frac{OD \ experiment}{OD \ control}\right) \times 100\%.$$

Confocal laser scanning microscope analysis

The biofilms of *A. baumannii* were formed in laser-scanning confocal Petri dishes, and the bacterial biofilms were fluorescently stained with a live/dead bacterial fluorescent dye kit to observe the survival of the biofilms under a CLSM.

In vitro antibacterial mechanism of Imi@ ZIF-8

Reactive oxygen species (ROS) measurement

PBS, Imipenem, ZIF-8 and Imi@ZIF-8 solutions were added to the bacterial suspension (1 \times 10⁶ CFU), co-cultured for 24 h and then resuspended in sterile saline. 1 µL of 2,7-dichlorodihydrofluorescein diacetate (DCFH-DA) fluorescent dye was added in the dark and incubated at 37°C for 1 h. The fluorescence value (excitation/emission wavelength 485/535 nm) was measured with a fluorescence enzyme marker, with the fluorescence intensity of 2,7-dichlorofluorescein (DCF) being proportional to the level of ROS (Scott et al., 1998; Hsieh et al., 2001).

Determination of malondialdehyde (MDA)

A. baumannii was incubated with each group of drug culture solution for 24 h, centrifuged (12,000 rpm, 2 min), and the supernatant was discarded and resuspended in 1 mL of 2.5% (w/ v) trichloroacetic acid. The solution was centrifuged (12,000 rpm,

20 min, 4°C) again to collect the supernatant, and 1 mL of 5% thiobarbituric acid (TBA) solution was added for dilution. An equal volume of 20% (w/v) TCA was added to the mixture; the reaction was carried out in a water bath at 100°C for 30 min, and the mixture was centrifuged (12,000 rpm, 20 min, 4°C). The final absorbance value was measured at 532 nm, and the MDA content (pg/mL) was calculated based on the molar extinction coefficient (1.56 nM⁻¹cm⁻¹).

Construction of the mouse peritonitis model in *Acinetobacter baumannii*

The 6-week-old female BALB/c mice were randomly divided into four groups: the PBS group, imipenem group, ZIF-8 group, and Imi@ZIF-8 group. The mice were infected with an intraperitoneal injection of 150 μ L containing 1 \times 10⁶ CFU/L A. baumannii. After 12-24 h, inflammation, necrosis and infiltration of inflammatory cells (neutrophils, lymphocytes and macrophages) were observed in the liver tissue of the mice, indicating the successful establishment of the abdominal infection mouse model. After the successful construction of the mouse model, the drug was administered once daily, and the injection continued for 3 days. The dosing concentration for each group was calculated based on the imipenem concentration, achieving a final dose of 10 mg/kg per group. After the seventh day of administration, whole blood was collected from the mice, and the liver tissues were stained with hematoxylin and eosin (HE) to observe the inflammatory infiltration and necrosis of liver tissues. Immunohistochemical staining was performed to detect the pro-inflammatory factors interleukin-6 (IL-6) and the neutrophil-specific marker Ly6G.

Biocompatibility and biosafety of Imi@ZIF-8

The biocompatibility of Imi@ZIF-8 was assessed using the CCK-8 method. MLE-12 cells (2×10^3 /well) were inoculated in a 96-well plate and incubated for 24 h, washed 1–2 times with sterile PBS solution, and ZIF-8 was prepared at a concentration gradient of 10, 20, 40, 60, 80, 100, and 120 µg/mL (using DMEM medium as solvent). After incubation with MLE-12 cells for 24 h, CCK-8 solution (10 µL) was added to each well for 3 h, and



In vitro antibacterial effect of Imi@ZIF-8. (A) Different concentration of Imipenem and ZIF-8 inhibited on the growth of *A. baumannii*. (B) Corresponding inhibition zone diameters of imipenem, ZIF-8 and Imi@ZIF-8 against *A. baumannii*. (C) Photographs of agar plates showing the antibacterial activity for PBS, imipenem, ZIF-8 and Imi@ZIF-8 against *A. baumannii*. (D) CLSM imaging of death/live staining after *A. baumannii* exposure to various treatments (PBS, imipenem, ZIF-8 and Imi@ZIF-8) for 24 h. Scale bar is 20 μ m. Data are presented as mean \pm SD (*n* = 3). Compared to ZIF-8: ***p < 0.001.

the absorbance value was measured at 450 nm. Healthy female BALB/c mice at 6 weeks of age were randomly divided into four groups and injected with PBS, Imipenem, ZIF-8 or Imi@ZIF-8 *via* the tail vein. After 1 week, whole blood was collected, and blood biochemical parameters (RBC, WBC, PLT, CRP, ALT,

AST, BUN, and CREA) were measured. Mice were executed by cervical dislocation. The major organs (heart, liver, spleen, lungs and kidneys) were taken for hematoxylin and eosin (HE) staining, and the lesions were observed under the microscope.



Influence of Imi@ZIF-8 on the biofilm formation and damage of *A. baumannii*. (A) CV staining analysis after receiving different treatments for 24 h. The absorbance value was detected using an enzyme marker (570 nm) to reflect biofilm formation. (B) CV staining analysis. Relative percentage of biofilm inhibition. (C) CLSM images of *A. baumannii* biofilms treated with various treatments (PBS, imipenem, ZIF-8, or Imi@ZIF-8) for 24 h. Scale bar is 20 μ m. Data are indicated as mean \pm SD (n = 3). Compared to the PBS group: ***p < 0.001; Compared to the imipenem group: ***p < 0.001.



Antibacterial mechanism of Imi@ZIF-8. (A) The fluorescence intensity of *A. baumannii* was measured using a fluorometer to reflect ROS formation. (B) MDA contents of *A. baumannii* treated with respective materials (PBS, imipenem, ZIF-8 or Imi@ZIF-8) for 24 h represents the extent of membrane damage. Data are presented as mean \pm SD (n = 3). Compared to the PBS group and the imipenem group: ***p < 0.001.

Statistical analysis

All data were expressed as mean \pm standard deviation, and statistical analysis was performed using SPSS 20.0 software. Differences between groups were analyzed using oneway ANOVA, followed by further analysis using Tukey's post-test (*p < 0.05, **p < 0.01, ***p < 0.001).

Results and discussion

Construction and characterization of the Imi@ZIF-8 nano-drug delivery system

In this study, the new antibacterial nano drug delivery system Imi@ZIF-8 was designed and synthesized. The particle size, morphology and dispersion of the prepared nano-drug delivery system were studied by TEM. The positively-charged ZIF-8 synthesized in this experiment was spherical, with an average particle size of about 80 nm and uniform distribution (Figure 1A). The overall morphology of the imipenem-loaded Imi@ZIF-8 showed no significant change, with a spherical shape and a particle size of around 80 nm (Figures 1B, E). SEM confirmed the TEM results, showing both ZIF-8 and Imi@ZIF-8 with particle sizes of around 80 nm and good dispersion (Figures 1C, D). DLS allows analysis of the hydrated particle size of the nanoparticles, complementing the TEM results on nanoparticle size. As shown in Figure 1F, the average hydrated particle size of ZIF-8 and Imi@ZIF-8 was about 80 nm. Moreover, the ζ-potential analysis of imipenem, ZIF-8, and Imi@ZIF-8 was performed separately, as displayed in Figure 1H. The ζ-potential of ZIF-8 after PEI modification showed a positive charge of 28.59 ± 1.39 mV, while imipenem was negatively charged (-8.57 ± 0.59 mV). The zeta potential of Imi@ZIF-8 after loading imipenem was 21.35 ± 1.35 mV. This lower value indicated that ZIF-8 was successfully loaded with imipenem and that the Imi@ZIF-8 nano-delivery system had good stability and dispersibility. Subsequently, UV-vis spectroscopy was performed to verify the encapsulation of imipenem in the ZIF-8 nanoparticles. As illustrated in Figure 1G, ZIF-8 showed a characteristic absorption peak at 216.5 nm, while imipenem had a characteristic absorption peak at 299 nm. Imi@ZIF-8 exhibited both characteristic absorption peaks in UV-vis, corresponding to imipenem (299 nm) and ZIF-8 (216.5 nm), indicating that ZIF-8 was successfully loaded with imipenem. These results provide strong evidence for the successful preparation of the Imi@ZIF-8 nanodrug delivery system.

In vitro release study of the Imi@ZIF-8 nano drug delivery system

ZIF-8 is an excellent nano-delivery vehicle due to its large specific surface area and porous structure, and the ZIF-8 nanoparticles were further modified to be positively charged, improving imipenem release. EE and LE of imipenem were determined using a UV-vis spectrophotometer at 299 nm. As shown in Figure 2A, the EE and LE of imipenem in the nanodelivery system were 76.38% \pm 2.05% and 33.68% \pm 1.53%, respectively. To investigate the pH response of the drug delivery system, the release of imipenem from Imi@ZIF-8 in neutral (pH = 7.4) and acidic environments (pH = 6.5) were evaluated at different time periods. Due to the anaerobic fermentation of bacteria at the site of infection, large amounts of organic acids are produced, creating an acidic microenvironment (Qiao et al., 2019). In this experiment, pH = 6.5 was used to simulate the acidic microenvironment of bacterial infection in vivo. The 24 h cumulative release efficiency of imipenem was 48.58% ± 1.49% at pH 7.4% and 59.36% ± 1.39% at pH 6.5 (Figure 2B), indicating that the acidic microenvironment resulted in a higher imipenem release rate. The facilitated release of imipenem from the nano-delivery system at the site of bacterial infection leads to superior antibacterial activity. This facilitated release is due to the protonation of the 2methylimidazole ligand in ZIF-8 under acidic conditions, resulting in the breaking of the ligand bond between zinc and 2methylimidazole, releasing imipenem from the nano-loaded system (Zheng et al., 2016; Soomro et al., 2019). The above experimental results indicated that ZIF-8 is a good drug delivery



Anti-infection effects of Imi@ZIF-8 *in vivo*. After successful establishment of the mouse peritonitis model in *A. baumannii*, BALB/c mice were intraperitoneal injected with PBS, imipenem, ZIF-8, or Imi@ZIF-8 (10 mg imi/kg). (A) Bacterial counting of liver tissues after treated with the respective materials. Data are indicated as means \pm SD (n = 3). (B) Representative images of liver tissues after H&E staining at 7 days after intraperitoneal injected with PBS, imipenem, ZIF-8, or Immunohistochemistry staining of Ly6G in liver tissues at 7 days after intraperitoneal injected with PBS, imipenem, ZIF-8, or Imi@ZIF-8, respectively. (D) Immunohistochemistry staining of IL-6 in liver tissues. Scale bar is 100 µm.

vehicle, and Imi@ZIF-8 is responsive to acidic microenvironments, providing superior antibacterial drug release at the site of bacterial infection.

In vitro antibacterial and anti-biofilm effects of Imi@ZIF-8

MIC is a measure of the antimicrobial performance of an antimicrobial agent. A broth dilution method was used to determine the MIC value of the free drugs. As shown in Figure 3A, the MIC value of imipenem and ZIF-8 against *A. baumannii* was 4 μ g/mL and 256 μ g/mL, respectively. Based on the results, the fractional inhibitory concentration (FIC) index of imipenem and ZIF-8 against *A. baumannii* was calculated to be less than 0.5, which illustrates the synergistic antibacterial effect of imipenem and ZIF-8 against *A. baumannii*. Furthermore, the inhibition activity of free imipenem, ZIF-8 and Imi@ZIF-8 were compared using the disc diffusion method, showing that Imi@ZIF-8 had the largest inhibition circles of 15 μ g Imi@ZIF-8, Imipenem

and ZIF-8 were 24.81 \pm 0.73 mm, 21.39 \pm 0.51 mm and 11.64 \pm 0.63 mm, respectively (Figure 3B). Figure 3C shows the bacterial counts of A. baumannii after 24 h treatment with PBS, ZIF-8, imipenem, and Imi@ZIF-8. A lower number of colonies was observed in the ZIF-8 group compared to the PBS group, but the inhibition effect was not significant. In contrast, the bacterial growth in the free imipenem and Imi@ZIF-8 groups was significantly inhibited, with the Imi@ZIF-8 group showing the most obvious effect. The above results indicate that ZIF-8 can enhance the antibacterial ability of imipenem against A. baumannii. Subsequently, the DMAO/EthD-III live/dead bacterial staining kit was used to further observe the bacterial survival status of the free drug and Imi@ZIF-8 groups. Each treatment group underwent livedead fluorescence staining and was observed under a CLSM. The DMAO fluorescent dye stains dead and alive bacteria green, whereas EthD-III only stains dead bacteria with a red fluorescent dye. The PBS group showed basically no red fluorescence signal, indicating that A. baumannii was alive (Figure 3D). However, the ZIF-8, Imipenem, and Imi@ZIF-8 treated groups demonstrated an increasing intensity of red fluorescence, indicating increasing A. baumannii death. After 24 h of Imi@ZIF-8 treatment, nearly all A.



The toxicity assessment of Imi@ZIF-8. (A) MLE-12 cells was treated with various concentration of ZIF-8 for 24 h; the cell viability (%) was assayed by CCK-8. (B–J) Blood routine and blood biochemical indices of mice at 7 days after intraperitoneal injection of PBS, imipenem, ZIF-8 and Imi@ZIF-8. Data are indicated as means \pm SD (n = 3).

baumannii were dead. In summary, ZIF-8 and imipenem exerted synergistic antibacterial effects against *A. baumannii*, with the Imi@ ZIF-8 nano-delivery system being more effective than free imipenem and ZIF-8. Therefore, the nano-delivery system was effective against *A. baumannii* infection.

Bacterial biofilms are associated with bacterial drug resistance, and the ability to inhibit biofilm formation is an important indicator in determining the clinical application of antimicrobial agents (Sharahi et al., 2019). Crystalline violet staining (CV) is a commonly used assay for early and mid-stage biofilm formation, with higher absorbance values indicating higher amounts of biofilm production by bacteria. As shown in Figure 4A, the absorbance value of the PBS group at 570 nm after 24 h of treatment is 0.313 ± 0.0097 , while the Imi@ZIF-8 group had an absorbance value of only 0.092 ± 0.006 ; Figure 4B shows the percentage of biofilm inhibition of *A. baumannii* by each treatment group after 24 h of treatment. The Imi@ZIF-8 group demonstrated a biofilm inhibition rate of about 71%, much higher than that of the imipenem group (20.93%) and the ZIF-8 group (49.74%), indicating that Imi@ZIF-8 exerted the strongest inhibitory effect on *A. baumannii* biofilm formation. To investigate the killing ability of the Imi@ZIF-8 nano-delivery system on *A. baumannii* biofilms, the "Live/Dead Bacterial Staining Kit" was used, and the survival of the biofilms was assessed under CLSM. The strongest red fluorescence was observed in the Imi@ZIF-8 group, indicating a basically all-dead biofilm, showing a



significantly stronger killing effect on *A. baumannii* than the PBS, imipenem and ZIF-8 groups (Figure 4C).

Antimicrobial mechanism of Imi@ZIF-8

Although small amounts of ROS are important for maintaining the life cycle of cells, excessive ROS can lead to oxidative stress, disrupting the integrity of bacterial cell membranes and interfering with a range of normal physiological activities, ultimately inducing bacterial death (Hui et al., 2020). Therefore, a DCFH-DA fluorescent probe was used to detect intracellular ROS production in A. baumannii. As shown in Figure 5A, the four groups, PBS, Imipenem, ZIF-8 and Imi@ZIF-8, showed increasing ROS production by A. baumannii, with the Imi@ZIF-8 group inducing a significantly higher amount of ROS production than the other treatment groups. In addition, nanomaterials may trigger lipid oxidation reactions in bacteria, and the MDA method was used to detect the lipid oxidation levels in A. baumannii to reflect the extent of bacterial damage. A significant difference in MDA content was observed between the different treatment groups (Figure 5B), with the highest MDA content produced by A. baumannii after Imi@ZIF-8 treatment. These findings suggested that Imi@ZIF-8 induced the most severe damage to A. baumannii.

The above results indicated that the Imi@ZIF-8 nano-delivery system effectively eliminated *A. baumannii* by interfering with the

normal physiological activities of bacterial cells, catalyzing the production of ROS, inducing lipid oxidation and other antibacterial mechanisms.

In vivo anti-infective effect of Imi@ZIF-8

To further investigate the in vivo anti-amastigotes effect of Imi@ ZIF-8, a celiac disease mouse model was constructed by infection with A. baumannii. The mice with peritonitis were randomly divided into the PBS group, imipenem group, ZIF-8 group and Imi@ZIF-8 treatment groups. After 3 days of continuous tail vein administration, the mice were removed and executed, and the liver tissues were collected under aseptic conditions for bacterial counting. The different treatment groups showed significantly different results. Compared with the other treatment groups, the Imi@ZIF-8 group had the lowest number of A. baumannii in the liver tissues, followed by the imipenem group, the ZIF-8 group, and the PBS group (Figure 6A). The pathological hepatic changes in mice with peritonitis were observed by HE staining. Furthermore, the HE section results also provided good evidence of the in vivo antibacterial effect of Imi@ZIF-8. As shown in Figure 6B, severe inflammatory cell infiltration and congestion were visible in the liver tissue sections of the PBS group, demonstrating a significant inflammatory response. However, the liver tissue in the Imi@ ZIF-8 group was significantly milder than the other treatment groups, with no significant inflammatory cell infiltration and mostly normal

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liver morphology. Among all the treatments, Imi@ZIF-8 was the most effective in combating *A. baumannii* infection. Subsequently, immunohistochemical staining was used to detect inflammation in the liver tissue following *A. baumannii* infection by determining the expression levels of IL-6 and Ly6G, which was positive when brownish-yellow particles were present. The immunohistochemical images revealed that the increased expression of IL-6 and Ly6G in the imipenem, ZIF-8 and PBS groups, in order, which were significantly higher than those in Imi@ZIF-8 group (Figures 6C, D). These results indicated that Imi@ZIF-8 effectively inhibited the infiltration and inflammatory response of leukocytes and had excellent therapeutic efficacy in mice with celiac disease caused by *A. baumannii*. Collectively, the above *in vivo* and *in vitro* results demonstrated that Imi@ZIF-8 could be an effective novel nano-delivery system against *A. baumannii*.

Biocompatibility and biosafety of Imi@ZIF-8

To investigate the feasibility of ZIF-8 as a nanodrug carrier for antimicrobial therapy, the biocompatibility and biosafety of the Imi@ ZIF-8 nanodrug delivery system were analyzed. MLE-12 cells were treated with different concentrations (10, 20, 40, 60, 80, 100, and 120 $\mu g/mL)$ of ZIF-8 for 24 h, and the cell viability of the MLE-12 cells was assayed using the CCK-8 method. The results are shown in Figure 7A. MLE-12 cell activity decreased as the concentration of ZIF-8 was increased, but its cell survival rate remained higher than 80%, indicating the good biocompatibility of Imi@ZIF-8. These results have laid a foundation for clinical antibacterial treatment. To assess the in vivo safety and toxic effects of ZIF-8 and Imi@ZIF-8, female BALA/c mice were randomly divided into four groups and injected with PBS, Imipenem, ZIF-8, and Imi@ZIF-8, respectively. Hematological parameters were measured to detect the toxicity of the nanomaterials in mice 1 week after tail vein administration, including whole blood parameters such as white blood cells (WBC), red blood cells (RBC), hemoglobin (HGB), and platelets (PLT), C-reactive protein (CRP), liver function parameters alanine aminotransferase (ALT) and glutathione aminotransferase (AST), and renal function parameters such as blood urea nitrogen (BUN), and creatinine (CREA). As shown in Figures 7B-J, no significant difference was observed in all hematological indicators between the groups, indicating no significant toxic effect from ZIF-8 and Imi@ZIF-8. In addition, the pathological changes in the major organs (heart, liver, spleen, lung and kidney) were assessed by HE staining to further clarify the in vivo toxicity of ZIF-8 and Imi@ZIF-8, as shown in Figure 8, no inflammation, fibrosis, necrosis or histological abnormalities were observed. The results suggest that ZIF-8 and Imi@ZIF-8 induce no significant damage to the heart, liver, spleen, lungs and kidneys and that they have good biosafety and low toxic side effects. In conclusion, the Imi@ZIF-8 nano-delivery system has good biocompatibility and biosafety.

Conclusion

The above experimental studies demonstrated the promising potential of our novel antimicrobial nano-delivery system (Imi@ZIF-8) in the treatment of *A. baumannii* infections, showing good

biocompatibility, high biosafety, pH acid responsiveness and efficient antimicrobial action. Due to its response to acidic pH, the Imi@ZIF-8 nano-delivery system features an improved release of the loaded imipenem at the acidic bacterial infection sites. Therefore, a synergistic antibacterial effect is achieved between the ZIF-8 and imipenem in the nano-delivery system, further enhancing its antibacterial activity against A. baumannii. When the loaded imipenem concentration reaches 20 µg/mL, Imi@ZIF-8 is highly effective against A. baumannii in vitro. Moreover, Imi@ZIF-8 exerts excellent therapeutic efficacy against A. baumannii at imipenem concentrations of 10 mg/kg and facilitates the inhibition of inflammatory response and leukocyte infiltration in mice with peritonitis, which promotes tissue recovery after bacterial infection. Overall, the Imi@ZIF-8 nano-delivery system provides a new clinical treatment strategy for the antimicrobial treatment of A. baumannii and provides a strong foundation for nano-antimicrobial therapy. More studies on the preventive effect of this nanomaterial on bacterial infections can be conducted in the future.

Data availability statement

The original contributions presented in the study are included in the article/Supplementary Material, further inquiries can be directed to the corresponding author.

Ethics statement

The animal study was reviewed and approved by Ethics Committee of The Third Affiliated Hospital of Xinxiang Medical University.

Author contributions

GSM and LXS conceived and designed the project. NXQ supervised this project. LXS conducted the experiments and analyzed experimental data. FMM and HLWH performed in vivo mice test. LH performed the statistical analysis. GSM wrote the manuscript, and all authors discussed the results and proofread this paper.

Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Unlocking cellular barriers: silica nanoparticles and fullerenol conjugated cell-penetrating agents for enhanced intracellular drug delivery

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The limited delivery of cargoes at the cellular level is a significant challenge for therapeutic strategies due to the presence of numerous biological barriers. By immobilizing the Buforin II (BUF-II) peptide and the OmpA protein on magnetite nanoparticles, a new family of cell-penetrating nanobioconjugates was developed in a previous study. We propose in this study to extend this strategy to silica nanoparticles (SNPs) and silanized fullerenol (F) as nanostructured supports for conjugating these potent cell-penetrating agents. The same molecule conjugated to distinct nanomaterials may interact with subcellular compartments differently. On the obtained nanobioconjugates (OmpA-SNPs, BUF-II-PEG₁₂-SNPs, OmpA-F, and BUF-II-PEG₁₂-F), physicochemical characterization was performed to evaluate their properties and confirm the conjugation of these translocating agents on the nanomaterials. The biocompatibility, toxicity, and internalization capacity of nanobioconjugates in Vero cells and THP-1 cells were evaluated in vitro. Nanobioconjugates had a high internalization capacity in these cells without affecting their viability, according to the findings. In addition, the nanobioconjugates exhibited negligible hemolytic activity and a low tendency to induce platelet aggregation. In addition, the nanobioconjugates exhibited distinct intracellular trafficking and endosomal escape behavior in these cell lines, indicating their potential for addressing the challenges of cytoplasmic drug delivery and the development of therapeutics for the treatment of lysosomal storage diseases. This study presents an innovative strategy for conjugating cell-penetrating agents using silica nanoparticles and silanized fullerenol as nanostructured supports, which has the potential to enhance the efficacy of cellular drug delivery.

KEYWORDS

nanobioconjugate, Buforin II, OmpA, silica nanoparticles, fullerenol, cellular uptake, endosomal escape

1 Introduction

One of the significant obstacles to the safe and efficient delivery of pharmacological agents to the desired tissues or cells is the development of carriers that can pass through different biological barriers, such as the cellular membrane, while avoiding the immune response, side-target effects, or degradative pathways, to ultimately reach the target site while maintaining high availability of the therapeutic cargo (McNeil, 2011; McNeil, 2018).

Carriers based on nanoparticles (NPs) have been evaluated due to their multifunctionality, which results from their easily modifiable particle shape and size, material composition, and structure, according to the requirements of both the different cargoes and the target sites, achieving not only high biocompatibility, bioavailability, and biodistribution, but also on-target effects (Hossen et al., 2018; Karabasz et al., 2020). For instance, Planque et al. (2011) reported that membrane permeability and integrity are highly dependent on the size and surface chemistry of the NPs. Silica nanoparticles (SNPs) are one of the preferred nanomaterials for drug delivery due to their many advantageous properties. This material is an excellent candidate for drug carriers due to its high thermal stability, chemical inertness, high hydrophilicity and biocompatibility, simple functionalization and high loading capacity, and inexpensive synthesis (Gonçalves, 2018; Esim et al., 2019). Recently, SNPs have been utilized for the diagnostic and therapeutic delivery of contrast agents and drugs, biosensors, DNA carriers, and enzyme immobilization (Kim et al., 2019). Fullerenes, on the other hand, are an emerging class of carbon-based nanomaterials for cellular-level cargo delivery (Bolskar, 2013). These materials exhibit a structure with unique physicochemical properties and a highly symmetric cage with different sizes. The C₆₀ fullerene has the most symmetrical structure. Fullerene-based systems have been used to investigate the release of chemotherapeutic agents to eliminate the side effects of drugs such as doxorubicin and paclitaxel, photosensitizers for the activation of reactive oxygen species for the treatment of cancer cells, nucleic acid release, drugs with anti-HIV-1 activity, transdermal release, fullerenols with antioxidant activity, cardiovascular drugs and release in the brain (Kazemzadeh and Mozafari, 2019).

In addition to the use of nanomaterials, known cell penetration agents, such as the protein OmpA (López-Barbosa et al., 2019) or the antimicrobial peptide BUF-II (Cuellar et al., 2018), are also used to increase the membrane permeability of drugs. These agents have the ability to translocate across biological barriers such as the cell membrane or even the blood-brain barrier (Komin et al., 2017). However, these molecules lack stability and have a short lifetime in biological systems, a problem that can be resolved by immobilizing them on nanomaterials (Alves and Olívia Pereira, 2014). Over the past few years, we have developed a dual strategy to engineer the surface of nanocarriers. This strategy involves functionalizing them with cell-penetrating agents and combining their attributes to create carriers that are more stable and have a higher loading capacity for therapeutic agents. By doing so, we aim to enhance the release of therapeutic agents from these carriers. The purpose of this study is to examine the effect of changing the nanostructured support on the translocation capacity and endosomal escape ability of cellpenetrating agents. To accomplish this, we intend to combine our knowledge of SNPs and fullerenol as potential nanostructured supports for conjugating these agents. Our goal is to determine if the resulting nanobioconjugates have the potential for efficient cell penetration and endosomal escape, which is essential for the success of many drug therapies.

Overall, the objective of our research is to determine the efficacy of various nanostructured supports in enhancing the performance of cell-penetrating agents. By investigating the translocation and endosomal escape ability of these agents, we hope to gain insights that will lead to the future development of more effective drug therapies.

2 Materials and methods

2.1 Materials

Tetraethylorthosilicate (TEOS) (98%), methanol, ammonia solution (30%-32%), tetramethylammonium hydroxide (TMAH) (25%), (3-Aminopropyl)triethoxysilane (APTES) (98%), (25%), amine-PEG₁₂-propionic glutaraldehyde acid. N-hydroxysuccinimide (NHS) (98%), N-[3-dimethylammino)propyl]-N'-ethyl carbodiimide hydrochloride (EDC) (98%), dimethyl sulfoxide (DMSO), Fullerene C_{60} , Tetra-nbutylammonium hydroxide (TBAH) (40% in water), toluene, hydrogen peroxide (H₂O₂), glacial acetic acid, 2-propanol, diethyl ether, and hexane were purchased from Sigma-Aldrich (MO, United States). Buforin Π (BUF-II-TRSSRAGLQFPVGRVHRLLRK) was purchased from GL Biochem Shanghai (Shanghai, China). Vero Cells (ATCC[®] CCL-81) and THP-1 Cells (ATCC° TIB-202) were used for delivery MTT (3-[4,5-Dimethylthiazol-2-yl]-2,5assays. Diphenyltetrazolium Bromide), DAPI (4',6-diamidino-2phenylindole, dihydrochloride), and Lysotracker Green DND-26 was purchased from Thermo Scientific (MA, United States). Dulbecco's modification of Eagle medium (DMEM), Roswell Park Memorial Institute (RPMI) 1640 medium, and fetal bovine serum (FBS) were obtained from Biowest (MO, United States).

2.2 OmpA overexpression in E. coli

OmpA protein was obtained from overexpression in *Escherichia coli*, following the protocol developed by Aguilera-Segura et al. (2014). *E. coli* K-12 W3110/pCA24N OmpA+34 was grown in Luria-Bertani (LB) agar plates [yeast extract (5 g L⁻¹), bacto tryptone (10 g L⁻¹), NaCl (10 g L⁻¹)] supplemented with chloramphenicol (50 µg mL⁻¹), and incubated for 16 h at 37°C, 250 rpm. Fresh liquid LB medium (19.5 mL) was inoculated with 500 µL from the previous culture and incubated at 37°C, 250 rpm, until reaching an optical density of 0.7 at 600 nm (OD600 nm). OmpA was obtained by inducing with IPTG (isopropylthio- β -galactoside) (2 mM) and by culturing for 3 more hours.

2.3 OmpA purification and characterization

The culture was centrifugated to obtain a pellet of OmpA overexpressed *E. coli*. The pellet was resuspended in buffer lysis in a ratio of 4 mL g^{-1} , sonicated at 4° C for 40 min and 37% amplitude, and centrifuged at 13,000 rpm and 4° C for 15 min.



Since OmpA protein was cloned with a His-tag, purification was attained by exposing the recovered supernatant to the Dynabeads[®] TALON[®] kit (Invitrogen). Purified OmpA protein was verified by SDS-PAGE, which showed a single 31 kDa band that agrees well with the molecular weight of OmpA. Concentration was measured via NanoDrop Spectrophotometer (Thermo Fisher Scientific) at 280 nm.

2.4 Synthesis and silanization of SNPs

SNPs were synthesized based on a Stober-like approach. The method involves hydrolysis and polycondensation of TEOS in an alcohol, water, and ammonia solution (Figure 1A) (Stober et al., 1968; Edrissi et al., 2011). Briefly, Ultrapure (type I) water (ultrapure water with а resistivity> 18 MΩ-cm and conductivity $<0.056 \ \mu\text{S cm}^{-1}$) (1.5 mL) and methanol (66.3 mL) were mixed. TEOS (0.9 mL) was then added and sonicated for 20 min using an ultrasonic bath (Elmasonic EASY 60H, 37 kHz, 150 W), then 30% ammonia in an aqueous solution (4.5 mL) was added, and the mixture was left in ultrasound for another 60 min in which a cloudy white suspension formed. The SNPs were centrifuged (Z-216, Hermle Labortechnik GmbH, German) and washed with Ultrapure (type I) water $(3 \times 20 \text{ min}, 14,500 \text{ rpm})$. Silanized SNPs were synthesized using a ratio of TEOS 95% and APTES 5% (Figure 1B) (Shafqat et al., 2019). Silanization with APTES renders aminopropyl functionalities on the surface of the NPs, which can be used to conjugate further BUF-II and OmpA or crosslinkers to generate reactive groups for coupling them. The silanized SNPs were centrifuged (Z-216, Hermle Labortechnik GmbH, German) (4 \times 20 min, 14,500 rpm) and washed with Ultrapure (type I) water. BUF-II and OmpA were conjugated according to the calculations presented in Supplementary Data S1 (Rangel-Muñoz et al., 2020).

2.5 Synthesis and silanization of fullerenol

Fullerenol was prepared from fullerene C_{60} by hydroxylation with H_2O_2 and TBAH as a phase transfer catalyst under organicaqueous bilayer conditions (Kokubo et al., 2011). Briefly, to a solution of fullerene C_{60} (100 mg) in toluene (50 mL), an aqueous solution of 30% H_2O_2 (10 mL) and TBAH (40% in water, 500 µL) was added and stirred for 16 h at 60°C. Subsequently, to eliminate residual TBAH, the aqueous phase containing the fullerenol was separated, and fullerenol was precipitated with a mix of 2-propanol, diethyl ether, and hexane (7:5:5, 85 mL). Then, to complete the purification, we combined dialysis (cellulose membrane dialysis tubing) and freeze-drying (Conversion: 100%, yield after purification: 75%) (De Santiago et al., 2019). Next, fullerenol (50 mg) was dissolved in 15 mL of Ultrapure (type I) water. TMAH solution (500 µl, 25% (v/v)) and



glacial acetic acid (25 μ L) were then added to the solution and sonicated for 10 min. APTES solution (500 μ l, 20% (v/v)) was added to the fullerenol solution for silanization. The silanized fullerenol was washed with Ultrapure (type I) water to remove the APTES that was not covalently attached to the fullerenol.

2.6 BUF-II and OmpA bioconjugation

Briefly, 100 mg of silanized SNPs or fullerenol were suspended in 30 mL of Ultrapure (type I) water and sonicated for 10 min (Elmasonic EASY 60H, 37 kHz, 150 W). This was followed by adding 2 mL of glutaraldehyde 2% (v/v) and by letting the mixture left to react in an orbital shaker for 1 h at 220 rpm. The amine-PEG12-propionic acid spacer was utilized to impart flexibility to BUF-II conjugated to SNPs or fullerenol, thereby increasing the probability of interaction with the target sites. After adding 10 mg of amine-PEG₁₂-propionic acid, the mixture was shaken for 24 h at 220 rpm. Finally, 100 mg of functionalized nanomaterial was resuspended in 30 mL of type I Ultrapure water. BUF-II was conjugated to the carboxyl groups of the spacer by its N-terminal using two equivalents of EDC and two equivalents of NHS (concerning the carboxyl groups) (Figure 2; Supplementary Figure S1). BUF-II (1 mg BUF-II per 100 mg of functionalized nanomaterial) was added and the mixture was shaken at 220 rpm for 24 h. The obtained nanobioconjugates were centrifuged (Z-216, Hermle Labortechnik GmbH, German) $(4 \times 20 \text{ min}, 14,500 \text{ rpm})$ and washed with Ultrapure (type I) water (Cuellar et al., 2018; Perez et al., 2019; Ramírez-Acosta et al., 2020).

Crosslinking of amine-terminal groups in the protein with aminopropyl groups on the surface of silanized SNPs or fullerenol facilitated by the addition of glutaraldehyde as the crosslinking agent enabled immobilization of OmpA on SNPs or fullerenol (Figure 3; Supplementary Figure S2) (López-Barbosa et al., 2019; Rangel-Muñoz et al., 2020). Briefly, 100 mg of silanized SNPs or fullerenol were suspended in 30 mL of ultrapure (type I) water and sonicated for 10 min (Elmasonic EASY 60H, 37 kHz, 150 W). This was followed by adding 2 mL of glutaraldehyde 2% (v/v) and by letting the mixture left to react in an orbital shaker for 1 h at 220 rpm. Then, OmpA (30 mg OmpA/100 mg functionalized nanomaterial) was added and shaken for 24 h at 220 rpm. The obtained nanobioconjugates were centrifuged (Z-216, Hermle Labortechnik GmbH, German) (4 \times 20 min, 14,500 rpm) and washed with ultrapure (type I) water.

2.7 Labeling of nanobioconjugates with rhodamine B

For confocal microscopy evaluation of cellular uptake and endosomal escape, the nanobioconjugates were labeled with the fluorescent probe rhodamine B. This was accomplished through the formation of amide bonds between the carboxylate group of rhodamine B and the free amine groups of nanobioconjugates.



Briefly, under dark conditions, 15 mg of EDC, 7.5 mg of NHS, and 1 mL of DMF were added to 5 mL of type I ultrapure water. Subsequently, 2 mg of rhodamine B was added, and the solution was heated to 40°C for 15 min with continuous magnetic stirring. This enables the activation of the carboxylate groups of rhodamine B to form amide bonds with the free amine groups of nanobioconjugates. The mixture was then allowed to cool to room temperature before being combined with 50 mg of nanobioconjugates. To prevent photobleaching, it was stirred for 24 h at 220 rpm using a shaker at room temperature and in complete darkness. The labeled nanobioconjugates were centrifuged (Z-216, Hermle Labortechnik GmbH, German) (20 min, 14,500 rpm) and washed several times with ultrapure (type I) water until no rhodamine B was detected in the supernatant (López-Barbosa et al., 2019).

2.8 Characterization of the nanobioconjugates

Infrared spectra were collected from 4,000-500 cm⁻¹ with a spectral resolution of 2 cm⁻¹ using a spectrometer ALPHA II FTIR Eco-ATR (Bruker Optik GmbH, Ettlingen, Germany) and an IRAffinity-1 spectrometer (Shimadzu Corporation). The diameter and ζ hydrodynamic potential of the nanobioconjugates were determined via Dynamic Light Scattering (DLS) and Electrophoretic Moobility (Zeta-Sizer Nano-ZS; Malvern Instruments, Malvern, UK). Thermogravimetric analysis (TGA, TA Instruments, New Castle, DE, United States) was used to estimate the amount of material conjugated to the SNPs and the fullerenol, implementing a linear temperature ramp at a rate of 10°C min⁻¹ from 25°C to 890°C under an inert atmosphere. Focused Ion Beam Scanning Electron Microscope (TESCAN LYRA3 FIB-SEM, Czech Republic) and Transmission Electron Microscope (TEM, FEI TECNAI G2 F20 Super Twin TMP, Hillsboro, OR, United States) were used to obtain information on the size, composition, and morphology of the nanomaterials. XPS spectra were obtained using a SPECS near-ambient pressure X-ray photoelectron spectrometer (NAP-XPS) with a PHOIBOS 150 1D-DLD analyzer, using a monochromatic source of Al-Ka (1,486.7 eV, 13 kV, 100 W) (SPECS GmbH, Berlin, Germany). The X-ray source and monochromator were aligned to get a 0.49 eV peak-resolution under a vacuum pressure of the chamber below 10⁻⁹ m bar. The samples were previously mounted on a non-conductive tape. The control of surface potential was achieved by an electron flood gun at 3 kV over a tantalum mesh with a nominal aperture of 0.43 mm. The spot size was 200 nm of diameter, the energy pass was fixed at 20 eV and the scan number for the high-resolution measurements was 20. The signals were calibrated to a binding energy of 284.6 eV for adventitious carbon and a Ta4f7/2 peak from the tantalum mesh was employed as reference. XPSPeak software was used for fitting the XPS spectra using a Shirley-type single-peaks background with a simultaneous GL peak-shape of 30% and full-width at half maximum (FWHM) data from literature.

Delivery of nanobioconjugates in mammalian cells was monitored using a confocal laser scanning microscope (Fluoview FV1000, Olympus, Tokyo, Japan). The images were obtained with a UPLSAPO 20x/0.75 NA objective and a PlanApo \times 60/1.35 NA objective. Excitation/Emission wavelengths of 405/422, 488/520, and 559 nm/575 nm were used to detect DAPI (nuclei), LysoTracker green (acidic organelles: endosomes/lysosomes), and rhodamine B (nanobioconjugates), respectively. Colocalization within biologically relevant regions of interest (ROIs) was analyzed using the plugin Coloc 2 of the Fiji[®] software (https://imagej.net/Fiji/Downloads). At least 30 images were taken for each treatment (about 10 cells per image were analyzed).

2.9 *In vitro* analysis of the nanobioconjugates' hemolytic effect

Hemolysis is the rapid destruction of erythrocyte membranes, which results in the release of intra-erythrocyte contents into the blood plasma. The hemolytic activity of the nanobioconjugates was determined using the method described by previously Muñoz-Camargo et al. (2018). Briefly, blood from healthy donors was collected in BD Vacutainer® blood tubes using EDTA as an anticoagulant. The samples were obtained with the approval of the Ethical Committee at the Universidad de los Andes (minute number 928-2018). Blood was centrifuged at 1,800 rpm for 5 min (Micro Centrifuge Z 360, Hermle Laboratories GmbH) to collect red blood cells, and the hematocrit level (lower layer, red) and plasma (upper layer, yellow) were marked. The plasma was then removed, and the tube was refilled to the mark with 150 mM NaCl, inverting the tube gently to mix, and centrifuged again. Subsequently, the supernatant was removed and replaced with PBS (Phosphate Buffered Saline) (1X). A red blood cells stock was prepared by adding 1 mL of isolated red blood cells $(4.3 \times 10^6 \text{ red blood cells})$ μL^{-1}) in 9 mL of PBS (1X). Serial dilutions of nanobioconjugates (300, 150, 75, 37.5, and 18.75 μ g mL⁻¹ in PBS) were prepared for the test in a 96-well microplate. 100 µL of each treatment and 100 µL of the diluted red blood cells were incubated for 1 h at 37°C and 5% CO₂. Finally, the microplate was centrifuged at 1,800 rpm for 5 min, and 100 µL of each supernatant was measured (oxyhemoglobin, 450 nm) in a microplate reader spectrophotometer (Multiskan[™] FC, Thermo Fisher Scientific Inc., United States). PBS (1X) and Triton X-100 (1%) were used as negative and positive controls, respectively.

2.10 *In vitro* assessment of nanobioconjugates' blood coagulation effect

The effect of the nanobioconjugates on platelet aggregation was tested using platelet-rich plasma (PRP). Blood was obtained from healthy donors in BD Vacutainer[®] tubes, anticoagulated with sodium citrate. PRP was obtained by centrifugation of a human blood sample at 1,000 rpm for 15 min (Micro Centrifuge Z 360, Hermle Labortechnik GmbH), and the PRP was collected and transferred to a fresh tube. Serial dilutions of nanobioconjugates (300, 150, 75, 37.5, and 18.75 µg mL⁻¹ in PBS) were prepared for the test in a 96-well microplate. Subsequently, 100 µL of each treatment and 100 µL of PRP were incubated for 15 min at 37°C and 5% CO₂. Thrombin (6U) was used as a positive control, while PBS (1X) was a negative reference. Finally, the aggregation was measured by optical density (OD) at 620 nm in a microplate reader spectrophotometer (MultiskanTM FC, Thermo Fisher Scientific Inc., United States) (Lopez-Barbosa et al., 2020).

2.11 MTT cytotoxicity test

The MTT (3-[4,5-dimethylthiazol-2-yl]-2.5 diphenyl tetrazolium bromide) assay is based on the metabolic reduction of MTT into formazan crystals by viable cells (Meerloo et al., 2011). Briefly, Vero cells (ATCC^{*} CCL-81) and THP1 cells (ATCC^{*} TIB-

202) were plated in 96-well culture plates in DMEM $(1.0 \times 10^6$ cells $100 \ \mu L^{-1}$ per well) and incubated at 37°C and 5% CO₂ for 24 h. Culture media was removed from wells, and DMEM 1% penicillin/ streptomycin (90 μ L) (without FBS) was added to each well. Subsequently, 100 μ L of each treatment (300, 150, 75, 37.5, and 18.75 μ g mL⁻¹ in PBS) were added and incubated at 37°C, 5% CO₂ for 24 h, and 48 h. The medium was removed, and DMSO (500 μ L) was used to dilute formazan crystals. Absorbance was read at 595 nm (reference 650 nm) in a microplate reader spectrophotometer (MultiskanTM FC, Thermo Fisher Scientific Inc., United States) and compared to the controls (Lopez-Barbosa et al., 2020).

2.12 Cell translocation and endosome escape

Vero Cells were seeded in a sterile glass slide previously placed in a 6-well microplate and incubated in DMEM medium supplemented with 10% (v/v) FBS at 37°C and 5% CO_2 for 24 h. Next, cells were exposed to fluorescently labeled nanobioconjugates (18.75 µg mL⁻¹), and the samples were incubated for 30 min and 4 h at 37°C and 5% CO₂. Supplemented culture medium was removed, and then, the cells were washed three times with DMEM medium and exposed for 10 min to DAPI (1 µL, 1:1,000) used to stain nuclei, and Lysotracker Green DND-26 (0.1 µL, 1:10,000) that labels acidic organelles (lysosomes/endosomes) before capturing confocal images (López-Barbosa et al., 2019). THP-1 Cells exposed to fluorescently labeled nanoconjugates (18.75 µg mL⁻¹) were incubated for 30 min and 4 h at 37°C and 5% CO₂. Then, the samples were exposed for 10 min to DAPI (1 µL, 1:1,000) and Lysotracker Green DND-26 (0.1 µL, 1: 10,000) before capturing confocal images (López-Barbosa et al., 2019).

2.13 Statistical analysis

Values (Hemolysis, platelet aggregation, cell viability) are expressed as the means \pm SDs of triplicates. Significance tests were analyzed by nonparametric—the normality of data distributions was assessed using the Shapiro–Wilk test—one-way ANOVA (Kruskal–Wallis test) and Dunn's multiple comparison test, using the GraphPad Prism 8.0.1[®] software (GraphPad Software, La Jolla, CA, United States). *p* values <0.05 were considered statistically significant.

3 Results and discussion

3.1 Physicochemical characterization of SNPs and nanobioconjugates based on SNPs

Figure 4A shows a schematic of the chemical structure of silanized SNPs and BUF-II-PEG₁₂-SNPs, and OmpA-SNPs nanobioconjugates. Figure 4B compares the FT-IR spectra of bare SNPs, silanized SNPs, free OmpA, free BUF-II, and nanobioconjugates. The bare SNPs exhibit distinctive absorptions at around 1,100 cm⁻¹ (Si-O *st as*, asymmetrical stretching), 970 cm⁻¹



Spectroscopic and thermal analyses of silanized SNPs and the nanobioconjugates (A) Schematic of the chemical structure of silanized SNPs and the nanobioconjugates (B) FT-IR spectra of (1) bare SNPs, (2) silanized SNPs, (3) BUF-II-PEG₁₂-SNPs nanobioconjugates, (4) free BUF-II, (5) OmpA-SNPs, and (6) free OmpA (C) DLS histogram for the size intensity distribution (D) TGA thermogram of SNPs, silanized SNPs, and nanobioconjugates (E) SEM and TEM images of the silanized SNPs, and nanobioconjugates.

(Si-OH st as), and 801 cm⁻¹ (Si-O st sy, symmetrical) (Pretsch et al., 2009; Edrissi et al., 2011). New bands were observed at 2,925 cm⁻¹ (C-H st as), 2,852 cm⁻¹ (C-H st sy), and 1,639 cm⁻¹ (N-H b, bending vibration), evidencing the presence of propylamine groups on the surface of the silanized SNPs (Shafqat et al., 2019). On the free OmpA, OmpA-SNPs, free BUF-II, and BUF-II-PEG₁₂-SNPs spectra, the presence of amide vibrational modes known as amide I (1,700-1,600 cm⁻¹) and amide II (1,570-1,540 cm⁻¹) and other vibrations specific that are absent on bare SNPs suggests correct conjugation of protein or peptide (Pretsch et al., 2009; Tatulian, 2013; López-Barbosa et al., 2019). The amide I mode is generated mostly by contributions of the C=O st, CN st, CCN ob (out-of-plane bending), and by the NH ib (in-plane bending) modes (Tatulian, 2013). The amide II mode includes contributions of the NH ib, CN st, CO ib, CC st, and NC st (Tatulian, 2013). Other vibrational modes of the amide group of protein or peptide and crosslinking agents overlap with the vibrational modes of SNPs. Figure 4C shows particle size distribution by intensity determined by DLS. Bare SNPs exhibited a mean hydrodynamic diameter (Z-average size) of 41 nm with a Polydispersity Index (PdI) of 0.1. A ratio of 95% of TEOS and 5% of APTES produced two populations of silanized SNPs with mean hydrodynamic diameters around 40 and 220 nm (Z-average size: 176 nm, PdI: 0.2). Regarding this, Li et al. (2019) demonstrated that the total uptake of SNPs in Hela cells was higher in co-exposures of large and small SNPs — 50/ 100 and 80 nm/150 nm-than in single exposures of the same. It can also be observed that the Z-average size of the SNPs increased after peptide and protein conjugation for BUF-II-PEG₁₂-SNPs (Z-average size: 212 nm, PdI: 0.05) and OmpA-SNPs (Z-average size: 230 nm, PdI: 0.05). There is no consensus regarding the optimal size that maximizes cellular uptake and maintains cell viability. A number of experimental studies indicate that particle size reduction does not necessarily increase cellular uptake (Barisik et al., 2014). However, nanocarriers based on NPs in the size range of 10-200 nm are frequently used to facilitate the delivery of cargoes at the cellular level. These nanocarriers are not easily excreted by any of the physiological systems designed for this purpose and therefore reach target organs and tissues in sufficient concentrations (Selby et al., 2017; Chenthamara et al., 2019). Thermogravimetric analysis (TGA) was used to estimate the amount of material conjugated to the SNPs (Figure 4D). The bare SNPs exhibited excellent thermal stability, losing only about 1.1% of their weight when heated from room temperature to 890°C. In contrast, weight loss was observed for silanized SNPs and nanobioconjugates in three temperature ranges (silanized SNPs: room temperature to 140°C, 140°C-450°C, and >450°C; nanobioconjugates: room temperature to 140°C, 140°C-340°C, and >340°C). Silanized SNPs, BUF-II-PEG₁₂-SNPs, and OmpA-SNPs showed a first weight loss of 8.5%, 8.8%, and 6.2%, respectively, that can be attributed to water loss. A second weight loss of 5.6% was observed for the silanized SNPs and OmpA-SNPs, whereas for BUF-II-PEG₁₂-SNPs, it was 8.3%. These losses can be assigned to the presence of non-hydrolyzed ethoxy groups of APTES and residual alcohol within the silica nanostructure (Kunc et al., 2019). Finally, the weight loss at the highest temperatures can be assigned to the loss of aminopropyl groups (7.2%) for the silanized SNPs, and the detachment of BUF-II, OmpA, and crosslinking agents in the case of the BUF-II-PEG₁₂-SNPs (11.0%) and OmpA-SNPs (11.6%) nanobioconjugates (López-Barbosa et al., 2019; Perez et al., 2019). SEM and TEM images were consistent with the data obtained by DLS regarding the size and the presence of two size populations of particles. In addition, it can be observed that the nanoparticles have a predominantly spherical morphology. Apparently, after conjugation, the roughness of the nanoparticles changes; this could also affect their interaction with cells and their loading capacity (Niu et al., 2015; Alan et al., 2020) (Figure 4E). Additionally, ζ potential is indicative of the stability of the suspension; if all the particles in suspension have ζ potentials above +25 mV or below -25 mV, they repel each other, and therefore show no tendency for aggregation, coagulation, or flocculation (Shnoudeh et al., 2019). The ζ potential measurements of the SNPs indicate that in aqueous media-pH close to 7- it acquires a negative surface charge of -37.6 ± 4.91 mV; this value indicates good SNPs stability in water. The ζ -potential reached values of 4.41 ± 3.27 mV, 7.34 ± 3.36 mV, and 18.1 \pm 5.13 mV for the silanized SNPs, and BUF-II-PEG₁₂-SNPs, and OmpA-SNPs nanobioconjugates (in aqueous media-pH close to 7), respectively. At physiological pH, these nanobioconjugates tend to precipitate.

The chemical surface characterization of the nanobioconjugates was evaluated by XPS. The detailed experimental set-up carried out for the samples is shown in Section 2.8. Here, SNPs and fullerenol nanocarriers were considered. Figure 5 shows the high-resolution (HR) spectra for the SNPs and the corresponding nanobioconjugates under examination. The peak components from the decomposition analysis are denoted from high to low binding energy, and colored zones clearly distinguish them. The binding energy (BE) values for all components that are part of the overall fitting, marked as a red line over black dots related to the experimental recorded data, are shown in Table 1. Starting at the C1s core-level, four mean sub-peaks for functionalizing samples were fitted, which correspond to O-C-O/C=O (red); C-O/C-N (blue); C-C (green) and C-Si (magenta) bonds. For the silanized SNPs, there were no highly oxidized species, which clearly indicate the successful conjugation of OmpA and BUF-II on the NPs' surface. Since these biomolecules and their intermediate states are too complex due to their chemistry and molecular weight (MW), it is not possible to establish a stoichiometric ratio between the species; nevertheless, the counts (Y-axis) for each core-level were normalized prior to their placement on the plots. Conversely, a qualitative analysis can be done. As a result, it is possible to determine that the oxidizing species for C1s core-level in the OmpA-SNPs system are greater than those in the BUF-II-PEG12-SNPs nanobioconjugates. In contrast, the BUF-II-PEG12-SNPs exhibited a high (C-N/C-O)/C-C ratio due to their low molecular weight and high C-N/C-O terminal bonds. Consistent C-Si bonds were found in the studied systems, allowing us to conclude that the SNPs were properly silanized and that the biomolecules are bound to the inorganic nanoparticles via C-Si-O covalent bonds. The O1s core-level was deconvoluted into four mean sub-peak components associated with chemisorbed OHmolecules: O-C-O/Si-O, C-O, and O=C species. Slight shifts in binding energies and high similitudes of the integral intensity of every peak as calculated from the area under the curve were observed. Moreover, if we compare the overall peak intensity with that of C1s for each compound, the corresponding ratio for the nanobioconjugates is lower than for the SNPs. This is most likely due to the lower concentration of C/O species on the surface of



XPS spectra of the C1s, O1s, N1s, and Si2p (left to right) core-level regions of silanized SNPs, BUF-II-PEG12-SNPs nanobioconjugates, and OmpA-SNPs nanobioconjugates samples (bottom to top). Peak components for the XPS lines are differentiated by colors from high to low binding energy values (left to right).

TABLE 1 Binding energy (BE) of the different XPS lines for C1s, O1s, N1s, and S2p peak components from silanized SNPs, BUF-II-PEG12-SNPs nanobioconjugates, and OmpA-SNPs nanobioconjugates samples.

Sample	C1s (BE- eV)	O1s (BE- eV)	N1s (BE- eV)	Si2p (BE- eV)
Silanized SNPs	283.4	530.3	400.8	102.0
	284.6	531.7	_	103.2
	285.5	532.7	_	104.1
	286.5	533.6	_	104.9
BUF-II-PEG ₁₂ -SNPs	283.3	530.5	397.2	101.8
	284.6	531.6	400.1	102.8
	285.5	532.5	402.5	103.7
	286.8	533.3	—	104.6
SNPs-OmpA	283.8	530.5	396.9	102.3
	284.6	531.7	399.8	103.2
	285.4	532.6	401.9	104.0
	286.1	533.,3	_	104.7

SNPs, where only the APTES chain is present. As peptide and protein are conjugated, the proportion of atomic C species increases dramatically, as does the C/O ratio. Now passing through the N1s core-level, three mean sub-peak components for nanobioconjugates were fitted. In the case of silanized SNPs, a weak signal of nitrogen from the conjugated APTES can be assigned to a primary amine

(Talavera-Pech et al., 2016). For the nanobioconjugates, protonated amines seem to be located at higher binding energies, followed by O-C-N and N=N-/N-H- bonds. The Si2p sub-peaks components corresponding to SiOx (red), Si-O- (blue), Si-O-C- (green) and Si-C- (magenta) bonds are shown from left to right for bare SNPs and nanobioconjugates (Talavera-Pech et al., 2016). Slight peak



Spectroscopic and thermal analyses of fullerenol and the nanobioconjugates (A) UV–VIS spectra of C_{60} in toluene and aqueous solution of fullerenol (B) Schematic of the chemical structure of fullerenol and the nanobioconjugates (C) FT–IR spectra of (1) fullerene, (2) fullerenol as produced (with TBAH residues), (3) purified fullerenol, and (4) silanized fullerenol (D) FT–IR spectra of (1) BUF–II–PEG12–F nanobioconjugates, (2) free fullerenol, silanized fullerenol, and nanobioconjugates (F) DLS histogram for the size intensity distribution (G) TEM images of the fullerenol, and nanobioconjugates. shifts and intensity changes can be attributed to conjugation of OmpA and BUF-II and are likely related to conformational changes upon conjugation.

3.2 Physicochemical characterization of fullerenol and fullerenol-based nanobioconjugates

Figure 6A shows a schematic of the chemical structure of silanized fullerenol, OmpA-F, and BUF-II-PEG₁₂-F nanobioconjugates. Figure 6B shows UV-visible absorption spectra of fullerene C60 in toluene and fullerenol in water. C60 fullerene dissolved in toluene has characteristic absorption bands with maxima at 283, 335, and 408 nm, followed by a broad absorption band in the range of 430-650 nm with reduced absorptions for the blue region and red; this combination gives the compound its distinctive purple color (Ajie et al., 1990). Fullerenol dissolved in water is yellow and almost transparent in the visible region due to its considerable perturbation of the π conjugation upon hydroxylation (Kokubo et al., 2008). Figure 6C compares the FT-IR spectra of (1) fullerene, (2) fullerenol as synthesized (with TBAH residues), (3) purified fullerenol, and (4) silanized fullerenol. Fullerene C60 has four active infrared modes at 1,429, 1,182, 573, and 525 $\rm cm^{-1}$ due to C-C bonds (Krätschmer et al., 1990). In the as-synthesized fullerenol, the two peaks observed at 2,963 and 2,873 cm⁻¹ (C-H st) were attributed to residual TBAH (Kokubo et al., 2011). Purified fullerenol showed a broad band at around 3,424 cm⁻¹(O-H st) and four characteristic bands at 1,598 cm⁻¹ (C=C st), 1,410 cm⁻¹ (O-H b), 1,352 cm⁻¹ (C-O-H b), and 1,112 cm⁻¹ (C-O st), which agree well with previously reported data (Kokubo et al., 2011; Ravelo-Nieto et al., 2020). Silanization was confirmed by the presence of new bands at 2,964 cm⁻¹ (C-H st as), 2,934 cm⁻¹ (C-H st sy), 2,875 cm⁻¹ (H-C (-N) st), 1,564 cm⁻¹ (N-H b), and 1,344 cm⁻¹ (C-N st), which can be attributed to propylamine groups. Moreover, absorption bands at 1,653 cm⁻¹ (N-H b), 1,110 cm⁻¹ (Si-O st), 1,052 cm⁻¹ (Si-O-Si st), and 690 cm⁻¹ (Si-C st) overlap with the vibrational modes of fullerenol (Cuellar et al., 2018). Figure 6D compares the FT-IR spectra of (1) OmpA-F, (2) free OmpA, (3) BUF-II-PEG₁₂-F, and (4) free BUF-II (Shafqat et al., 2019). The free OmpA, OmpA-F, free BUF-II, and BUF-II-PEG₁₂-F spectra showed the amide I and amide II vibrational modes along with other specific vibrations that are absent in the nonfunctionalized fullerenol. However, these signals overlap with the vibrational modes of fullerenol. Thermal stability of fullerenol and nanobioconjugates was studied by TGA (Figure 6E). TGA results of purified fullerenol show three main a weight loss stages: room temperature to 100°C, 100°C-570°C, and >570°C. The initial weight loss (~8.1%) corresponds to dehydration of the samples (Kokubo et al., 2011). The second weight loss (~54.0%) corresponds to the dehydroxylation before the structural degradation of the fullerene nucleus that occurs at temperatures above 570°C (~37.9% residual weight) (Goswami et al., 2004). Then, using the method described by Goswami et al.(2004), the number of -OH groups per fullerene could be estimated at 30, which is similar to results reported by others previously (Kokubo et al., 2011; Kovač et al., 2018; De Santiago et al., 2019). Four main weight loss steps are observed in silanized fullerenol, and BUF-II-PEG₁₂-F, OmpA-F

nanobioconjugates: room temperature to 120°C, 120°C-340°C, 340°C-570°C and >570°C. Silanized fullerenol, BUF-II-PEG₁₂-F, and OmpA-F presented a first weight loss of ~12.1%, ~11.5%, and ~12.1%, respectively. These can be attributed to the dehydration of the samples. The second weight loss of ~34% was observed for the silanized fullerenol, whereas for BUF-II-PEG₁₂-F and OmpA-F were ~21% and ~22%, respectively. Silanized fullerenol, BUF-II-PEG₁₂-F, and OmpA-F presented a third weight loss of ~15.4%, ~14.1%, and ~15.0%, respectively. These losses in the temperature range of 150°C-570°C can be assigned to the decomposition of aminopropyl groups for the silanized fullerenol and the detachment of the aminopropyl groups, conjugating agents and the BUF-II and OmpA for the nanobioconjugates (Goswami et al., 2004; Cuellar et al., 2018; Perez et al., 2019). Figure 6F shows the particle size distribution by intensity determined by DLS. The fullerenols should have a diameter of ~1.0 nm but tend to form clusters in water easily (Brant et al., 2007; Kokubo et al., 2011). The synthesized fullerenol, redispersed by sonication, exhibited two populations of clusters with mean hydrodynamic diameters at around 2 nm and 14 nm (Z-average size: 8 nm, PdI: 0.2). After peptide and protein conjugation, the polydispersity of the samples increased and rendered them unsuitable for DLS measurements; consequently, we performed TEM analysis. (Figure 6G). Fullerenol TEM images were consistent with DLS data regarding cluster size and the presence of two cluster population sizes. Furthermore, a change in the morphology of the nanobioconjugates is evidenced after peptide and protein immobilization, as well as aggregate formation. The aggregation may be attributable to the use of glutaraldehyde, a bifunctional reagent with a propensity for uncontrolled polymerization during the conjugation process (Hermanson, 2013). The ζ potential measurements of the fullerenol indicate that in aqueous media-pH close to 7- it acquires a negative surface charge of -20.4 ± 7.47 mV. The ζ potential varied to $-12.9 \pm 0.40 \text{ mV}$ and $-19.9 \pm 0.65 \text{ mV}$ for BUF-II-PEG₁₂-F and OmpA-F nanobioconjugates in aqueous media-pH close to 7 -, respectively. These nanobioconjugates tend to precipitate at physiological pH.

Figure 7 shows the high-resolution (HR) spectra of fullerenol and the corresponding nanobioconjugates under examination. The decomposed peak components are labeled from high to low binding energy and depicted by colored zones. The color code employed is similar to that utilized for SNPs. Table 2 presents the binding energy (BE) values for all components integrated in the overall fitting, represented as a red line over black dots related to the experimental recorded data. Starting with the C1s core-level, the deconvolution revealed a sub-peak at higher energies and the presence of the C-C cage at 284.6 eV (Nurzynska et al., 2022). Following silanization and conjugation of BUF-II or OmpA, the peak at the lowest energy became weak or null. This may be due to the photoelectrons' inability to escape the outermost surface layer. In contrast, the pristine fullerenol sample exhibited an energy shift, most likely due to the presence of highly oxidized species associated with the hydroxyl binding onto conjugated pi bonding systems. A modified fullerenol energy sub-peak was identified at 289 eV, which can be also attributed to highly oxidized bonds such as O-C-OO (marine blue) (Nurzynska et al., 2022). The presence of these bonds may result from the chemisorption of oxygen molecules onto C-O-



XPS spectra of the C1s, O1s, N1s and Si2p (left to right) core-level regions of Fullerenol, Silanized fullerenol (F), BUF-II-F and OmpA-F nanobioconjugates samples (bottom to top). Peak components for the XPS lines are differentiated by colors from high to low binding energy values (left to right).

TABLE 2 Binding energy (BE) of the different XPS lines for C1s, O1s, N1s and S2p peak components from Fullerenol w/o silanization, F, BUF-II-F and OmpA-F
nanobioconjugates samples.

Sample	C1s (BE- eV)	O1s (BE- eV)	N1s (BE- eV)	Si2p (BE- eV)
Fullerenol w/o silanization	283.3	529.8	_	_
	284.6	531.1	—	—
	285.9	532.2	—	—
	287.5	533.2	—	—
Silanized fullerenol (F)	282.9	530.7	397.3	101.6
	284.6	531.7	400.0	102.6
	287.1	532.8	402.3	103.4
	289.6	533.9	—	104.3
BUF-II-F	282.6	531.0	397.6	102.0
	284.6	532.2	400.1	102.9
	285.8	533.1	402.5	103.6
	287.5	534.1	—	104.5
	289.8	_	—	_
OmpA-F	284.6	531.1	396.7	101.3
	285.7	532.3	400.0	102.2
	287.3	533.2	402.2	103.1
	289.5	534.,2	_	104.1



24 and 48 h (**B**) *In vitro* evaluation of the hemocompatibility. Assessment of the hemolytic effect of nanobioconjugates (Positive control: Triton X-100, negative control: PBS. In all cases, hemolysis was below 3%; thus, the nanobioconjugates are not hemolytic; and Assessment of nanobioconjugates effects on blood coagulation (Positive control: Thrombin, negative control: PBS). There is no significant percent platelet aggregation induced by the nanobioconjugates.

radicals arising from the cleavage of C=C bonds and/or from C-Obonds present in the activated hydroxyls before silanization. Crucially, BUF-II-PEG₁₂-F and OmpA-F nanobioconjugates exhibited a clearly differentiated C1s high-resolution spectra, confirming the successful conjugation of BUF-II and OmpA. The decreasing C-C/(C-O/C-N) ratio upon conjugation provided further evidence of the superior conjugation efficiency of BUF-II.

Concerning the O1s-core level, a sharper peak with a slight shift to higher energy was observed for silanized and nanobioconjugates samples compared to the reference. The larger full-width of half-maximum (FWHM) value of the pristine sample is likely associated with the overall electric field's spread on the outer C_{60} -cage surface due to defects that are absent on functionalized samples. This permits the favored ejection of O1s-photoelectron at lower kinetic energies. No evidence of work function alteration due to surface charge artifacts was found since the C1s-main peak of the cage was located at 284.6 eV. Nevertheless, the C1s-peak of the reference was also broader, without any evidence of a change in the peak asymmetry compared to the functionalized samples. In contrast, the main three N1s-subpeak components were detected for silanized and nanobioconjugates samples, with a higher intensity detected for the former due to the protonated amine species of the covalently attached APTES molecules. This contrasts with the observations for the SNPs discussed above. The area under the curve for the nitrogen binding energy was lower for the nanobioconjugates compared to the silanized samples due to the lower C/N ratio after conjugation of the peptide and protein molecules. Finally, the main Si2p-subpeak components provided further evidence of the successful silanization of fullerenol.

3.3 Biocompatibility

Biocompatibility is a crucial property in the development of nanocarriers for biomedical applications. A material is considered biocompatible if it does not elicit an undesired response from the organism. Therefore, the assessment of biocompatibility is a fundamental step in the design and



development of nanocarriers for drug delivery and diagnostic purposes. It is imperative to ensure that the nanocarriers are not toxic to the body and do not cause any adverse reactions (Soares et al., 2018). In order to ensure the biocompatibility of nanomaterials, multiple tests are required as per established standards such as the ISO 10993 series and ASTM F1903. Hemocompatibility and cytotoxicity tests are among the several necessary evaluations. The hemolytic properties, effects on blood coagulation, and cytotoxicity of the tested samples were assessed *in vitro*.

As shown in Figure 8A, cell viability was evaluated in THP-1 cells—a human leukemia monocytic cell line—and Vero cells—a monkey kidney epithelial line—after 24 and 48 h of exposure to the SNPs-based treatments. The outcomes demonstrated a concentration-dependent decrease in cell viability for all treatments and cell lines. Notably, at low doses of 18 and 37 μ g/mL, no significant reduction in cell viability was observed for either cell type, implying the treatments' safety profile at lower concentrations. Moreover, it was observed that the cytotoxic potential of OmpA-SNPs was comparatively lower than that of

BUF-II-PEG₁₂-SNPs. The viability of cells treated with OmpA-SNPs nanobioconjugates remained above 70% (dotted line) even at concentrations as high as 75 µg mL-1. Conversely, at the same concentration, the viability of THP-1 and Vero cell lines treated with BUF-II-PEG₁₂-SNPs nanobioconjugates was found to decrease. (International Organization for Standardization., 2009). In all cases, the covalent conjugation of BUF-II peptide and OmpA protein to these nanostructured materials involves the use of surface spacers (APTES, amine-PEG₁₂-propionic acid, glutaraldehyde, EDC, NHS), which has rendered the nanobioconjugates less cytotoxic than the bare SNPs. Surface functionalization modified the properties of the nanoparticles—e.g., the Z-average size, the ζ potential, the roughness-and thus the interactions between nanoparticles and biological components, such as proteins and cell membranes, ultimately reducing cytotoxicity (Kim et al., 2013). Figure 8B shows in vitro evaluation of the hemocompatibility. The treatments revealed platelet aggregation values between 2% and 16% above the negative reference-values higher than 20% are considered that induce platelet aggregation (dotted line) (Potter et al., 2018). The silanized SNPs or nanobioconjugates induced no



times (30 min and 4 h).

significant hemolytic effect, and the hemolysis values remained below 3% — Percent hemolysis less than 2 means the test sample is not hemolytic (dotted line); 2%–5% hemolysis means the test sample is slightly hemolytic; and >5% hemolysis means the test sample is hemolytic (Neun et al., 2018).

Similarly, in the case of nanobioconjugates that rely on fullerenol, cell viability remained uncompromised in both cell types, even at low doses of 18 and 37 µg mL-1 across all treatments. There was no significant reduction observed in the viability of the cells (Figure 9A). Figure 9B shows *in vitro* evaluation of the hemocompatibility. The results of the treatments showed platelet aggregation values that were 2%-15% higher than the negative reference in the fullerenol and BUF-II-PEG₁₂-F treatment. Conversely, there was no significant difference in platelet aggregation between the negative control and the OmpA-F nanobioconjugates treatment (Potter et al., 2018). The silanized fullerenol or nanobioconjugates did not induce a significant hemolytic effect, as evidenced by the hemolysis values remaining below 3% (Neun et al., 2018).

We recognize that the dosages used in our work may not exhibit toxicity towards target cells, such as cancer or infected cells, and that altering the dosage might change the mechanism of cellular uptake. As such, future studies should investigate the potential biological actions of these nanobioconjugates using a drug model to better understand their efficacy and safety in diverse cellular contexts. This will involve characterizing how the nanobioconjugates and their cargoes are trafficked inside cells and determining the appropriate cargo release concentrations for specific cell lines. Our current research serves as a foundation for developing conjugation strategies with known cell penetration agents and for exploring the potential of silica nanoparticles and fullerenol as nanostructured supports in targeted drug delivery applications.

3.4 Cellular uptake and endosomal escape

Figure 10 shows the cellular uptake of silanized SNPs, OmpA-SNPs, and BUF-II-PEG₁₂-SNPs nanobioconjugates by Vero cells. Rhodamine B-labeled nanobioconjugates (red) were observed to be homogeneously distributed within the cells, without significant penetration of the cell nucleus (blue). The colocalization of the nanobioconjugates with acidic organelles, such as endosomes/



(B) Confocal microscopy images of effective cellular uptake of silanized SNPs and OmpA-SNPs in THP-1 cells. A zoomed view of individual cells is shown on the insets in the right panels (B) Confocal microscopy images of effective cellular uptake of silanized SNPs and OmpA-SNPs in THP-1 cells. A zoomed view of individual cells is shown on the insets in the right panels (C) Endosomal escape study via colocalization analysis (D) Intracellular area percentage coverage by silanized SNPs and the nanobioconjugates for THP-1 cells after the two exposure times (30 min and 4 h).

lysosomes, was determined quantitatively through correlation analysis, based on Pearson's coefficient (PC). The PC value ranges from 1 to -1, where 1 represents complete and positive correlation between the intensity of fluorescence signals, -1 denotes perfect but negative correlation, and 0 indicates no correlation (Adler and Parmryd, 2010; Dunn et al., 2011). Intracellular area percentage coverage by silanized SNPs and the nanobioconjugates was also determined.

CP values after 30 min in Vero cells were 0.76 ± 0.07 ; 0.71 ± 0.10 ; and 0.66 ± 0.10 for the silanized SNPs, BUF-II-PEG₁₂-SNPs, and OmpA-SNPs, respectively. These CP values decreased to 0.72 ± 0.09 ; 0.55 ± 0.12 ; and 0.54 ± 0.13 ; for the silanized SNPs, BUF-II-PEG₁₂-SNPs, and OmpA-SNPs, respectively, after 4 h of incubation. A reduced level of colocalization with the endosomal/lysosomal marker (green) is indicative of the propensity of nanobioconjugates to evade endosomes in Vero cells. While the precise mechanisms involved in endosomal escape are not fully understood, it is likely that this occurs either through the formation of temporary pores or via the proton sponge effect, as posited in prior studies (Cho et al., 2009; Cardoso et al., 2019; López-Barbosa et al., 2019; Lopez-Barbosa et al., 2020).

In the case of THP-1 cells (Figure 11), CP values after 30 min were 0.76 \pm 0.07; 0.70 \pm 0.06; and 0.90 \pm 0.05 for silanized SNPs, BUF-II-PEG₁₂-SNPs, and OmpA-SNPs, respectively. After 4 h of incubation, the CP values approached 0.69 \pm 0.77; 0.78 \pm 0.06; and 0.91 ± 0.05 ; for the silanized SNPs, BUF-II-PEG₁₂-SNPs, and OmpA-SNPs, respectively, indicating a low tendency of nanobioconjugates to escape from endosomes in THP-1 cells. This behavior holds potential significance for the investigation of enzyme replacement therapies in the management of lysosomal storage diseases. Such diseases necessitate periodic intravenous infusions of human recombinant lysosomal enzymes, produced through recombinant DNA techniques. Following administration of the treatment, the recombinant enzymes disperse throughout the tissues, undergo internalization by cells, and are directed to the lysosomal compartment for the purpose of substituting the deficient protein in the patients (Parenti et al., 2013). Finally, the percentages of the area covered by the nanobioconjugates were higher in THP-1 cells than in Vero cells and increased with incubation time.

Figure 12 provides evidence of the cellular uptake of OmpA-F and BUF-II-PEG₁₂-F nanobioconjugates by Vero cells. The rhodamine B-labeled nanobioconjugates (red) were observed to



Cellular uptake and endosomal escape of the fullerenol-nanobioconjugates in Vero cells (A) Confocal microscopy images of effective cellular uptake of BUF-II-PEG₁₂-Fnanobioconjugates in Vero cells. A zoomed view of individual cells is shown on the insets in the right panels (B) Confocal microscopy images of effective cellular uptake of OmpA-F nanobioconjugates in Vero cells. A zoomed view of individual cells is shown on the insets in the right panels (B) Confocal microscopy images of effective cellular uptake of OmpA-F nanobioconjugates in Vero cells. A zoomed view of individual cells is shown on the insets in the right panels (C) Endosomal escape study via colocalization analysis (D) Intracellular area percentage coverage by the nanobioconjugates for Vero cells after the two exposure times (30 min and 4 h).

be homogeneously distributed within the cells but did not significantly reach the cell nucleus (blue). In contrast to SNPbased nanobioconjugates, fullerenol-based nanobioconjugates tend to aggregate and form clusters, which could have significant implications for their biological applications. CP values after 30 min were 0.23 \pm 0.11; and 0.28 \pm 0.11 for BUF-II-PEG₁₂-F and OmpA-F, respectively. After 4 h of incubation, the CP values approached 0.37 ± 0.17 ; and 0.29 ± 0.08 ; for BUF-II-PEG₁₂-F and OmpA-F, respectively. CP values less than 0.5 indicate a low degree of colocalization between the nanobioconjugates and the endosomes/lysosomes. The extensive coverage of the cytoplasmic area by the particles provides evidence for the internalization of the nanobioconjugates. These results suggest that the mechanism of entry of the nanobioconjugates into Vero cells is likely nonendocytic.

In the case of THP-1 cells (Figure 13), CP values after 30 min were 0.83 ± 0.07 ; and 0.77 ± 0.11 for BUF-II-PEG₁₂-F and OmpA-F, respectively. These CP values decreased to 0.74 ± 0.16 ; and 0.74 ± 0.12 ; for BUF-II-PEG₁₂-F and OmpA-fullerenol, respectively, after 4 h of incubation. This suggests a tendency of BUF-II-PEG₁₂-F nanobioconjugates to escape from endosomes in THP-1 cells, which

is not observed for OmpA-F nanobioconjugates. The high coverage of the cytoplasmic area confirms the internalization of the nanobioconjugates.

In addition to the cell penetration and endosomal escape capabilities of our nanobioconjugates, it is essential to highlight their potential for targeted drug delivery to specific subcellular compartments. By functionalizing the nanobioconjugates with appropriate ligands, such as specific peptide sequences or small molecules, these drug delivery systems can be tailored to exhibit a high affinity for the desired organelle. This customization enables enhanced specificity in targeting organelles such as mitochondria or others of interest. Moreover, we acknowledge the potential of non-endocytic mechanisms as a more benefits straightforward route for targeting subcellular compartments. While endosomal escape is a crucial step in ensuring efficient delivery of cargo to the cytosol, non-endocytic routes might offer alternative advantages in achieving more targeted delivery to specific organelles. Further exploration of these strategies and the development of suitable ligands will be essential in optimizing the nanobioconjugates for specific therapeutic applications.



uptake of BUF-II-PEG₁₂-F nanobioconjugates in THP-1 cells. A zoomed view of individual cells is shown on the insets in the right panels (**B**) Confocal microscopy images of effective cellular uptake of OmpA-F nanobioconjugates in THP-1 cells. A zoomed view of individual cells is shown on the insets in the right panels (**C**) Endosomal escape study via colocalization analysis (**D**) Intracellular area percentage coverage by the nanobioconjugates for THP-1 cells after the two exposure times (30 min and 4 h).

4 Conclusion and outlook

In summary, our study demonstrates a comprehensive approach for immobilizing translocating biomolecules on SNPs and fullerenol. The success of this strategy was confirmed by a range of analytical techniques including FT-IR, TGA, DLS, Electrophoretic Mobility, SEM, TEM, and XPS. The resulting nanobioconjugates, including OmpA-SNPs, BUF-II-PEG₁₂-SNPs, OmpA-F, and BUF-II-PEG₁₂-F, exhibited high biocompatibility in both Vero and THP-1 cell lines. Moreover, our evaluations of hemolytic effects and platelet aggregation demonstrated their safety at the tested concentrations. Our confocal microscopy studies revealed efficient internalization of the different nanobioconjugates in both Vero and THP-1 cells, with notable differences in endosomal escape. In particular, OmpA-SNPs and BUF-II-PEG₁₂-SNPs showed a tendency to escape from endosomes in Vero cells, while remaining trapped in THP-1 cells. On the other hand, OmpA-F and BUF-II-PEG₁₂-F were effectively internalized by both cell lines, with a superior tendency to escape from endosomes in Vero cells. These findings are significant, as they provide evidence for the potential of our nanobioconjugates to enhance the stability and half-life of translocating biomolecules and cross biological membranes without affecting cell viability. The ability to develop highly tunable cargo delivery systems is crucial for meeting the needs of specific treatments and targeting cell or organelle types. Overall, our study highlights the promise of our nanobioconjugates as a platform for the development of innovative therapeutic approaches. In our forthcoming research, we aim to elucidate the intracellular trafficking mechanisms of these nanobioconjugates and their cargoes, as well as the targeted release of cargo within specific cell lines at precise concentrations.

Data availability statement

The original contributions presented in the study are included in the article/Supplementary Material, further inquiries can be directed to the corresponding author.

Ethics statement

Human blood samples were collected under the permission obtained from the ethics committee at Universidad de los Andes, minute number 928-2018 (17 September 2018). The patients/participants provided their written informed consent to participate in this study.

Author contributions

Conceptualization, JCC, ER-N, and AD-R; Methodology, data curation, and data analysis ER-N, JC, CO, AD-R and JCC; Formal analysis and investigation, ER-N, JC, PR, LR-G, VQ; Validation, CM-C, LR, JCC, and AD-R; Writing—original draft preparation, ERN; Writing—review and editing, JCC, JC, and AD-R; Supervision, CM-C, LR, JCC, and AD-R. All authors have read and agreed to the published version of the manuscript.

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Supplementary material

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fbioe.2023.1184973/ full#supplementary-material

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