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Extracellular vesicles in cardiovascular diseases: pathogenic mediators, diagnostic tools, and therapeutic vectors

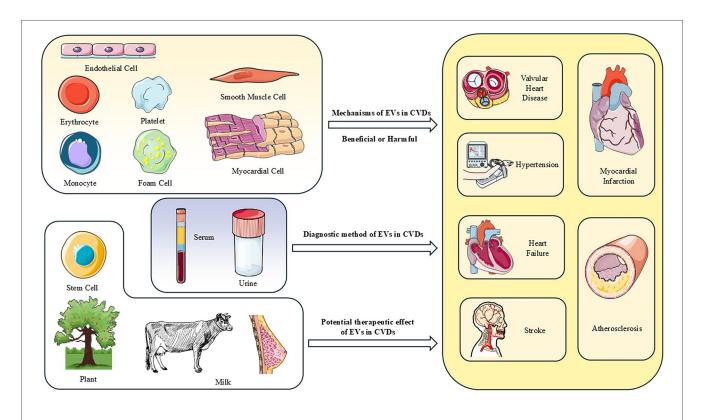
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Cardiovascular diseases (CVDs), the leading global cause of mortality, underscore an urgent need for innovative diagnostic and therapeutic strategies. Extracellular vesicles (EVs)—lipid-bilayer nanoparticles transporting bioactive cargo (microRNAs, proteins, lipids)—are critical mediators of intercellular communication in CVD pathogenesis. They exhibit functional duality: propagating pathology (inflammation, fibrosis, thrombosis) while facilitating tissue repair. This review synthesizes EV biogenesis mechanisms, isolation methodologies, source-specific functions, and multifaceted roles in atherosclerosis, myocardial infarction, heart failure, and stroke. We further diagnostic biomarkers, evaluate EV-based engineered applications, clinical translation challenges, and future directions.

KEYWORDS

extracellular vesicles, cardiovascular diseases, atherosclerosis, myocardial infarction, stroke, heart failure, hypertension, valvular heart disease



GRAPHICAL ABSTRACT

The roles of EVs in CVDs and the potential of acellular therapy. EVs derived from tissues and circulating sources target recipient cells through intercellular communication, influencing their biological functions and thereby impacting the progression of cardiovascular diseases. EVs from plant cells and stem cells carry molecules that promote protection and repair of the heart, significantly alleviating damage caused by cardiovascular diseases, indicating great potential as acellular therapies. EVs from plasma, serum and urine provide a potential avenue for the diagnosis and prognosis of future CVDs.

1 Introduction

In recent years, with the widespread spread of unhealthy living habits and the aggravation of the global aging trend, the morbidity and mortality of cardiovascular diseases (CVDs) have increased significantly, now accounting for approximately one-third of all deaths globally (1). CVDs encompass a diverse range of conditions affecting the heart and vascular system, categorized into several primary groups. Atherosclerosis and hypertension are prevalent conditions often regarded as major risk factors for various cardiovascular diseases, leading to complications such as myocardial infarction, heart failure, valvular heart disease, and stroke. Ischemic heart disease, characterized by insufficient blood supply to the heart, mostly refers to coronary heart disease (CHD), which can manifest as stable or unstable angina and myocardial infarction. Valvular heart disease involves damage to or a defect in one of the four heart valves, affecting blood flow through the heart. Finally, stroke, caused by interrupted blood supply or vessel rupture in the brain, encompasses ischemic and hemorrhagic types. This article primarily focuses on current research related to extracellular vesicles and their association with CVDs, as well as the closely related conditions in their pathogenesis. Despite significant advances

in cardiovascular disease research, there remains an urgent need for the development of more effective preventive strategies, diagnostic tools, and therapeutic interventions. Extracellular vesicles (EVs) exhibit a dualistic nature in cardiovascular diseases (CVDs), functioning as both essential mediators of physiological homeostasis and potential instigators of pathological progression. Under physiological conditions, EVs serve as critical vectors for intercellular communication within the cardiovascular system. By selectively transferring bioactive molecules-including proteins, miRNAs, and lipids-between cells, EVs orchestrate vital processes such as angiogenesis, metabolic coordination, and tissue regeneration. This molecular crosstalk maintains vascular integrity, supports cardiac repair mechanisms, and ensures functional synergy among diverse cell types. Conversely, pathological contexts trigger EVs to adopt detrimental roles. Stress-induced alterations in EV biogenesis lead to the packaging of pro-inflammatory, pro-fibrotic, and proarrhythmic cargoes. These aberrant EVs propagate cellular damage by facilitating endothelial dysfunction, amplifying inflammatory cascades, and accelerating fibrotic remodeling. Moreover, they disrupt electrophysiological stability and promote maladaptive signaling across cardiac tissues. Systemically, EVs derived from injured cells disseminate pathological molecules to distant organs, exacerbating multi-organ crosstalk in conditions 10.3389/fcvm.2025.1666589

like heart failure. Furthermore, EVs demonstrate significant diagnostic utility (2). Circulating EVs encapsulate cell-specific molecular signatures shielded from degradation, offering a dynamic "liquid biopsy" platform (3). Quantitative and qualitative profiling of EV cargoes enables early detection of subclinical cardiovascular injury, stratifies disease severity, and predicts risks of major adverse events. Integration of multi-omics EV data with advanced analytics heralds a new era in precision cardiology, transforming EVs from biological messengers into powerful clinical tools. Given the breadth of cardiovascular phenotypes and the uneven volume of EV-specific evidence, this review focuses on atherosclerosis, myocardial infarction, heart failure, hypertension, and valvular heart disease; findings related to cardiomyopathies are discussed within the Heart Failure section where the clinical and pathophysiological features converge.

2 Overview of extracellular vesicles

2.1 Classification and biogenesis process of EVs

Extracellular vesicles mainly include microvesicles, exosomes, and apoptotic bodies. They exhibit significant differences in size, composition, and surface markers (4) (Figure 1). Furthermore, the EVs research landscape is expanding to include other nanoscale vesicular structures, such as nanovesicles, which are typically defined by their small size (<50–100 nm) and are of emergent research interest due to their potential as efficient delivery vehicles (5). The article mainly talks about microvesicles and exosomes.

Microvesicle (MV) biogenesis originates from the outward budding and fission of the plasma membrane. This process typically bypasses endosome formation and necessitates significant

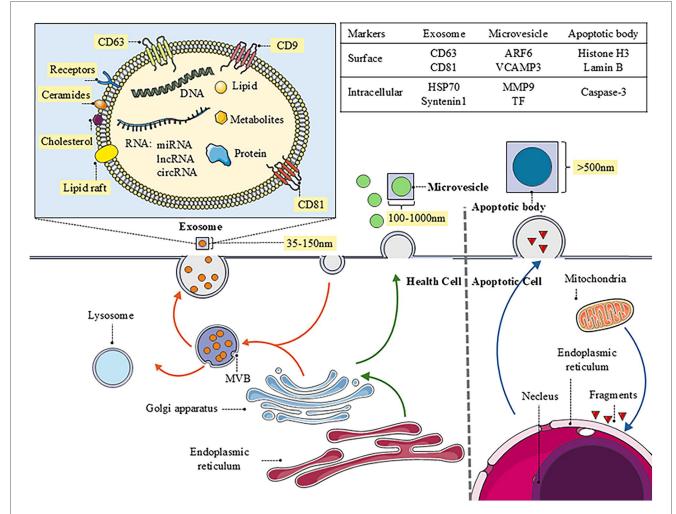


FIGURE 1

The biogenesis of EVs and the distinct characteristics of different types. The image provides a detailed depiction of the biogenesis and intricate molecular structure of exosomes, while briefly outlining the generation of microvesicles and apoptotic bodies. Additionally, it specifies the functions and markers associated with each type of vesicle, offering important insights into intercellular communication and their roles in biological processes.

molecular reorganization within the plasma membrane, involving alterations in lipid composition, protein distribution, and calcium ion (Ca²⁺) concentration. Key facilitators, such as aminophospholipid translocases (flippases/floppases), scramblases, and calpain proteases, disrupt membrane phospholipid asymmetry. This disruption promotes physical membrane curvature, remodels the actin cytoskeleton, and ultimately enables membrane budding and MV release (6). The involvement of cytoskeletal components and their regulatory machinery is essential for MV generation (7). Accumulating evidence indicates a strong association between MV formation and the modulation of cytoskeletal dynamics by small GTPases, notably the Rho (RAS homology) family and ADP-ribosylation factor (ARF). In malignant cells, ROCK and ARF6 GTPases critically mediate vesicle budding by regulating cytoskeletal dynamics.

The generation of exosomes mainly involves five steps: endocytosis and inward budding of the plasma membrane, formation of early sorting endosomes (ESE), maturation of late sorting endosomes (LSE), formation of multivesicular bodies (MVB), and finally, the release of exosomes. First, exosome formation begins with the endocytosis of the cell membrane, allowing external substances to enter the cell and form ESEs, which can exchange substances with the trans-Golgi network and endoplasmic reticulum. During this process, the inward-budded membrane fuses with the endoplasmic reticulum, Golgi apparatus, and even mitochondria. ESEs mature into LSEs, where secondary inward budding results in the formation of intraluminal vesicles (ILV), which contain cellular components such as proteins, nucleic acids, and lipids (8). Thus, the formation of ILVs is one of the core steps in exosome biogenesis and is dependent on the endosomal sorting complex required for transport (ESCRT) complexes (ESCRT-0, -I, -II, and -III) and their core proteins, such as ESCRT-III, ALIX, and tumor susceptibility gene 101 (TSG101) (9). These proteins promote ILV generation by regulating inward budding of membranes and separation of vesicles (10). Subsequently, mature MVBs can either interact with intracellular autophagosomes or lysosomes for degradation or approach the cell membrane through the cellular cytoskeleton and microtubule system. When fused to lysosomes, ILVs are degraded, conversely, when they fuse with cell membranes, endosomes release exosomes into the extracellular space (11). In addition to the classical ESCRT-mediated pathway, exosome formation can also occur through ESCRT-independent pathways, such as those involving lipids and Rab GTPases. lipid raftrich regions of the cell membrane may directly bend inward during endocytosis to form ILVs, a process that does not rely on ESCRT complexes but still requires specific lipids and membrane proteins, such as tetraspanins (e.g., CD63, CD81, etc.), which play essential roles in the formation and function of exosomes (12, 13). Similarly, Rab GTPases are involved in ESCRT-mediated exosome biogenesis and also regulate exosome formation independently of the ESCRT pathway. Rab GTPases, such as Rab27a, Rab35, and Rab11, promote the release of exosomes by regulating the trafficking of MVBs and their fusion with cell membranes.

Apoptotic bodies represent another category of EVs, generated through plasma membrane budding during programmed cell death (14). These vesicles are primarily cleared by phagocytes via efferocytosis. Regulation of apoptotic body formation involves specific molecular mediators governing apoptotic cell disassembly, including ROCK1 kinase, Pannexin 1 channels, and Plexin B2 receptors. Compared to exosomes and microvesicles, studies on apoptotic bodies as an EV subtype are limited (15).

From a functional and application perspective, EVs can be categorized into natural and artificial/synthetic classes. Natural EVs (like the aforementioned MVs and EXOs) carry bioactive molecules from their parent cells and act as crucial intercellular messengers, playing a dual role in the pathophysiology of cardiovascular diseases. In contrast, synthetic nanovesicles, including biomimetic vesicles based on cell membranes (e.g., platelet membrane, stem cell membrane vesicles) and synthetic liposomes, are primarily engineered to harness their potential as therapeutic delivery vehicles. By mimicking the membrane properties and targeting capabilities of natural EVs while allowing for precise engineering, these artificial nanovesicles can be efficiently loaded with drugs, nucleic acids (e.g., miRNA, siRNA), or functional nanoparticles for targeted delivery, overcoming the limitations of conventional therapies.

Evidence for their therapeutic potential is rapidly accumulating. Biomimetic nanovesicles designed for ischemic stroke therapy exemplify this promise. One innovative platform co-encapsulates thrombolytic tissue plasminogen activator (tPA) with neuroprotective melanin nanoparticles within a platelet membrane shell (16). This design leverages natural thrombustargeting abilities for site-specific accumulation, followed by near-infrared light-triggered release of both cargoes. This approach not only accelerates clot lysis through thermal enhancement of tPA activity but also sequentially addresses ischemia-reperfusion injury through scavenging of reactive oxygen species, demonstrating sophisticated spatiotemporal control. Similarly, in atherosclerosis, engineered mesenchymal stem cell-derived nanovesicles have been functionalized with a novel targeting peptide (GSPREYTSYMPH, or PREY) selected through phage display screening (17). These PREY-conjugated nanovesicles exhibit precise homing to disturbed flow sitesearly atherosclerotic lesions characterized by endothelial activation. Upon accumulation, they deliver their innate antiinflammatory and pro-reparative cargo, effectively reducing endothelial permeability and monocyte recruitment in preclinical models. This strategy highlights how synthetic nanovesicles can be engineered to target specific cardiovascular pathophysiological niches for localized therapeutic intervention.

These emerging studies underscore the significant potential of nanovesicles, particularly through engineered designs, as next-generation precision platforms for cardiovascular drug delivery, offering solutions to longstanding challenges in targeting, controlled release, and combination therapy.

2.2 Isolation and separation of EVs

The isolation and enrichment of EVs constitute a critical prerequisite for their clinical application as diagnostic biomarkers and therapeutic agents (Table 1). Current

TABLE 1 Isolation methods of EVs (22, 26-34).

Methods	Abbreviation	Advantages	Disadvantages	Reference title
Differential/ Ultracentrifugation	DC/UC	Cost-effective for large volumes	Long processing time	Improved recovery of urinary small extracellular vesicles by differential ultracentrifugation
		Simple protocols	Risk of EVs damage/ aggregation	
		Accessible equipment	Protein co-isolation	
Density Gradient	DGC	High purity	Very time-consuming	Isolation of human salivary extracellular vesicles by iodixanol
Centrifugation		Preserves integrity	Low throughput; specialized equipment	density gradient ultracentrifugation and their characterizations
		Reduces aggregation	Risk of sample loss	
Size-Exclusion	SEC	Rapid processing	Limited sample volume	A review of exosomal isolation methods: Is size exclusion
Chromatography		High EVs/protein ratio	EVs dilution	chromatography the best option?
		Preserves function	Co-elution of contaminants	
Ultrafiltration/Tangential	TFF	Rapid processing	Risk of membrane clogging	Extracellular vesicle isolation by a tangential-flow filtration-
Flow Filtration		Reduced clogging	Shear stress lead to EVs deformation	based large-scale purification method
		Preserves integrity	Yield loss	
Polymer-Based Precipitation	РВР	Simple protocols	Co-precipitation of proteins or polymers	Enrichment of astrocyte-derived extracellular vesicles from human plasma
		High concentration yield	Contaminant interference in downstream analysis	
Immunoaffinity Capture	-	High specificity and purity	Low yield	EV separation: release of intact extracellular vesicles immunocaptured on magnetic particles
		subpopulation isolation	Expensive cost of antibodies	
		Works with small volumes	Excludes antigen-negative Evs	
Microfluidic-Based	-	Minimal sample use	Specialized fabrication	Recent advances in microfluidic-based extracellular vesicle
Isolation		Automatable/high- throughput	Limited scalability	analysis
Field-Flow Fractionation	FFF	High integrity	Specialized instrument	Separation and characterization of extracellular vesicles from
		Isolates diverse EV subsets	Require pre-concentration	human plasma by asymmetrical flow field-flow fractionation
Hydrostatic Dialysis	HDI	Effective abundant	Requires validation	MiR-145 detection in urinary extracellular vesicles increase
Isolation		protein removal	Limited adoption	diagnostic efficiency of prostate cancer based on hydrostatic filtration dialysis method
Aqueous Two-Phase	ATPS	Pure and scalable	Higher cost and longer time	Next generation aqueous two-phase system for gentle, effective,
System		Gentle treatment	Limited clinical utility	and timely extracellular vesicle isolation and transcriptomic analysis

mainstream separation techniques include ultracentrifugation (UC), density gradient ultracentrifugation, size-exclusion chromatography (SEC), ultrafiltration (UF), polymer-based precipitation, field-flow fractionation, and immunoaffinity capture (IA). Studies have demonstrated that polymer-based precipitation and size-exclusion chromatography yield the highest total EV production (18). Notably, asymmetric flow field-flow fractionation (AsFIFFF/AF4) has emerged as a promising technique for isolating EVs subpopulations (19). Microfluidic platforms and combinatorial multi-step approaches also exhibit unique potential in this field. Given the inherent biological complexity, source heterogeneity, and functional diversity of EVs, integrated separation strategies are generally considered optimal (20). Current technical challenges primarily involve exogenous contamination, laborious operational procedures, and standardization deficiencies, necessitating the development of novel separation-enrichment systems with enhanced purity and efficiency. For the distinct sources discussed herein, common EV isolation methods are as follows, Serum/plasma commonly employs UC or SEC (21). Urine frequently uses UF combined with SEC or polymer-based precipitation (22). Tissue-derived EVs require initial enzymatic digestion/homogenization, followed predominantly by density gradient ultracentrifugation. Stem cell supernatant often relies on UC, polymer-based precipitation, or SEC. Milk typically favors SEC or optimized density gradient ultracentrifugation (23). Plant-derived EVs mainly depend on differential centrifugation combined with density gradient centrifugation (24). Combinatorial approaches (e.g., UC+SEC) are widely adopted to enhance purity and yield for specific sources, and emerging technologies like microfluidics show promise for advancing EV isolation (25). Optimizing source-specific isolation strategies remains a key research priority.

2.3 Different origins and functions of EVs

2.3.1 EVs derived from plasma and serum

Mammalian blood is a primary source of extracellular vesicles (EVs), which can be isolated from either plasma or serum. Plasma

is obtained from anticoagulated blood and retains coagulation factors, providing a representation of circulating EVs that is closer to the in vivo physiological state. In contrast, serum is derived from clotted blood, a process that potently activates platelets, leading to a significant ex vivo release of plateletderived EVs and concomitant consumption of coagulation factors (35). Due to minimized ex vivo cellular activation, plasma-derived EVs are generally considered more reliable for reflecting the native circulating EV profile, exhibiting less preanalytical variability and thus offering greater result reproducibility. Consequently, plasma is the preferred and most common source for EV research in cardiovascular diseases. However, serum serves a specific purpose in studies focusing on thrombosis. The very process of clot formation enriches serum with EVs released from platelets under low-flow and hypercoagulable conditions ex vivo, making it a valuable medium for investigating the role of platelet-derived EVs in thrombotic pathologies (36). Both plasma-EVs and serum-EVs originate from various cellular sources and serve as critical mediators of intercellular communication and clinically validated diagnostic biomarkers. They encapsulate cell-statespecific cargoes (e.g., miRNAs, proteins) that reflect real-time pathophysiological states. For instance, sEV-derived miR-1 and miR-133a demonstrate significant elevation within 3 h postinfarction, serving as early-warning signatures for acute myocardial infarction (AMI) prior to troponin detectability (37). Their intrinsic biocompatibility and circulatory stability further underscore their potential for targeted therapeutic delivery.

2.3.2 EVs derived from milk

Milk contains high amounts of EVs originating from different cell populations in the mammary glands (e.g., adipocytes, epithelial cells, stem cells, and immune cells) and plays important roles in interorganismal and cross-species communication. Because milk is considered scalable. economical, and safe source of EVs, milk-derived EVs (miEVs) have recently been introduced as novel drug carriers (38). A significant advantage of miEVs is their suitability for oral administration, which is primarily attributed to their innate resistance to degradation in the harsh conditions of the gastrointestinal tract (39). Milk contains a high concentration of casein micelles, and these phosphoproteins exhibit chaperone-like properties that can encapsulate and protect EVs during transit through acidic environments. This protective effect facilitates the efficient uptake of EVs by intestinal epithelial cells, thereby enabling the systemic distribution of their bioactive cargos. Studies using cardiosphere-derived cells EVs (CDC-EVs) have illustrated that the association with casein enhances EV uptake and biodistribution, leading to improved tissue targeting and therapeutic outcomes in animal models (40).

2.3.3 Evs derived from plant

The Plant-Derived Extracellular Vesicles (PDEVs) are nanosized vesicle structures ranging from 50 to 1,000 nm in diameter, featuring a spherical lipid bilayer. Importantly, the

human immune system does not detect these PDEVs, allowing them to achieve longer circulation times and higher bioavailability (41). Furthermore, PDEVs exhibit the ability to penetrate the intestinal mucus barrier and withstand gastrointestinal enzymes and bile, making them a reliable and advantageous choice for drug delivery. They carry small RNAs and miRNAs that facilitate intercellular communication among species across different kingdoms. As natural vectors, PDEVs transport bioactive compounds derived from their source plants. In addition to their functional roles, PDEVs demonstrate significant biological activities, such as antioxidant and anti-inflammatory effects (42). Notably, their outer lipid bilayer membrane acts as a protective barrier for bioactive molecules against external conditions. This structural feature not only allows PDEVs to sustain circulation within the body for extended periods but also promotes the continuous accumulation of drugs. Moreover, PDEVs are known for their strong safety profile, exhibiting low toxicity and good biocompatibility while minimizing side effects, which further enhances their potential as effective drug delivery systems.

3 Mechanisms of EVs in CVDs

3.1 Mechanisms of EVs in atherosclerosis

3.1.1 Endothelial dysfunction and vascular inflammation

Atherosclerosis is fundamentally a chronic inflammatory and fibroproliferative response to retained and modified lipoproteins. Classical risk factors reduce endothelial nitric oxide bioavailability and increase permeability and adhesion molecules, promoting monocyte recruitment transendothelial migration (43). Within the intima, monocytes differentiate into macrophages and foam cells; activated platelets amplify leukocyte recruitment and coagulation, whereas smooth muscle cells migrate and produce matrix, driving plaque growth, calcification, and instability (44). Throughout these steps, EVs shuttle miRNAs, proteins, and lipids that fineendothelial-monocyte/macrophage-platelet-smooth muscle cross-talk: some EVs propagate oxidative stress, inflammation, and thrombosis, while others favor resolution and repair (45, 46). This framework underpins the mechanistic and translational studies summarized below (Tables 2, 3).

3.1.1.1 Endothelial cell-derived EVs in AS

The inducing factors for EVs that cause endothelial cell injury partially to overlap with traditional cardiovascular disease (CVD) risk factors, including but not limited to: diabetes, hyperlipidemia, hypertension, and smoking. The exosomal cargo NEDD4l, released by vascular endothelial cells under high glucose and ox-LDL induction, enhances the ubiquitination of $I\kappa B-\alpha$ and PPAR γ , promotes macrophage M1 polarization and ox-LDL uptake, thereby exacerbating endothelial injury (47). Under high glucose and AngII stimulation, endothelial cell-derived EVs

TABLE 2 Damage effects of EVs in AS (47-49, 54, 56-64).

Source	Cargos	Signaling pathway/ target	Function	Reference title
ox-LDL-treated ECs	miR-505	SIRT3↓	$ \begin{array}{c} \text{Promotes NET} \\ \text{formation} \rightarrow \text{Aggravates endothelial} \\ \text{inflammation} \end{array} $	Exosome-encapsulated miR-505 from ox-LDL-treated vascular endothelial cells aggravates atherosclerosis by inducing NET formation
ox-LDL-treated ECs	HIF1A-AS2	miR- 455-5p \downarrow \rightarrow ESRRG \uparrow \rightarrow NLRP3 \uparrow	Induces endothelial pyroptosis and vascular inflammation	Endothelial cell-released extracellular vesicles trigger pyroptosis and vascular inflammation to induce atherosclerosis
ox-LDL-treated ECs	LINC01005	miR-128-3p↓ → KLF4↑	Drives VSMC phenotypic switching → Endothelial injury	Exosomal LINC01005 derived from oxidized low- density lipoprotein-treated endothelial cells regulates vascular smooth muscle cell phenotypic switch
High glucose-treated ECs	PDGF-BB	miR-296-5p↑ → Bcl-2↑Bak/Bax↓	Suppresses VSMC apoptosis → Abnormal angiogenesis	PDGF-BB carried by endothelial cell-derived extracellular vesicles reduces vascular smooth muscle cell apoptosis in diabetes
Obese visceral adipose tissue	miR-27b-3p	$PPAR\alpha \downarrow \rightarrow NF \text{-} \kappa B \uparrow$	Induces endothelial inflammation	Exosomal miR-27b-3p secreted by visceral adipocytes contributes to endothelial inflammation and atherogenesis
High Glucose + ox- LDL treated Ecs	NEDD4l	IκB-α/PPARγ↓	Promotes macrophage M1φ polarization and ox-LDL uptake	Exosomal NEDD4l derived from HG + oxLDL-induced vascular endothelial cells accelerates macrophage M1 polarization and oxLDL uptake by ubiquitinating $I\kappa B\alpha$ and PPAR γ .
High Glucose + AngII treated Ecs	-	eNOS↓and ERK↑	Reduces NO bioavailability → increases permeability	Glucose and angiotensin II-derived endothelial extracellular vesicles regulate endothelial dysfunction via ERK1/2 activation.
Radiation-treated Ecs	miR-126-5p/ miR-212-3p	Monocytes†	Triggers vascular inflammation	The miR-126-5p and miR-212-3p in the extracellular vesicles activate monocytes in the early stage of radiation-induced vascular inflammation implicated in atherosclerosis
Platelets	miR-92a-3p	PTEN/PIP3/Akt → Col8a1↑	Induces vascular stiffness and reduced elasticity	Platelet-Derived Extracellular Vesicles Increase Col8a1 Secretion and Vascular Stiffness in Intimal Injury
M1 macrophages	miR-185-3p	Smad7 $\downarrow \rightarrow TGF-\beta\uparrow$	Inhibits EC proliferation → Promotes EC apoptosis	M1 macrophages-derived extracellular vesicles elevate microRNA-185-3p to aggravate atherosclerosis
ox-LDL-treated macrophages	miR-19b-3p	JAZF1↓	Enhances VSMC migration/ proliferation	microRNA-19b-3p-containing extracellular vesicles derived from macrophages promote the development of atherosclerosis by targeting JAZF1
Mature dendritic cells	TNF-α	NF-κB↑	Induces endothelial injury	Exosomes derived from mature dendritic cells increase endothelial inflammation and atherosclerosis via membrane TNF-α mediated NF-κB pathway
Neutrophils	miR-155	BCL6↓	Exacerbates endothelial inflammation	Neutrophil microvesicles drive atherosclerosis by delivering miR-155 to atheroprone endothelium
Nicotine exposure (via MEVs)	miR-155	-	Worsens endothelial damage	Nicotine exacerbates endothelial dysfunction and drives atherosclerosis via extracellular vesicle-miRNA
NAFLD hepatocytes	miR-1	KLF4/NF-κB↑	Triggers endothelial inflammation	Hepatocyte-derived extracellular vesicles promote endothelial inflammation and atherogenesis via microRNA-1
Sleep deprivation	miR-182-5p↓	MYD88↓ → NF-κB/NLRP3↓	Promotes endothelial inflammation	Sleep deprivation promotes endothelial inflammation and atherogenesis by reducing exosomal miR-182-5p
TNF-α/shear stress- treated ECs	miR-92a	KLF4↓	Accelerates LDL uptake → Foam cell formation	Extracellular MicroRNA-92a mediates endothelial cell- macrophage communication
H. pylori Bacteria	Bacterial antigens	IL-6/IL-8↑	Mediates inflammation and endothelial injury	Extracellular vesicles and endothelial dysfunction in infectious diseases.

reduce NO bioavailability by inhibiting endothelial nitric oxide synthase (eNOS) activity and activating the ERK signaling pathway, thereby promoting endothelial dysfunction and increasing permeability (48). Endothelial cell-derived EVs containing miR-126-5p and miR-212-3p are released after radiation exposure, triggering vascular inflammation via monocyte activation (49) This early dysfunction facilitates leukocyte adhesion and trans-endothelial migration, initiating the inflammatory cascade in the arterial intima.

3.1.1.2 Monocyte cell-derived EVs in AS

During the development of atherosclerosis, following endothelial dysfunction, the next significant contributor to AS is foam cell formation. Under the influence of various adhesion and chemotactic factors, monocytes adhere to the damaged vascular endothelium, migrate across the vascular wall, and transform into macrophages. Macrophages can uptake erythrocyte-derived extracellular vesicles via endocytosis (50). Under inflammatory stimulation, their cargo heme can attenuate the ability of ox-LDL-

TABLE 3 Repair effect of EVs in AS (65-74).

Source	Cargos	Signaling pathway/ target	Function	Reference title
MSCs	miR-let7	HMGA2/IGF2BP1↓ → NF- κB/PTEN↓	Promotes M2 polarization → Anti- inflammation	Exosomes derived from mesenchymal stem cells attenuate the progression of atherosclerosis in ApoE(-/-) mice via miR-let7 mediated infiltration and polarization of M2 macrophage
MSCs	TGF-β1	miR-132↑ → Mycbp2/TSC2↑	Induces M2 polarization → Inhibits calcification	Mesenchymal stem cell-secreted extracellular vesicles carrying TGF- β 1 up-regulate miR-132 and promote mouse M2 macrophage polarization
MSCs	miR-145	JAM-A↓	Inhibits EC migration \rightarrow Reduces inflammation	Mesenchymal stem-cell-derived exosomal miR-145 inhibits atherosclerosis by targeting JAM-A
MSCs	FENDRR	miR-28↓ → TEAD1↑	Suppresses EC apoptosis/oxidative stress	Exosomes derived from mesenchymal stem cells ameliorate the progression of atherosclerosis in ApoE(-/) mice via FENDRR
Thrombin- activated platelets	miR-223	ICAM-1↓ → NF-κB/MAPK↓	Attenuates endothelial inflammation	Thrombin-activated platelet-derived exosomes regulate endothelial cell expression of ICAM-1 via microRNA-223
LSW-treated VSMCs	miR-145	PDCD4↓	Improves EC proliferation/ migration	Tripeptide Leu-Ser-Trp regulates the vascular endothelial cells phenotype switching by mediating the vascular smooth muscle cells-derived small extracellular vesicles packaging
Endothelial cells	miR-222	ICAM-1↓	Inhibits monocyte adhesion to ECs	Endothelial microparticles reduce ICAM-1 expression in a microRNA-222-dependent mechanism
M2 macrophages	miR-221-3p	Grb10↓	Reduces EC apoptosis/ inflammation	M2 macrophage-derived exosomes inhibit apoptosis of HUVEC cell through regulating miR-221-3p
Apoptotic ECs	miR-126	RGS16↓ → CXCR4/ CXCL12↑	Recruits progenitor cells → Endothelial repair	Delivery of microRNA-126 by apoptotic bodies induces CXCL12-dependent vascular protection
Perivascular adipose tissues	miR-382-5p	BMP4↑ → PPARγ-ABCA1/ ABCG1↑	Enhances cholesterol efflux → Reduces foam cells	Perivascular adipose-derived exosomes exert proatherogenic effects by regulating macrophage foam cell formation and polarization
Endothelial progenitor cells	miR-21-5p	SIPA1L2↓ → Autophagy↑	Promotes EC proliferation/ migration → Vascular repair	Endothelial colony-forming cell-derived exosomes regulate autophagic flux to promote vascular endothelial repair
HAL-loaded M2 macrophages	HAL (drug)	Heme synthesis↑ → CO/ bilirubin↑	Exerts anti-inflammatory effects	Molecularly engineered macrophage-derived exosomes with inflammation tropism and intrinsic heme biosynthesis for atherosclerosis treatment
Lemon-derived EVs	-	AhR/Nrf2↑	Antioxidant/anti-inflammatory effects	Lemon-derived nanovesicles achieve antioxidant and anti- inflammatory effects activating the AhR/Nrf2 signaling pathway
Paeonol-treated monocytes	miR-223	STAT3↓	Reduces EC adhesion/ inflammation	Paeonol attenuated inflammatory response of endothelial cells via stimulating monocytes-derived exosomal MicroRNA-223
Genetically modified ECs	anti-miR-33a- 5p	miR-33a-5p↓ → ABCA1↑	Enhances cholesterol efflux	Exosome-mediated transfer of anti-miR-33a-5p from transduced endothelial cells enhances macrophage and vascular smooth muscle cell cholesterol efflux
Foam Cells	Prosaposin	$GPR37\downarrow \rightarrow ERK/$ $AP1\uparrow \rightarrow Tim4\uparrow$	Enhances efferocytosis and accelerates inflammation resolution	Efferocytes release extracellular vesicles to resolve inflammation and tissue injury via prosaposin-GPR37 signaling
Avocado-derived EVs	ginkgetin and berberine	TNF-α/IL-6/IL-1β/CD36↓	Inhibits oxLDL-induced foam cell formation	Avocado-derived extracellular vesicles loaded with ginkgetin and berberine prevent inflammation and macrophage foam cell formation.
Quercetin-loaded EVs	Quercetin	SIRT1↑	Alleviates ferroptosis and aging- related endothelial damage	Therapeutic application of quercetin in aging-related diseases: SIRT1 as a potential mechanism.

treated macrophages to form foam cells. Foam cell-derived EVs express prosaposin, which binds to macrophage GRP37, increasing the expression of the efferocytosis receptor Tim4 through activation of the ERK-AP1 signaling axis (51). This leads to enhanced macrophage efferocytosis efficiency and accelerated inflammation resolution. Exosomes (BMDM-IL-4-exo) produced by bone marrow-derived macrophages (BMDMs) exposed to the M2-polarizing cytokine IL-4 contain abundant miR-99a/146b/378a (52). These exosomal miRNAs inhibit inflammation by targeting NF-κB and TNF-α signaling and further promote M2 polarization in recipient macrophages. Under high glucose induction, extracellular vesicles released by bone marrow cell-derived macrophages increase the number of circulating hematopoietic and myeloid cells. This leads to an increased macrophage population, exacerbating vascular inflammation and promoting atherosclerosis progression. Under nicotine stimulation, EVs derived from monocytes are secreted abundantly. These EVs encapsulate miR-155, which induces endothelial cell dysfunction and vascular inflammation by targeting BCL2, MCL1, TIMP3, BCL6 and activating the NF-κB pathway (53). During atherosclerosis progression, foam cell formation represents a pivotal pathological stage following endothelial dysfunction, wherein macrophages differentiated from migrated monocytes regulate inflammation and foam cell dynamics via EVs uptake; EVs derived from diverse cellular sources and their miRNA cargo exert bidirectional modulation, either resolving inflammation or exacerbating vascular damage through targeted signaling cascades, collectively driving disease advancement.

3.1.1.3 Platelet-derived EVs in AS

Platelet-derived EVs modulate vascular homeostasis through targeted miRNA delivery, exerting dual regulatory effects that

suppress endothelial inflammation under physiological conditions while promoting vascular stiffening and prothrombotic states in pathological contexts. Platelet-secreted EVs participate in delivering miR-92a-3p to vascular smooth muscle cells (VSMCs) (54). This then induces the production and secretion of Col8a1 via the PTEN/ PIP3/Akt pathway, leading to increased vascular wall stiffness and reduced elasticity (54). Thrombin-activated platelet-derived exosomes (P-EXO) release large amounts of miR-223, which subsequently inhibits ICAM-1 expression in endothelial cells (ECs). MiR-223 may prevent EC inflammation by modulating NF-kB and MAPK pathways (55). Simultaneously, they transport high levels of miR-25-3p, which is responsible for reducing ox-LDL-induced EC inflammation and lipid deposition, thereby inhibiting AS progression. Platelet-derived exosomes containing miRNAs such as miR-223 and miR-25-3p can, under certain conditions, modulate endothelial activation and reduce inflammation, however, in proinflammatory states, these EVs may also promote a procoagulant environment by delivering tissue factor (TF) and plasminogen activator inhibitor-1 (PAI-1).

3.1.1.4 Other sources EVs in AS

Steatotic hepatocyte-derived EVs promote endothelial inflammation by mediating miR-1-induced KLF4 pathway suppression and NF-κB pathway activation (75). Human aortic stenotic valve-derived extracellular vesicles can lead to endothelial dysfunction, pro-adhesive and procoagulant responses via the AT1R/NADPH oxidase/sodium-glucose cotransporter 2 (SGLT2) pro-oxidative pathway (76). Extracellular vesicles derived from human induced pluripotent stem cell-derived endothelial cells (hiPSC-ECs) reduce endothelial cell apoptosis and restore autophagy, suggesting that EV-based therapies offer a promising avenue for targeting endothelial dysfunction in similar vascular pathologies (77). During Helicobacter pylori infection, bacteriumderived EVs deliver bacterial antigens into vascular endothelial cells, mediating inflammation and endothelial injury (64). Notably, milk-derived EVs enriched with miR-30b-5p have been implicated in cardiovascular and metabolic diseases by modulating inflammation and oxidative stress. This miRNA suppresses TLR4 activity in endothelial cells, downregulates pro-inflammatory molecules, and thereby attenuates atherosclerotic progression (78). In atherosclerosis, EVs derived from heterogeneous cellular sources can pathologically exacerbate endothelial damage and procoagulant responses, while therapeutically attenuating apoptosis and restoring autophagic homeostasis.

3.1.1.5 Potential therapeutic effect of PDEVs in AS

Many studies indicate that plant-derived extracellular vesicles (PDEV) can deliver antioxidants as a therapeutic application for oxidative stress-related diseases. MiRNAs in green leafy vegetable-derived EVs (such as miR-156a) have been shown to exert protective effects against atherosclerosis by modulating gene expression in human cells. Furthermore, plant compounds with antioxidant activity known to be carried in these vesicles (e.g., polyphenols and flavonoids) can scavenge reactive oxygen species and upregulate protective pathways like Nrf2 (42). These effects may reduce endothelial oxidative stress and the

production of inflammatory mediators, both key to the pathogenesis of vascular inflammation. In a study investigating avocado-derived extracellular vesicles, researchers demonstrated a significant reduction in macrophage inflammatory cytokine expression (including TNF- α , IL-6, IL-1 β , and Cd36) and inhibition of oxLDL-induced foam cell formation (79). EVs serve as vehicles for quercetin loading and delivery, mediating SIRT1 activation to alleviate ferroptosis and prevent aging-related diseases (80). This study highlights the potential of EVs as signaling carriers for treating aging-related diseases.

3.1.2 Plaque formation, calcification & destabilization

Monocyte recruitment and foam cell accumulation secondary to endothelial dysfunction are hallmarks of atherosclerotic plaque growth. Following initial plaque formation and VSMC proliferation and migration, calcium deposition can begin, particularly within the fibrous cap covering the plaque. The calcification process plays a crucial role in plaque vulnerability.

Under endotoxin stimulation, monocyte-derived microvesicles can transmit cell death signals via encapsulated caspase-1, inducing programmed apoptosis in VSMCs, thereby promoting the calcification process and plaque formation (81). EV-derived circ_0001785 has been identified as a novel biomarker for atherosclerosis and has been demonstrated to reduce endothelial cell injury and delay plaque formation via the miR-513a-5p/ TGFBR3 ceRNA network mechanism (82). Engineering extracellular vesicles offers a potential EV-based therapeutic strategy for atherosclerosis generation. Small extracellular vesicles (psEVs) derived from carotid artery plaques provide insights into tissue- and disease-specific pathology. PsEVs have been demonstrated to induce inflammatory endothelial dysfunction in vitro and exacerbate atherogenesis in ApoEdeficient mice (83). During endothelial cell apoptosis, CXCL12 production mediated by miR-126, and released via apoptotic bodies, promotes progenitor cell recruitment through paracrine action and inhibits macrophage adhesion to the vascular wall, thereby limiting atherosclerosis progression (72). Mouse experiments have demonstrated that administering CXCL12-containing apoptotic bodies or miR-126 confers plaque-stabilizing characteristics in different atherosclerotic mouse models.

As plaques progress, they undergo complex remodeling processes, including calcification and fibrous cap destabilization. EVs released by VSMCs and macrophages are implicated in promoting calcification and altering plaque stability. Calcifying EVs are a subset of EVs enriched in calcium-binding proteins (such as annexins) and can serve as nucleation factors for hydroxyapatite crystals, thereby promoting microcalcification within plaques. A biomimetic nanocarrier modeled after natural grapefruit-derived extracellular vesicles was designed and manufactured (84). It is loaded with sodium thiosulfate (STS, an approved drug for treating vascular calcification—VC) and further modified with an elastin-specific targeting peptide (ESTP) for VC-targeted delivery of STS. In *in vitro* experiments,

ESTP nanomedicine demonstrated superior cellular uptake in calcifying vascular smooth muscle cells. Through the absorption of delivered STS, it inhibited VSMC calcification. Mechanistically, ESTP nanomedicine significantly prevented VC by driving M2 macrophage polarization, reducing inflammation, and inhibiting the bone-vascular axis.

Furthermore, EV-mediated delivery of matrix metalloproteinases (MMPs) and other proteolytic enzymes to the extracellular matrix can degrade collagen and other structural proteins within the fibrous cap. This degradation not only weakens the fibrous cap strength but also increases the risk of plaque rupture and subsequent thrombus formation. Plaque rupture is often the triggering event for acute thrombotic cardiovascular events. EVs promote the thrombotic cascade by providing a procoagulant surface rich in phosphatidylserine (PS) and tissue factor (TF). Specifically, platelet-derived EVs can enhance the coagulation process by transferring TF to the surfaces of endothelial cells and monocytes, thereby augmenting thrombin generation and local clot formation (85). EVs from monocyte/platelet aggregates can stimulate the release of proinflammatory cytokines from the exacerbating the local further prothrombotic microenvironment, secondary to other types of cardiovascular diseases (86). EVs exert dual regulatory effects both beneficial and detrimental on the inflammation-fibrosis-calcification axis during the development of atherosclerotic disease. On one hand, ox-LDL infiltrates endothelial cells, and various EVs released by endothelial cells recruit the mononuclear lymphatic system, promoting endothelial inflammation. This leads to subsequent endothelial fibrosis and eventual calcification detachment in the progression of atherosclerosis. On the other hand, EVs derived from endothelium and plaques suppress endothelial inflammation and plaque detachment, thus playing a beneficial role in disease progression.

3.2 Mechanisms of EVs in hypertension

3.2.1 Mechanisms of EVs in hypertension damage

The pathogenic role of EVs in hypertension shares several key mechanisms with atherosclerosis, particularly involving chronic inflammation, oxidative stress, and dysregulated vascular tone. In hypertension, EVs derived from activated vascular endothelial cells, platelets, and immune cells propagate vascular dysfunction by transferring proinflammatory cytokines, signaling receptors, and regulatory RNAs to recipient cells (87). For instance, endothelial cell-derived EVs impair eNOS activity, thereby reducing nitric oxide (NO) bioavailability, a key factor in maintaining vascular tone. Moreover, EVs enriched in molecules such as TNF-α and IL-1β can activate NF-κB signaling in vascular smooth muscle and endothelial cells, promoting a proinflammatory environment that contributes to vascular remodeling and stiffness. EVs also contribute to the dysregulation of the renin-angiotensin system (RAS) by transporting components like the angiotensin II type 1 receptor (AT1R), which can be transferred to recipient cells to potentiate the local RAS response and facilitate processes like hypertrophy and remodeling (88). Furthermore, beyond the vasculature, brainderived EVs have been shown to induce neuroinflammation and oxidative stress within key cardiovascular regulatory regions of the brain, such as the paraventricular nucleus (89). These EVs carry proinflammatory cytokines and enzymes that promote mitochondrial dysfunction and reactive oxygen species production, perpetuating a state of oxidative stress that increases sympathetic nervous system activity, a recognized driver of hypertension (90). The molecular cargo of EVs differs significantly between hypertensive and normotensive states, underscoring their active role in both perpetuating disease and potentially offering pathways for intervention.

3.2.2 Diagnostic potential of EVs in hypertension

At present, the diagnosis and clinical management of essential hypertension continue to rely predominantly on conventional blood pressure (BP) measurements (91). While these measurements remain fundamental to hypertension assessment, they cannot differentiate between hypertensive subtypes, identify underlying molecular mechanisms, or optimally monitor treatment efficacy (92). This has created a compelling need for novel biomarkers that facilitate early detection, enable patient stratification, and allow monitoring of cardiovascular complications. EVs have recently emerged as promising candidate biomarkers in cardiovascular research due to their stability in circulation, ease of isolation from accessible bodily fluids, and, most importantly, their molecular cargo that dynamically reflects cardiovascular stress responses and pathological remodeling (93).

Evidence from multiple studies indicates that plasma-derived and urinary-derived EVs carry distinctive molecular signatures associated with hypertensive cardiovascular damage (94). Clinical studies have established that circulating platelet-derived EVs show significant correlations with nocturnal blood pressure patterns, non-dipping status, and increased pulse wave velocity, a validated marker of arterial stiffness, establishing these EVs as integrated biomarkers of vascular health status (95). Furthermore, research has indicated that endothelial-derived EVs carrying angiotensin II type 1 receptors (AT1R) exhibit significantly elevated levels in hypertensive patients and potentially modulate vascular responses to neurohormonal activation (96).

The miRNA content of EVs provides particularly valuable insights into hypertension-related cardiovascular pathology. Upregulated miRNAs such as miR-320d and miR-423-5p have been mechanistically implicated in promoting vascular smooth muscle phenotype switching and vascular remodeling (97). Furthermore, studies have identified that EVs from injured endothelial cells can remodel the vessel wall in hypertension through these miRNA-mediated mechanisms (98). Proteomic analyses have also revealed significant alterations in EV protein cargo; for instance, hypoxia-induced EVs carry glucose-regulated protein 78 kDa (GRP78), which contributes to vascular smooth muscle cell calcification—a process relevant to hypertensionmediated cardiovascular damage (99). Another study demonstrated that circulating EV levels were significantly different between white coat hypertension and sustained hypertension phenotypes, suggesting their utility in hypertension subtyping (95).

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Emerging technologies are enhancing the diagnostic potential of EV-based biomarkers. Analysis of plasma small extracellular vesicles (sEVs) using surface-enhanced Raman scattering (SERS) combined with machine learning methods has demonstrated high sensitivity and specificity for cardiovascular disease detection (100). This interdisciplinary approach offers a promising strategy for non-invasive, precise early detection of hypertension-related cardiovascular impairment. Additionally, research into Mendelian hypertension forms has discovered EV-mediated alterations in vascular function, particularly through mechanisms involving vascular smooth muscle calcification and endothelial dysfunction (101).

The multifaceted information contained within EVs, reflecting various aspects of cardiovascular pathophysiology, offers a promising platform for improved risk stratification and dynamic monitoring of hypertensive patients beyond conventional BP assessment. Their ability to provide insights into vascular stiffness, myocardial stress, and vascular remodeling makes them particularly valuable for managing hypertension-related cardiovascular complications.

3.3 Mechanisms of EVs in myocardial infarction and ischemia/reperfusion injury

Over the past five years, while research on extracellular vesicles (EVs) in the AS (atherosclerosis) field has primarily focused on pathogenic mechanisms and cell-free therapies, the main research focus on EVs in myocardial ischemia, myocardial infarction (MI), and ischemia-reperfusion injury (I/R) has centered on stem cell therapy. The etiology of these three diseases is attributed to vascular occlusion, caused by various factors including atherosclerosis, thrombosis, and other pathological processes, ultimately leading to cardiac ischemia. Myocardial ischemia refers to the deficiency of oxygen and nutrients in the myocardium due to insufficient blood supply. If this state persists, it may progress to myocardial infarction, characterized by irreversible necrosis of cardiomyocytes. Furthermore, during reperfusion following myocardial infarction, the restoration of blood flow can induce ischemiareperfusion injury, which further exacerbates cardiomyocyte damage and impacts patient prognosis. Therefore, understanding these pathophysiological processes is crucial for developing effective clinical interventions.

3.3.1 Mechanisms of EVs in MI damage

Extracellular vesicles from CDC-EVs enhance regulatory T cell (Treg) proliferation and interleukin-10 (IL-10) production. BCYRN1, a long non-coding RNA (lncRNA) highly abundant in CDC-EVs, plays a significant role (102). In mouse models of myocardial infarction and reperfusion injury, administration of CDC-EVs, particularly those overexpressing BCYRN1, demonstrated cardioprotective effects by reducing infarct size and troponin I levels. M2 macrophage-derived extracellular vesicles containing miR-378a-3p in cardiomyocyte pyroptosis after MI disrupt NLRP3 and inhibit activation of the NLRP3/

Caspase-1/GSDMD pathway by suppressing ELAVL1 (HuR) expression and HuR translocation to the cytoplasm, thereby alleviating cardiomyocyte pyroptosis (103). Recent studies indicate that EVs derived from coronary injury sites in MI patients, particularly leukocyte-derived EVs carrying malondialdehyde (MDA+) oxidation epitopes, neutrophils via the TLR4/PAD4 signaling pathway, inducing NETosis, thereby exacerbating local thrombosis and myocardial injury (104). Naturally occurring human MDA-specific IgM antibodies (MDA-IgM) significantly inhibit NETosis. A strong negative correlation was verified between the CD45 + MDA + EVs/IgM ratio and worsening cardiac function. Findings suggest that M2 macrophage-derived small extracellular vesicles (SEVs) containing circUbe3a promote proliferation, migration, and phenotypic transformation of cardiac fibroblasts (CFs) by directly targeting the miR-138-5p/RhoC axis, which may also exacerbate myocardial fibrosis after acute myocardial infarction (105). The EVs/macrophage axis plays a role in exacerbating I/R injury, similar to its function in damaging vascular endothelium in AS. EVs can influence the progression of myocardial ischemia-reperfusion injury by affecting macrophage polarization. miR-155-5p in heart-derived EVs promotes macrophage M1 polarization by activating the JAK2/ STAT1 pathway, leading to local cardiac inflammation and even triggering systemic inflammation in distant organs (106). During MI progression, EVs orchestrate a paradoxical landscape of protective regeneration vs. pathological damage through spatiotemporally constrained immunomodulatory and cell death pathway regulation.

3.3.2 Diagnostic effect of EVs in MI

Currently, the standard biomarkers for myocardial infarction (MI) are high-sensitivity cardiac troponins, including cTn-I and cTn-T (107). Although highly effective in the clinical setting for diagnosing MI, high-sensitivity troponins focus solely on AMI diagnosis and not on prediction and early warning. Therefore, finding new biomarkers is necessary, as early diagnosis of MI is associated with improved outcomes for future MI development. EVs have garnered considerable interest as novel blood biomarkers (Table 4). EVs have become important diagnostic tools for ischemic heart disease (IHD) patients because they can be easily enriched in large quantities from bodily fluids, and their diverse cargo is resistant to degradation (108). The cargo of EVs reflects the true metabolic state of the originating cardiomyocytes. Studies have indicated that after acute myocardial infarction (AMI), compared to the control group, EVs in the blood of the AMI group are larger in size, with elevated circulating levels of CD144 but decreased levels of CCR6 and CXCR3 (109). Simultaneously, compared to peripheral blood after AMI, coronary artery blood showed a significant decrease in CCR6 levels. Multiple experiments have demonstrated that a hypoxic environment promotes cardiomyocyte secretion of EVs. Using surface antigen CD172a as a specific marker for cardiomyocyte-derived EVs, both in vitro and in vivo experiments showed that hypoxia increases EV yield (110). Small extracellular vesicles circulating in ST-segment

TABLE 4 EVs as biomarkers in CVDs (117-133).

EV Cargo	Source	Disease	Clinical outcomes	Ref.	
lncRNA Neat1	Cardiomyocytes	Cardiac	lncRNA Neat1 EV modulates the expression of P53 target	Long noncoding RNA-enriched vesicles secreted by	
		ischemia	genes, cell-cycle regulators and promoted cellular survival.	hypoxic cardiomyocytes drive cardiac fibrosis	
lncRNA CoroMarker	Monocytes	CAD	CoroMarker was stable in plasma because it was mainly in the EVs which is a stable, sensitive and specific biomarker for CAD	Plasma long non-coding RNA, CoroMarker, a novel biomarker for diagnosis of coronary artery disease	
miR-126	Plasma	CAD	Increased plasma EV miR-126 and miR-199a reduce the	MicroRNA expression in circulating microvesicles predicts	
miR-120	Fiasilia	CAD	risk of major cardiovascular outcomes in CAD patients	MicroRNA expression in circulating microvesicles predicts cardiovascular events in patients with coronary artery disease	
miR-155	Urine	CAD	Increased urinary EV miR-155 as a biomarker of CAD progression	MicroRNA-155 is decreased during atherosclerosis regression and is increased in urinary extracellular vesicles during atherosclerosis progression	
miR-92a	Endothelialium	CAD	EC-derived EV miR-92a is increased in CAD patients. miR-92a regulates angiogenesis in recipient EC	Atherosclerotic conditions promote the packaging of functional microRNA-92a-3p into endothelial microvesicles	
miR-1a	Plasma	STEMI	In STEMI patients, miR-1, -133a, -133b, and -499-5p were	Circulating microRNAs are new and sensitive biomarkers	
miR-133a/b			up-regulated, where as miR-122 and -375 were already	of myocardial infarction	
miR-499-5p			lower than in healthy subjects		
miR-122					
miR-375					
miR-192	Plasma	AMI	The 3 p53-responsive miRNAs that were upregulated in	Circulating p53-responsive microRNAs are predictive	
miR-194			the serum of post-AMI patients who experienced	indicators of heart failure after acute myocardial infarction	
miR-34a			development of HF within 1 year of AMI onset		
miR-24	Plasma	CAD	The plasma concentrations of exosomes and their cargo of	Coronary artery-bypass-graft surgery increases the plasma	
miR-208a/b			cardiac miRs increased in patients undergoing CABG and	concentration of exosomes carrying a cargo of cardiac MicroRNAs: an example of exosome trafficking out of the	
miR-122			were positively correlated with hs-cTnI		
miR-210				human heart with potential for cardiac biomarker discovery	
miR-9	Serum AIS		serum exosomal miR-9 and miR-124 are promising	MiR-9 and MiR-124 in the serum exosomes of acute	
miR-124	•		biomarkers for diagnosing AIS and evaluating the degree of damage caused by ischemic injury	ischemic stroke patients	
miR-134	Serum	AIS	RT-qPCR analysis showed that exosomal miR-134 was significantly increased in AIS patients within 24 h after stroke onset compared with that of control group	Increased serum exosomal miR-134 expression in the acute ischemic stroke patients	
miR-152-3p	Serum	AIS	Serum exosome miR-152-3p in patients with AIS was significantly lower than in healthy controls	Decreased Serum Exosomal miR-152-3p Contributes to the Progression of Acute Ischemic Stroke	
miR-21-5p	Plasma	IS	The plasma-derived exosomal miR-21-5p and miRNA-	Diagnosis of hyperacute and acute ischaemic stroke: the	
miR-30a-5p			30a-5p in combination are promising biomarkers for diagnosing IS and distinguishing among HIS, SIS, and RIS, especially miRNA-30a-5p for the diagnosis of the HIS phase	potential utility of exosomal MicroRNA-21-5p and MicroRNA-30a-5p	
miR-422a	Plasma	IS	Plasma exosomal miR-422a and miR-125b-2-3p may serve	Plasma Exosomal miR-422a and miR-125b-2-3p serve as	
miR-125b-			as blood-based biomarkers for monitoring and diagnosing in IS patients, with plasma exosomal miR-422a showing	biomarkers for ischemic stroke	
2-3p			the best diagnostic value		
miR-223	Serum	AIS	The level of miR-223 in circulating exosomes was elevated after onset of acute ischemic stroke, stroke patients with poor outcomes inclined to have a greater exosomal miR-223 expression	Increased circulating exosomal miRNA-223 is associated with acute ischemic stroke	
let-7b-5p	Plasma	AIS	In AIS patient, hsa-let-7b-5p, hsa-miR-16-5p, and hsa-	Differentially expressed miRNA profiles of serum derived	
miR-16-5p			miR-320c were upregulated, whereas hsa-miR-548a-3p	extracellular vesicles from patients with acute ischemic	
miR-320c	1		and hsa-miR-6808-3p	stroke	
miR-548a-3p	1				
miR-6808-3p					
miR-140-5p	Plasma	AIS	Within the first 6 h of symptom onset, three specific	MicroRNA expression profile in acute ischemic stroke	
miR-210-3p			miRNAs exhibited significant differential expression		
miR-7-5p			compared to other time points and healthy controls		
CD31/	Plasma	CAD	Increased plasma CD31/Annexin 5 EVs as an independent	Circulating CD31+/Annexin V+ microparticles correlate	
Annexin 5			predictor of cardiovascular events in CAD patients	with cardiovascular outcomes	
C1Q1A	Plasma	Myocardial	Plasma EV proteins as predictive biomarkers and	Plasma-derived extracellular vesicles contain predictive	
C5		infarction	therapeutic targets in myocardial infarction	biomarkers and potential therapeutic targets for Myocardial Ischemic (MI) injury	
GP1BA					
PPBP					

(Continued)

TABLE 4 Continued

EV Cargo	Source	Disease	Clinical outcomes	Ref.
APOC3				
CD144	Plasma	Myocardial injury	Increased plasma of CD144-EVs as predictor of cardiovascular complications	Significance of a multiple biomarkers strategy including endothelial dysfunction to improve risk stratification for
CD144	Endothelialium	Myocardial infarction	Endothelium-derived microparticles can improve prediction of future cardiovascular events in patients at high risk for CHD	cardiovascular events in patients at high risk for coronary heart disease
mCRP	Monocytes	CAD	mCRP in monocyte-derived EVs as biomarker of inflammatory process in CAD patients	Crp is transported by monocytes and monocyte-derived exosomes in the blood of patients with coronary artery disease
mCRP	Endothelialium	Myocardial infarction	EV transport and delivery of pro-inflammatory mCRP in AMI patients	Serum extracellular vesicle protein levels are associated with acute coronary syndrome
mCRP	Endothelialium	PAD	EC-derived EV mCRP is increased in patients with PAD, and was suggested as a predictor of vascular disease risk	Plasma Levels of Endothelial Microparticles Bearing Monomeric C-reactive Protein are Increased in Peripheral Artery Disease
CD11b	Urine	CAD	Increased CD45+ and CD11b+ and decreased CD16+ in	MicroRNA-155 Is Decreased During Atherosclerosis
CD16			urinary EVs are associated with CAD progression	Regression and Is Increased in Urinary Extracellular
CD45				Vesicles During Atherosclerosis Progression
Sphingolipid	Plasma	STEMI	EV lipid signature discriminates STEMI patients and may be used as therapeutic strategy	Analysis of urinary exosomal metabolites identifies cardiovascular risk signatures with added value to urine analysis

elevation myocardial infarction (STEMI) reflect the severity of myocardial injury in patients (111). By analyzing monocyte transcriptome data, a strong positive correlation was observed between PKIG and naïve B cells in AMI-monocyte-derived exosomes, RPL23 monocyte-derived exosomes showed a positive correlation, and OST4 potentially interacts with the p53 signaling pathway to mediate cardiomyocyte apoptosis (112). Through analysis of EVs in postmortem body fluids, significantly elevated levels of miR-486-5p were found in patients with high-grade atherosclerotic plaques, suggesting its potential as a biomarker for diagnosing acute myocardial infarction triggered by coronary atherosclerosis, including within the forensic field (113). Proteomic profiling of urinary EVs revealed reduced expression levels of UMOD protein in CAD patients' urinary EVs, this study contributes to the future use of uEVs as an emerging biomarker for the early, non-invasive diagnosis of CAD through protein differentials (114). Proteomic identification of plasma exosomes also revealed differences in the types and levels of protein expression in both STEMI and NSTEMI (115). Analysis of human plasma small extracellular vesicles (sEVs) using surface-enhanced Raman scattering (SERS) measurement technology combined with various machine learning methods demonstrated high sensitivity and specificity, suggesting that interdisciplinary fusion provides a promising strategy for non-invasive, safe, and high-precision early detection of CAD (116). Transcending the limitations of conventional MI biomarkers, EVs leverage their cellular origin specificity and molecular stability to redefine the paradigms of early warning, precise diagnosis, and dynamic monitoring in acute coronary syndromes through multi-omics dimensions.

3.3.3 Mechanisms of stem cell-derived EVs in MI treatment

The EV-based regenerative toolkit is reshaping post-injury myocardial functional restoration through synergistic engineered

delivery strategies and endogenous repair mechanisms (Table 5). Hypoxia and tumor necrosis factor-alpha (TNF-α) modulate extracellular vesicle release from human induced pluripotent stem cell-derived cardiomyocytes, this research aids in developing novel pharmacological strategies for MI (77). Extracellular vesicles from pluripotent stem cells or specially processed endogenous human serum exert cardioprotective effects post-injury (134). Mesenchymal stem cell-derived EVs (MSC-EVs) can rescue myocardial I/R injury by inducing cardiomyocyte autophagy via the AMPK and Akt pathways, reducing apoptosis and myocardial infarct size while improving cardiac function (135). Culturing MSC-EVs using a hollow fiber bioreactor-based three-dimensional (3D) system yields higher quantities with similar biological functions compared to traditional two-dimensional (2D) culture (136). Monocyte mimics confer stronger recruitment characteristics to MSC-EVs in I/R models through Mac1/LFA1-ICAM-1 adhesion molecule interactions, enhancing their targeting efficiency to injured myocardium and improving EV delivery to ischemic-damaged myocardium (137). Exosomes derived from adipose stem cells (ASC-Exos) promote angiogenesis in ischemic hindlimbs and hearts in mice by delivering miR-31 via the miR-31/FIH1/ hypoxia-inducible factor-1α (HIF-1α) signaling pathway, alleviating ischemic heart disease (138). ASC-Exos-delivered miR-205 significantly reduced cardiomyocyte apoptosis while promoting angiogenesis and microvascular endothelial cell proliferation to improve cardiac function after myocardial infarction (139). Adipose stem cell-derived exosomal miR-196a-5p and miR-425-5p prevented ischemia-induced mitochondrial dysfunction and reactive oxygen species (ROS) production in cardiomyocytes, increased angiogenesis, and polarized macrophages towards the anti-inflammatory immunophenotype (140). Furthermore, miR-196a-5p reduces and reverses myofibroblast activation and lowers collagen expression to inhibit myocardial fibrosis. ASCs-EVs-miR-221

TABLE 5 Potential treatment of EVs in MI (138, 139, 145-157).

Source	Cargos	Signaling pathway/target	Function	Reference title
Mesenchymal stem cell derived EVs	miR-210	Efna3↓	proangiogenic effects	Mesenchymal stem cells-derived extracellular vesicles, via miR- 210, improve infarcted cardiac function by promotion of angiogenesis
	_	NF-κB↓ → Nrf2/HO- 1↑	anti-inflammation and polarizing M2 macrophage	Exosomes secreted by FNDC5-BMMSCs protect myocardial infarction by anti-inflammation and macrophage polarization via NF-κB signaling pathway and Nrf2/HO-1 axis
	miR-223	P53/S100A9↓	relieved myocardial fibrosis and inflammation infiltration	Human umbilical cord mesenchymal stem cell-derived extracellular vesicles loaded with miR-223 ameliorate myocardial infarction through P53/S100A9 axis
	miR-212-5p	NLRC5/VEGF/TGF- beta 1/SMAD↓	attenuate MI/cardiac fibrosis	MSCs-Derived Extracellular Vesicles Carrying miR-212-5p Alleviate Myocardial Infarction-Induced Cardiac Fibrosis via NLRC5/VEGF/TGF-β1/SMAD Axis
Adipose stem cell derived EVs	miR-31	FIH1/HIF-1α↑	proangiogenic effects	Exosomes from adipose-derived stem cells alleviate myocardial infarction via microRNA-31/FIH1/HIF-1α pathway
	miR-205	HIF-1α/VEGF↑	proangiogenic effects	ADSC-derived exosomes attenuate myocardial infarction injury by promoting miR-205-mediated cardiac angiogenesis
	miR-24-3p	Nrf2↑	cardioprotective effects	Adipose stem cell-derived nanovesicles for cardioprotection: production and identification of therapeutic component
Induced pluripotent stem cell derived EVs	miR-106a-363	Jag1/Notch3/Hes1↓	myocardial repairment	miR-106a–363 cluster in extracellular vesicles promotes endogenous myocardial repair via Notch3 pathway in ischemic heart injury
	miR-145/let-7/ miR-302a-5p	BMP-4/PDGFα/ VEGF-C↑	proangiogenic effects	Induced Pluripotent Stem Cell (iPSC)–Derived Extracellular Vesicles Are Safer and More Effective for Cardiac Repair Than iPSCs
Hypoxia treated stem cell derived EVs	miR224-5p	$TXNIP\downarrow \rightarrow HIF-1\alpha\uparrow$	protect myocardial cells from hypoxic vulnerability	Extracellular vesicles from hypoxia-preconditioned mesenchymal stem cells alleviates myocardial injury by targeting thioredoxin-interacting protein-mediated hypoxia-inducible factor- 1α pathway
Blueberry-treated stem cell derived EVs	-	-	anti-inflammatory and anti- apoptotic effects	The Therapeutic Effects of Blueberry-Treated Stem Cell-Derived Extracellular Vesicles in Ischemic Stroke
Engineered stem cell derived EVs	p38α antagonistic peptides	Mapk14↓ → α-SMA/ collagen I↑	Inhibitory effect on CFs fibrosis	Sodium Alginate Hydrogel Infusion of Bone Marrow Mesenchymal Stem Cell-Derived Extracellular Vesicles and p38α Antagonistic Peptides in Myocardial Infarction Fibrosis Mitigation
	miR-210-3p	EFNA3↓	promoted angiogenesis	Extracellular Vesicles from NMN Preconditioned Mesenchymal Stem Cells Ameliorated Myocardial Infarction via miR-210-3p Promoted Angiogenesis
	Melatonin	ROS↓	inhibiting oxidative stress	Melatonin Engineered Adipose-Derived Biomimetic Nanovesicles Regulate Mitochondrial Functions and Promote Myocardial Repair in Myocardial Infarction
Human umbilical cord mesenchymal stem cell- derived Evs	-	PI3 K/Akt↑	alleviate ER stress-induced apoptosis	Extracellular Vesicles Derived from Human Umbilical Cord Mesenchymal Stromal Cells Protect Cardiac Cells Against Hypoxia/Reoxygenation Injury by Inhibiting Endoplasmic Reticulum Stress via Activation of the PI3K/Akt Pathway

significantly enhanced proliferation and expression of antiapoptotic proteins in H9C2 cells (a cardiomyocyte cell line derived from the left ventricle of Sprague Dawley rats) (141). Plasma-derived extracellular vesicles from myocardial infarction patients inhibited TNF-α-induced cardiomyocyte death by suppressing TNF-α expression. yREX3 alleviates myocardial ischemic injury through selective DNA methylation (142). The non-coding RNA yREX3 encapsulated within EVs mediates the epigenetic silencing of the protein interacting with C kinase 1 (Pick1) via methylation of upstream CpG sites by triggering widespread transcriptomic changes in macrophages. Simultaneously, yREX3 interacts with polypyrimidine tractbinding protein 3 (PTBP3) to methylate the Pick1 gene locus. Inhibiting Pick1 in macrophages enhances Smad3 signaling and boosts exocytosis, thereby minimizing cardiac necrosis in rats with myocardial infarction. The neonatal mammalian heart possesses the ability to regenerate after injury by inducing cardiomyocyte proliferation. Experiments demonstrated that EVs from regenerating neonatal mouse heart tissue after apical resection surgery (AR-Neo-EVs) exhibited stronger proproliferative, anti-apoptotic, and pro-angiogenic activities compared to EVs from neonatal mouse heart tissue (Neo-EVs) (143). Subsequent studies confirmed that delivering AR-Neo-EVs via sodium alginate hydrogel microspheres is an effective method for treating myocardial infarction. Conversely, vesicles derived from myocardial tissue of postnatal day 8 mice (P8-EVs) significantly promoted M1-like macrophage polarization, enhanced phagocytosis, and influenced macrophage factor secretion by activating the Pak2-Erk1/2 axis, thereby inhibiting neonatal cardiomyocyte proliferation. Human amniotic fluid stem cell-derived EVs demonstrated potential for cardiomyocyte renewal (144). It shows that stem cell derived

EVs offer enhanced translational potential as delivery vehicles for myocardial regeneration therapy by improving cardiac targeting, coordinating multiple repair pathways, and preserving bioactive molecule stability.

3.3.4 Potential therapeutic effect of engineered EVs in MI

Source platelet lysate-derived EVs (SCPL-EVs) carry a range of trophic factors and multiple recognized cardioprotective miRNAs (85). Experiments demonstrated their protection of rodent and human cardiomyocytes from I/R injury, stimulation of angiogenesis in human cardiac microvascular endothelial cells, and reduction of scar formation, thereby improving cardiac function. miR-4496 and miR-4691-5p are highly enriched in human embryonic stem cell-derived EVs (hESC-eEVs) (158). Overexpression of miR-4496 or miR-4691-5p led to increased endothelial cell (EC) tube formation and wound closure in vitro, confirming the novel pro-angiogenic functions of these miRNAs. CD47 signaling can help evade macrophage clearance by binding to signal regulatory protein alpha (SIRPα) (159). Purified CD47-EVs were encapsulated with miR-21a (a specific anti-apoptotic miRNA) via electroporation to construct Electro-CD47-EVs. effectively This approach improved biodistribution of these miR-21a-containing EVs in the heart, prolonged their retention time in circulation, and reduced phagocytic clearance, providing new insights for potential therapeutic tools for myocardial I/R injury. EVs were functionalized on their surface with cardiac-targeting peptides (CTP) via genetic modification to generate cardiac-targeted EVs (CTP-EVs). Curcumin was then loaded into these CTP-EVs (CTP-EV-Cur) to specifically deliver curcumin to the heart, it higher bioavailability and enhancing cardioprotective efficiency (160). Interestingly, after validating that miR-144-3p is a primary contributor to the therapeutic effects mediated by curcumin, co-loading curcumin and miR-144-3p into the CTP-EVs (CTP-EVs-Cur) retained their active targeting ability while exhibiting cardioprotective effects both in vitro and in vivo. Targeting miR-222-engineered extracellular vesicles (TeEVs), tailored with CTPs, are developed as ischemic TeEV therapeutics. These TeEVs are encapsulated within mechanical hydrogels to create injectable TeEV-loaded cardiac patches, enabling minimal invasiveness to attenuate IRI. The injectable patches facilitate the precise targeting of TeEVs for the efficient rescue of damaged cells. Persistent delivery of TeEVs into the infarcted region alleviates acute IRI and mitigated remodeling post IRI. This is linked to focal adhesion activation, cytoskeleton force enhancement, and nuclear force-sensing preservation. These findings may pave the way for force-sensing approaches to cardiac therapy using bioengineered therapeutic patches (161). Evidence demonstrates that engineered EVs via peptide-directed targeting, therapeutic cargo loading, and hydrogel-based delivery systems, significantly enhance cardiac drug accumulation, prolong circulation half-life, and enable sustained release in injured areas, offering novel precision strategies for myocardial I/R injury.

3.4 Mechanisms of EVs in heart failure and cardiac remodeling

Heart failure (HF) represents a complex clinical syndrome characterized by progressive cardiac remodeling following injury. Extracellular vesicles have emerged as pivotal mediators in this process, influencing pathological remodeling, serving as biomarker sources, and offering novel therapeutic opportunities. This section delineates the multifaceted roles of EVs in HF, spanning their contributions to disease mechanisms, diagnostic potential, and therapeutic applications.

3.4.1 Mechanisms of EVs in HF damage and remodeling

Chronic heart failure (CHF) is associated with redox imbalance. In CHF induced after MI, EVs enriched with cardiac miRNAs can mediate enhanced oxidative stress and heightened cardiac and central sympathetic nerve excitation by targeting and downregulating the Nrf2/antioxidant signaling pathway (162). The miR-155-5p in heart-derived EVs promotes macrophage M1 polarization by activating the JAK2/STAT1 pathway, leading to local cardiac inflammation and even triggering systemic inflammation in distant organs (106). Conversely, some EVs exhibit protective properties. In EVs derived from human induced pluripotent stem cell-derived cardiomyocytes (hCM-EVs), transfection of human cardiac fibroblasts (hCFs) with miR-24-3p prevented TGF-β1-mediated induction of FURIN, CCND1, and SMAD4-miR-24-3p target genes involved in TGFβ1-dependent fibrosis—regulating the transformation of hCFs to myofibroblasts (163). Furthermore, the long noncoding RNA Tcf21 antisense RNA inducing demethylation (lncRNA-TARID), enriched in EVs, was found to upregulate Tcf21 expression, which suppresses transforming growth factor beta (TGF- β) signaling and myofibroblast differentiation to inhibit cardiac fibrosis (164). Circular RNA circWhsc1 was found enriched in neonatal mouse hearts, particularly in cardiac ECs, and further upregulated in ECs and EC-derived EVs under hypoxic conditions. EV-derived circWhsc1 activates TRIM59 by enhancing its phosphorylation, thereby strengthening TRIM59 binding to STAT3, phosphorylating STAT3, and inducing cardiomyocyte (CM) proliferation (165). These findings highlight the dual role of EVs in heart failure-mediating both detrimental processes such as oxidative stress, inflammation, and sympathetic overactivation, as well as protective mechanisms including antifibrotic and pro-regenerative effects-suggesting their potential as therapeutic targets in HF management.

3.4.2 Diagnostic potential of EVs in HF

The molecular cargo of EVs provides a rich source for discovering biomarkers to monitor HF progression and predict outcomes. Temporal changes in the ratio of coagulation and fibrinolysis pathway proteins in low-density lipoprotein-associated EVs (LDL-EVs) may be more advantageous than currently used plasma biomarkers for predicting reverse left ventricular remodeling after acute myocardial infarction (AMI) (166).

Furthermore, downregulated circCEBPZOS was detectable in serum EVs from patients with adverse cardiac remodeling. CircCEBPZOS acts as a competitive endogenous RNA (ceRNA) by directly binding miR-1178-3p, thereby promoting transcription of phosphoinositide-dependent kinase-1 (PDPK1), and its reduced levels are associated with deleterious post-MI remodeling and worse cardiac function (167). In summary, EVs offer a promising platform for the identification of novel biomarkers that reflect dynamic pathophysiological changes in HF, enabling early risk stratification, precise monitoring of disease progression, and improved prediction of clinical outcomes beyond conventional diagnostic tools.

3.4.3 Therapeutic potential of EVs in HF

The therapeutic application of EVs, particularly from stem cells and engineered sources, holds significant promise for mitigating cardiac remodeling and improving function in HF. Stem cells from multiple sources offer potential for cardiac remodeling after myocardial infarction and heart failure. EVs from human umbilical cord mesenchymal stem cells (hUCMSCs) pre-treated with NMN, highly expressing miR-210-3p, significantly enhanced tube formation, migration, and proliferation capabilities in human umbilical vein endothelial cells (HUVECs) (153). N-EVs (NMNpretreated EVs) promoted infarct healing through miR-210-3p targeting EFNA3 to improve angiogenesis. Hypoxia-induced adipose-derived stem cell EVs (ADSC-EVs) exhibited stronger proliferation, migration, and tube formation capabilities in HUVECs compared to normoxic ADSC-EVs and demonstrated neovascularization in mouse models Intrapericardial injection of engineered EVs containing lncRNA-TARID upregulated Tcf21 expression in epicardium-derived cells and improved cardiac function and histology in mice and porcine MI models (164). In particular, milk EVs derived from colostrum contain various anti-inflammatory factors facilitating the transition from inflammation to proliferation phase, as well as factors for tissue remodeling and angiogenesis (168). Bovine milk-derived exosomes (BM-Exos) alleviate fibrosis through pro-angiogenic mechanisms, thereby improving cardiac fibrosis. In both in vivo models of isoproterenol (ISO)-induced cardiac fibrosis and in vitro OGD-treated HUVECs, BM-Exos were demonstrated to reduce extracellular matrix (ECM) deposition, improve cardiac function, and enhance pro-angiogenic growth factors (91).

Collectively, EVs play a dual role in heart failure, both as drivers of pathological remodeling and as vectors for innovative diagnostics and therapeutics. Understanding their intricate mechanisms and harnessing their potential paves the way for novel precision medicine approaches in cardiovascular disease.

3.5 Mechanisms of EVs in stroke

Acute ischemic stroke (AIS) is the most common type of stroke. Despite significant advances in stroke treatment, including but not limited to reperfusion therapies, the limited time window after stroke onset, the effectiveness of treatments, and associated risks highlight the necessity of researching novel diagnostic and therapeutic strategies.

3.5.1 Mechanisms of EVs in stroke damage and repairment

Analogous to the peripheral vascular system, under hypoxic conditions, cerebral microvascular endothelial cell (CEC)derived exosomes promote microglial M1 polarization and endothelial cell injury by delivering circ-0000495 (169). Conversely, ischemic CEC-derived small extracellular vesicles (CEC-sEVs), enriched in miR-27a, play a crucial role in the ischemic brain repair process by promoting axonal remodeling and improving neurological outcomes (170). sEVs derived from activated neurons, enriched in miR-100-5p, can activate the NFκB pathway, causing abnormal activation of adjacent neurons and expanding the scope of neuronal damage, while also driving microglial activation to exacerbate neuroinflammation (171). Apoptosis induced by AIS releases substances that stimulate microglia to activate pro-inflammatory pathways and enhance EV production, or have the opposite effects. M2-polarized microglia-derived EVs, enriched in miR-124, reduce astrocyte proliferation by decreasing Signal Transducer and Activator of Transcription 3 (STAT3) gene signaling, thereby reducing glial scar formation and promoting post-stroke recovery (172). Furthermore, M2 microglia-derived small EVs, enriched in miR-93-5p and miR-25-3p, downregulated the expression of TGFBR, PTEN, and FOXO3 in neural stem cells (NSCs), increasing NSC proliferation and neuronal differentiation. Inhibition of the extracellular matrix protein Sema3A in astrocyte-derived EVs during the subacute phase of stroke suppresses activated astrocytes in the peri-infarct cortex and enhances neurological functional recovery (173). Exosomes secreted by ischemic astrocytes treated with a Sema3A inhibitor further increase Prostaglandin D2 Synthase (PTGDS) expression and contribute to post-stroke axonal growth and elongation, as well as functional recovery (174). In spontaneous intracerebral hemorrhage (ICH), red blood cell-derived microparticles inhibit hematoma growth in ICH (175). EVs in brain injury bidirectionally regulate neuroinflammation and regeneration via non-coding RNA delivery including endothelial and neuron-EVs exacerbate microglial polarization excitotoxicity, whereas ischemia-adapted endothelial and glial vesicles promote axonal remodeling and suppress scarring.

3.5.2 Diagnostic effect of EVs in stroke

To date, stroke diagnosis primarily relies on clinical symptom assessment and neuroimaging, suitable blood biomarkers for stroke detection via blood tests remain undeveloped. Extracellular vesicles (EVs), on the one hand, can cross the blood-brain barrier (BBB), and more importantly, reflect the real-time status of the secreting cells, thereby offering the possibility of serving as diagnostic biomarkers for brain diseases (Table 4). Examination of plasma EV markers (CD9+) alongside neuronal markers (L1CAM, CD171), astrocytic markers (EAAT1), and oligodendrocyte markers (MOG) demonstrated elevated levels of astrocyte-derived extracellular vesicles (ADEs) within the first month after stroke in clinical settings (176). Multiple studies have proposed that EVs derived from plasma,

through analysis of their miRNA and proteomics, confirm their role in the pathophysiology of acute stroke and offer potential biomarkers for stroke diagnosis.

3.5.3 Potential therapeutic effect of stem cell-derived EVs in stroke

After AIS, both bone marrow mesenchymal stem cell-derived EVs (BMSC-EVs) and brain endothelial cell-derived EVs (BEC-EVs) protect blood-brain barrier (BBB) integrity, with BMSC-EVs enhancing efficacy in improving neurological function. Both EVs attenuate BBB permeability by inhibiting the Cav-1/CD147/ VEGFR2/MMP pathway and restore tight junction protein expression by counteracting abnormal endocytosis, repairing and enhancing BBB integrity (177). MSC-derived EVs in the treatment of ischemic cerebrovascular disease exhibit mechanisms and efficacy similar to the previously mentioned MSC-EVs in repairing vascular endothelium during myocardial ischemia. Adipose-derived MSCs, via their contained miR-760-3p, inhibit CHAC1 expression to suppress the progression of ferroptosis, a key process in ischemic stroke, effective delivery to the brain via intranasal (IN) administration alleviates cerebral ischemia/reperfusion injury (178). Dental pulp stem cell-derived EVs, enriched in miR-877-3p, interact with Bclaf1, reducing neuronal apoptosis, decreasing infarct size, and improving cerebral edema (179). Extracellular vesicles derived from blueberry-treated mesenchymal stem cells (MSCs) demonstrated significantly greater protection against neuronal cell death in oxygen-glucose deprivation (OGD) cells and middle cerebral artery occlusion (MCAo) animal models compared to the untreated group, both EV treatments increased cell viability, reduced cerebral infarction area in animal experiments, and improved behavioral assessments (151). EVs derived from Panax notoginseng, a commonly used Chinese herbal medicine, contain lipid components that suppress the inflammatory response by shifting the microglial phenotype from M1 to M2, alleviating cerebral ischemia/reperfusion injury (CI/R) (180). Furthermore, studies found that these lipids act on the PI3K/ Akt pathway to reduce cerebral infarction volume and exert therapeutic effects. Momordica charantia (bitter melon) exosomes, via miR-5266, activate the AKT/GSK3β pathway, inhibit neuronal apoptosis in stroke, alleviate ischemiareperfusion injury, and maintain BBB integrity (181). Application of NSC-derived exosomes (NSC-Exos) to an in vitro mouse neuronal injury model before and/or during OGD combined injury verified their significant protective effect on astrocytes. Mice treated with intravenously injected NSC-Exos after stroke and reperfusion injury all exhibited significantly reduced infarct volumes. On one hand, NSC-Exos promote neuroprotection by transferring miR-150-3p, which targets CASP2, inhibiting neuronal apoptosis after brain injury, on the other hand, they enhance the nuclear translocation of Nrf2 to counteract oxidative stress and reduce inflammatory responses (182). Simultaneously, NSC-Exos demonstrated functions promoting neuronal axon elongation and angiogenesis in HUVECs, offering a potential therapeutic avenue for AIS. Coincubation of BDNF protein (a widely distributed neurotrophic factor) with hNSC-Exos for 24 h loaded BDNF into the exosomes, resulting in BDNF-hNSC-Exos, which exhibited stronger anti-apoptotic and pro-differentiation capabilities than hNSC-Exos. In a rat ischemic stroke model, BDNF-hNSC-Exos effectively suppressed microglial expression, inhibited inflammation, promoted endogenous NSC differentiation, thereby reducing infarct size and improving neurological function (80). Experimental data confirm that engineered extracellular vesicles subjected to molecular preconditioning, bioactive cargo loading, or delivery route optimization synergistically enhance barrier restoration, neuroprotection, and regenerative efficacy, with significant improvement neurological outcomes in stroke models.

3.5.4 Potential therapeutic effect of engineered EVs and PDEVs in stroke

Utilizing the property of bacterial-derived outer membrane vesicles (OMVs) to target neutrophils, pioglitazone (PGZ) was encapsulated into OMVs. The resulting OMV@PGZ nanoparticles inherited the functions associated with the bacterial outer membrane, facilitating targeted uptake by neutrophils (183). Results indicated OMV@PGZ enhanced PGZ delivery to the brain for treating ischemic stroke, while inhibiting nucleotide-binding oligomerization domain-like receptor protein 3 (NLRP3) inflammasome activation and ferroptosis, reducing inflammatory responses after reperfusion injury to exert neuroprotective effects. Experiments in tMCAO mice treated with Lactobacillus plantarum-derived EVs (LEVs) confirmed their efficacy in reducing apoptosis of ischemic neurons, potentially by inhibiting neuronal apoptosis via the miR-101a-3p/c-Fos/TGF-β axis, thereby preventing ischemiainduced brain injury (184). Engineered EVs in cerebrovascular diseases are mostly used to deliver therapeutic components. Utilizing the targeting ability of RGD peptide and the cellpenetrating ability of Angiopep-2 peptide, dual-modified adipose-derived stem cell EVs demonstrated high transcellular permeability across the BBB in vitro, while targeting ischemic blood vessels, in vivo, intravenous administration achieved rapid accumulation of active components in the ischemic lesion area. By modifying the surface of M2 microglia-derived EVs with rabies virus glycoprotein peptide 29 (RVG29), miR-221-3p and miR-423-3p within them exert anti-apoptotic effects on neurons via the p38/ERK signaling pathway in stroke (185). Momordica charantia-derived exosome-like nanoparticles (MC-ELNs) were demonstrated to cross the blood-brain barrier and accumulate in ischemic regions. They exerted neuroprotection by delivering miR-5266 to suppress matrix metalloproteinase 9 (MMP-9) expression which preserving BBB integrity and activating the AKT/GSK3β signaling pathway, ultimately reducing cerebral infarct volume and improving neurological deficits in MCAO rats (181). Researches show that engineered EVs and plantderived EVs, via targeting modifications and therapeutic cargo loading, breach the blood-brain barrier for targeted delivery to cerebral ischemic foci, exerting neuroprotection through suppression of neuroinflammation and programmed cell death.

3.6 Mechanisms of EVs in valvular heart disease

Valvular heart disease (VHD) encompasses a range of structural and functional abnormalities of heart valves, leading to impaired blood flow. Extracellular vesicles play critical roles in the pathogenesis, diagnosis, and treatment of VHD, serving as key mediators of intercellular communication and potential vehicles for therapeutic intervention.

3.6.1 Mechanisms of EVs in VHD pathogenesis

Valvular cells under stress or injury release EVs carrying a distinct molecular cargo that reflects the underlying pathophysiological processes. EVs derived from diseased valves exhibit increased levels of inflammatory cytokines, IL-6, TNF-α and calcification-promoting factors such as alkaline phosphatase, osteopontin, bone morphogenetic protein 2 (BMP2), and MMPs, which drive inflammation, calcification, and tissue remodeling in VHD (186). Additionally, dysregulation of specific miRNAs within EVs-such as downregulation of miR-143 and miR-145 which involved in valvular smooth muscle cell homeostasis and upregulation of miR-21, contributes to disease progression by promoting fibrosis and inhibiting apoptosis (187, 188). Lipidomic analyses further reveal elevated levels of ceramides, sphingomyelin, and oxidized phospholipids in EVs from diseased valves, which are associated with oxidative stress and apoptosis (189). EVs from calcified human aortic stenotic valves (AS-EVs) are highly enriched in TF and activate valvular endothelial cells via the AT1R/NADPH oxidase/SGLT2 pathway. This triggers a pronounced pro-oxidant, pro-inflammatory, and pro-thrombotic response, leading to endothelial dysfunction and heightened thrombogenicity, a process that can be mitigated by SGLT2 inhibition (190). Overall, these findings underscore that valvular-derived EVs are not merely bystanders but active contributors to the pathogenesis of valvular heart disease, driving a cascade of inflammatory, calcific, thrombotic, and fibrotic processes.

3.6.2 Diagnostic potential of EVs in VHD

EVs offer a promising non-invasive approach for early detection and risk stratification of VHD. Their molecular cargo, including proteins, miRNAs, and lipids, provides a snapshot of valvular health and disease activity. For instance, EVs associated proteins such as MMPs and osteopontin can serve as early biomarkers before structural changes are detectable via echocardiography (191). miRNA signatures, such as miR-21, miR-143 and miR-145 within EVs can differentiate between VHD subtypes and predict disease progression (187). Multimarker EV panels integrating protein, miRNA, and lipid profiles are under development to enhance diagnostic accuracy and prognostic value.

3.6.3 Therapeutic potential of EVs in VHD

EVs hold significant promise as therapeutic agents in VHD, offering targeted and minimally invasive treatment strategies.

Mesenchymal stem cell-derived EVs (MSC-EVs) exhibit potent anti-inflammatory effects by delivering anti-inflammatory cytokines (e.g., IL-10, TGF-β) and miRNAs (e.g., miR-146a), which attenuate valvular inflammation and fibrosis (192). Engineered EVs loaded with anti-calcific miRNAs (e.g., miR-302/367 cluster) can inhibit osteogenic differentiation of valvular interstitial cells and reduce calcification in preclinical models (193). EVs derived from telocytes (TCs) were shown to ameliorate calcific aortic valve disease (CAVD) in mice by delivering miR-30b, which targets Runx2 and inhibits the Wnt/ β-catenin pathway, thereby reducing VICs osteogenic differentiation and calcium deposition (194). EVs derived from endothelial progenitor cells (EPCs) promote endothelial repair and angiogenesis, thereby mitigating fibrosis and improving valve function (195). Furthermore, EVs can be functionalized to deliver targeted therapeutics including statins and anti-fibrotic drugs directly to valvular tissue, enhancing treatment efficacy and reducing systemic side effects (196).

Collectively, EVs play dual roles in VHD, as mediators of disease progression and as innovative tools for diagnosis and therapy. Harnessing their potential through further research and clinical validation may lead to novel precision medicine approaches for valvular heart disease.

4 Challenges and development directions in the future

4.1 Technical hurdles

One of the primary challenges hindering the clinical application of EVs in CVDs is rooted in the technical limitations associated with their isolation, purification, and characterization. Owing to the inherent heterogeneity of EV populations, which are secreted by various cell types and exist in multiple subtypes such as exosomes, microvesicles, and apoptotic bodies, classical isolation methods often yield preparations that are contaminated with protein aggregates, lipoproteins, or other non-vesicular materials, thereby compromising the purity and, consequently, the reproducibility of downstream therapeutic applications (197).

Additional technical challenges involve limitations in cargo loading methodologies. The efficient incorporation of therapeutic molecules—such as specific miRNAs, proteins, or drugs—into EVs demands innovative strategies that preserve vesicle integrity while ensuring sufficient encapsulation efficiency. While techniques like passive incubation, electroporation, and sonication have been employed, each exhibits intrinsic drawbacks including cargo aggregation, potential vesicle disruption, and low loading efficiency, collectively constraining their utility. Critically, these challenges compound broader technical barriers spanning from the reproducible isolation of homogeneous EV populations to the standardization of characterization methods essential for evaluating therapeutic efficacy and pharmacokinetics. Collectively, these hurdles represent a fundamental impediment to developing reliable EV-based cardiovascular therapeutics.

TABLE 6 Clinical trials involving EVs in CVDs.

Disease	Source	Meanings	Reference
Myocardial Infarction	Exosomes	Determine the miRNA expression profile in peripheral blood exosomes of patients with myocardial infarction	NCT04127591
Myocardial Infarction	MSC-Exosomes	Protect the heart against the harmful effects of ischemia and also IR, preserve cardiac function, reduce the zone of MI	NCT05669144
Acute Ischemic Stroke	MSC-Exosomes	Improves functional recovery and enhances neurite remodeling, neurogenesis, and angiogenesis after AIS	NCT03384433
Acute Ischemic Stroke	GD-iExo-003	Explore the effect of exosomes derived from HIPSC in AIS	NCT06138210
Acute Ischemic Stroke	MSC-Exosomes	Safety of cultured allogeneic adult umbilical cord derived MSC Exosomes intranasal instillations for the treatment of stroke	NCT05158101
Acute Ischemic Stroke	Exosomes	Characterization of circulating EVs/Exosomes as new predictive biomarkers in Stroke patients	NCT05370105
Acute Ischemic Stroke	RBC-Evs	RBC deformability will serve as a biomarker of the conditioning response and predictor of the clinical outcome in stroke patients	NCT04266639
Acute Ischemic Stroke	NSC-EVs	Clinical trial to assess the safety and efficacy of NouvSoma001 in the treatment of ischemic stroke	NCT06612710
Acute Ischemic Stroke	EVs	EVs profiles may change during acute stroke, in the chronic stroke phase, and according to the level of cSVD, may act as disease biomarkers	NCT06257823
Acute Ischemic Stroke	Serum EVs	Evaluation of predictive circulating biomarkers for rehabilitation recovery in ischemic and haemorrhagic stroke patients	NCT06871800
Stroke	EVs	Investigating the association of EVs and dysregulated coagulation in the prediction of Stroke	NCT05645081
Heart Failure	EVs	Investigate the expression of EVs RNAs during admission for and decongestion of ADHF patients and determine their diagnostic and prognostic potential	NCT06169540
Atherosclerosis	Fat tissue EVs	Research about EVs signaling in obesity and cardiometabolic diseases	NCT06408961
Hypertension	Urinary- Exosomes	Determine the concentrations and variabilities of urinary exosomal sodium channels and plasma angiotensins in patients with difficult-to-treat arterial hypertension and to investigate their dependency on clinical parameters and sampling conditions	NCT03034265
Hypertension	Endothelial-EVs	Assessing the prognostic significance of endotehlial dysfunction in hypertension	NCT06316271
Heart Failure	Plasma-Exosomes	Further explore the treatment's impact on heart failure-related inflammatory markers	NCT06501768
Atrial Fibrillation	Fat tissue EVs	Investigate the role of epicardial fat derived exosomes in patients who suffer from atrial fibrillation	NCT03478410
Heart Surgery	Blood MVs/ Exosomes	Study of role of blood microvescicles and exosomes in patients with graft occlusion after aortocoronary bypass surgery	NCT05411445

4.1.1 Limitations of EV-based biomarkers in CVDs

Biological variability presents further obstacles. Fluctuations in EV molecular cargo driven by physiological states including circadian rhythms or exercise, and comorbidities such as diabetes or renal dysfunction, obscure disease-specific signatures (198, 199). This instability is intensified by inconsistent pre-analytical variables, particularly variations in sample collection protocols, storage conditions, and processing delays across research cohorts. Critical validation deficiencies hinder clinical implementation (200). Most proposed EV biomarkers emerge from small discovery cohorts without external validation through multicenter trials. Longitudinal evidence demonstrating prognostic utility in diverse populations remains scarce, limiting risk stratification capabilities. Furthermore, the significant cost and technical demands of EV profiling compared to conventional biomarkers create economic barriers. Regulatory pathways for EV-based diagnostics are currently undefined. These collective challenges impede the translation of EV biomarker research into cardiovascular clinical practice (201).

4.2 Clinical trials involving EVs in CVDs

Over the last five years, EVs have emerged as promising biomarkers and therapeutic agents in cardiovascular disease

owing to their ability to reflect the state of the parent cell and modulate regenerative processes. Their potential has been investigated across a wide spectrum of CVDs such as myocardial infarction, heart failure, ischemia/reperfusion injury, stroke, and atherosclerosis. A growing number of clinical trials have been initiated to explore the translational value of EVs.

A detailed analysis of the current clinical landscape, as summarized in Table 6, reveals several key trends. First, research on stroke, particularly acute ischemic stroke, constitutes a predominant focus of registered EV clinical trials in the CVDs field (e.g., NCT05877016, NCT04202783, NCT05191381). This highlights the urgent clinical need for novel diagnostic and therapeutic strategies in neurovascular diseases. Second, the majority of these registered studies are in early phases (I or II) and are primarily observational in nature. These trials aim to correlate specific EV signatures (e.g., surface proteins, miRNA content) with disease diagnosis, severity, or prognosis outcomes, laying the groundwork for EV-based liquid biopsies.

Regarding the studied populations, most trials enroll patients with acute ischemic stroke, post-myocardial infarction, or chronic heart failure, reflecting the focus on acute events and subsequent maladaptive remodeling. In terms of interventional strategies, the few existing therapeutic trials mainly investigate the safety and efficacy of allogeneic MSC-EVs. For instance, the NCT05191381 trial employs intravenous infusion of allogeneic MSC-EVs for

ischemic stroke treatment, while the NCT04327635 trial explores the intramyocardial injection of MSC-EVs to treat heart failure following myocardial infarction. In contrast, interventional clinical trials specifically targeting atherosclerosis itself remain scarce, indicating a significant gap and a future direction for translating mechanistic insights into therapeutic applications.

4.3 Difficulties in clinician translational

Beyond technical challenges, significant difficulties in translational medicine impede the clinical implementation of extracellular vesicle (EV)-based therapies for cardiovascular conditions. A pivotal aspect of translation is the scalability of EV production, inherently linked to the donor cell source and culture conditions. Many preclinical studies employ mesenchymal stem cells (MSCs) or cardiac progenitor cells to harvest EVs. However, these cells exhibit variable secretion rates, and the EV yield is often insufficient for human therapeutic dosing, necessitating the development of bioreactor-based systems and optimized cell expansion protocols. This lack of scalability is compounded by the difficulty in ensuring batch-to-batch consistency in EV composition and bioactivity, which remains sensitive to factors such as donor cell condition, culture medium, and processing variations.

Another major translational hurdle concerns the vivo biodistribution, clearance, and targeted delivery of EVs. Intravenously administered EVs are frequently cleared rapidly by the mononuclear phagocyte system (MPS), leading to suboptimal accumulation at intended cardiovascular sites. This is partly due to nonspecific uptake by organs like the liver, spleen, and lungs. Consequently, high dosages are required, which may be infeasible from a manufacturing standpoint and unacceptable from a safety perspective, especially given potential off-target effects. Furthermore, the dosing paradigms for EVs therapeutics remain inadequately defined; the effects of administered EV concentrations are influenced not only by the absolute dose but also by the interplay between EV stability, tissue targeting, and payload diversity. Due to their natural biodistribution, selective homing of EVs to diseased cardiovascular tissues remains suboptimal, and engineering modifications to enhance targeting have not yet yielded consistent improvements in in vivo delivery efficiency.

Safety concerns are also paramount. Despite the reduced immunogenicity of exosome-based therapies relative to cell-based approaches, potential risks persist, including the inadvertent transfer of oncogenic or pro-inflammatory genetic material, immunomodulatory effects that could provoke adverse reactions, and the possibility of horizontal gene transfer, necessitating rigorous long-term safety evaluations.

4.4 Frontier exploration

In response to the aforementioned challenges, recent research (primarily from the past five years) has focused on revolutionizing EV-based cardiovascular therapies through bioengineering and advanced technology integration. A key frontier involves enhancing EVs targeting specificity, cargo efficiency, and stability. Researchers now conjugate targeting ligands (e.g., peptides, antibodies) to direct EVs to damaged cardiac tissues, improving cardiomyocyte uptake while minimizing off-target effects. Notably, cardiac homing peptides show enhanced retention in infarcted myocardium in preclinical models (160). Parallel strategies employ rabies virus glycoprotein (RVG) antigens to achieve brain-directed EVs delivery, demonstrating the versatility of this targeting approach (185). Genetic and metabolic engineering further enable cargo customization with cardioprotective miRNAs/mRNAs to stimulate angiogenesis (202).

Advances in imaging represent another critical frontier. Stateof-the-art modalities (e.g., PET/MRI radiolabeling) track EVs biodistribution and clearance kinetics in vivo, while live-cell reporters enable real-time monitoring (203). These technologies elucidate EVs pharmacokinetics and validate targeting efficacy under physiological conditions. Innovative biomaterial integration overcomes systemic delivery limitations. Incorporating EVs into hydrogels or cardiac patches sustains localized release at myocardial sites, improving bioavailability and leveraging biomaterial mechanical properties for tissue repair. Concurrently, synthetic EVs mimetics offer tunable composition and scalability for future therapies. Multi-omics and machine learning herald a new era: Integrated proteomic/transcriptomic/lipidomic profiling deciphers EVs molecular signatures from cardiac cells, identifying novel CVDs biomarkers and therapeutic targets. This facilitates precision medicine applications where individualized EVs profiles guide patient-specific interventions.

5 Conclusion

Cardiovascular diseases (CVDs) represent a persistent global health challenge, demanding innovative approaches for diagnosis and therapy. This review consolidates the multifaceted roles of extracellular vesicles (EVs) in CVDs, highlighting their dual capacity to propagate pathological processes while orchestrating tissue repair. Critically, EVs derived from diverse sources—including circulating biofluids, stem cells, and plants which carry unique molecular signatures that enable their utility as dynamic biomarkers for early detection and risk stratification. The emergence of engineered EVs and plant-derived EVs further expands their therapeutic potential, offering targeted strategies for myocardial and cerebrovascular injury. However, clinical translation requires overcoming challenges in standardization of isolation protocols, scalable production, and precise tissue delivery. Future research must integrate multi-omics technologies with advanced bioengineering platforms to unlock the full potential of EVs in precision cardiology.

Author contributions

TL: Conceptualization, Investigation, Methodology, Supervision, Validation, Visualization, Writing – original draft,

Writing – review & editing. WW: Formal analysis, Methodology, Supervision, Validation, Writing – review & editing. ZQ: Visualization, Writing – original draft. YC: Investigation, Validation, Visualization, Writing – review & editing. KZ: Validation, Visualization, Writing – original draft. HL: Supervision, Validation, Writing – review & editing. JS: Conceptualization, Visualization, Writing – original draft. HZ: Supervision, Validation, Writing – review & editing.

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MRI

magnetic resonance imaging

Glossary

MSC mesenchymal stem cell acute myocardial infarction AMI MVmicrovesicles AS atherosclerosis MVB multivesicular bodies ASC adipose stem cell NF-κB nuclear factor kappa B CEC cerebral microvascular endothelial cell NMN nicotinamide mononucleotide CFs cardiac fibroblasts NSC neural stem cell CHD coronary heart disease OGD oxygen-glucose deprivation CI/R cerebral ischemia/reperfusion injury Ox-LDL oxidized low-density lipoprotein CM cardiomyocyte P-EXO platelet-derived exosomes CTP cardiac-targeting peptides PDEVs plant-derived EVs **CVDs** cardiovascular diseases PET positron emission tomography EC endothelial cell PS phosphatidylserine eNOS endothelial nitric oxide synthase plasma small extracellular vesicles psEVs **ESCRT** endosomal sorting complexes required for transport **ROCK** rho-associated coiled-coil containing kinases **ESTP** elastin-specific targeting peptide ROS reactive oxygen species EVs extracellular vesicles rabies virus glycoprotein peptide RVG **EXO** exosomes **SERS** surface-enhanced Raman scattering HF heart failure sEVs serum EVs HIF-1α hypoxia-inducible factor-1α ST-segment elevation myocardial infarction STEMI I/R ischemia-reperfusion injury TF tissue factor **ICH** intracerebral hemorrhage TGF-β tumor growth factor β **IHD** ischemic heart disease TNF-α tumor necrosis factor α lncRNA long non-coding RNA Treg regulatory T cell ΜI myocardial infarction **VEGF** vascular endothelial growth factor microRNA miR **VSMC** vascular smooth muscle cell MMP matrix metalloproteinase MPS mononuclear phagocyte system.