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Elucidating the mechanism of corneal epithelial cell repair: unraveling the impact of growth factors

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The repair mechanism for corneal epithelial cell injuries encompasses migration, proliferation, and differentiation of corneal epithelial cells, and extracellular matrix remodeling of the stromal structural integrity. Furthermore, it involves the consequential impact of corneal limbal stem cells (LSCs). In recent years, as our comprehension of the mediating mechanisms underlying corneal epithelial injury repair has advanced, it has become increasingly apparent that growth factors play a pivotal role in this intricate process. These growth factors actively contribute to the restoration of corneal epithelial injuries by orchestrating responses and facilitating specific interactions at targeted sites. This article systematically summarizes the role of growth factors in corneal epithelial cell injury repair by searching relevant literature in recent years, and explores the limitations of current literature search, providing a certain scientific basis for subsequent basic research and clinical applications.

KEYWORDS

cornea, corneal epithelial cells, growth factors, growth factor receptor, repair mechanism

1 Introduction

The cornea, in direct contact with the external environment, stands out as one of the body's tissues with the most dense innervation (1). Its structure is crucial for preserving the health and functionality of the ocular surface (Figure 1 shows the anatomical structure of the cornea). Damage to the corneal epithelium can result in corneal infections, ulcers, scar formation and vision loss ultimately. In recent years, factors such as surgical trauma, drug use, and infection have contributed to a rising number of patients with corneal epithelial injuries. As a result, the repair of corneal epithelial injuries has emerged not only as a prominent topic in basic scientific research but also as an urgent clinical problem that requires attention.

The corneal epithelium plays a crucial role in safeguarding the eye's barrier, stabilizing the tear film and maintaining the microenvironment of the ocular surface (2). Currently, the model governing corneal epithelial homeostasis relies on the XYZ hypothesis. According to this framework, the migration of limbal stem cells (LSCs) towards the central region of the cornea (X), combined with the vertical proliferation and differentiation of basal cells (Y), balances with the shedding of squamous cells (Z) from the epithelial surface. Healthy eyes are continuously bathed in tears containing growth factors, essential substances for maintaining



Histology of cornea. The cornea is structurally divided into five layers. The anterior epithelial layer comprises 5–7 layers of renewable epithelial cells. Behind the epithelial basement membrane lies the Bowman membrane, consisting of collagen fibers. The thickest layer, the stroma, primarily consists of keratocytes and collagen fibers, crucial for maintaining corneal transparency. Descemet, generated by the endothelium, is a transparent, elastic thin film with no distinct structure but possesses strong resistance. The endothelium is formed by a layer of hexagonal endothelial cells, incapable of regeneration.

the normal function of ocular surface tissue (3). Due to the unique anatomical location of the cornea, it is particularly susceptible to various injuries (4). When damaged, there is an upregulation of growth factors in tears, which target the cornea through relevant signaling pathways, thereby promoting corneal epithelial repair (5) and maintaining corneal epithelial homeostasis. Therefore, growth factors play a pivotal role in repairing corneal epithelial injuries and maintaining the normal microenvironment of the corneal epithelium.

In consideration of the aforementioned, this review aims to elucidate the roles played by various growth factors in the repair of corneal epithelial cells and comprehensively analyze their respective mechanisms of action. The overarching objective is to establish a robust scientific foundation that can serve as a springboard for subsequent basic research endeavors and clinical applications. The ensuing discussion will delve into the intricate interplay of growth factors within the context of repair mechanisms.

2 Growth factors involved In repair mechanisms

2.1 Epidermal growth factor

Epidermal growth factor (EGF) stands as one of the earliest identified single-chain peptides recognized for its ability to stimulate cell growth, playing a pivotal role in wound healing and maintaining tissue homeostasis by regulating cell survival, growth, motility and differentiation (6). In instances of corneal epithelial cell damage resulting from trauma, surgery or infection, EGF facilitates the migration and proliferation of corneal epithelial cells through the activation of its receptor EGFR/ErbB and subsequent binding. Consequently, this process promotes the effective repair of corneal epithelial injuries (7, 8). During the initial phases of corneal epithelial damage healing, EGFR1/ErbB1 tyrosine kinase instigates cellular signaling, activating downstream effectors such as the type III phosphoinositide 3-kinase (PI3K)—protein kinase B (Akt) axis and extracellular signal-regulated kinase (ERK). This orchestrated activation contributes significantly to the overall repair mechanism of corneal epithelial injuries (9). Furthermore, substance P also emerges as a noteworthy contributor to corneal epithelial injury repair, operating through the activation of EGFR and downstream signaling molecules, such as Akt (10). Abnormal activation of the EGFR-PI3K-AKT and ERK signaling pathways may result in increased cell apoptosis, decreased cell proliferation and delayed wound closure (11). The EGFR signaling pathway can further activate nuclear factor kappa-B (NF- κ B) and histone deacetylase 6 (HDAC6). NF- κ B, in turn, activates the transcription inhibitor CCCTC binding factor (CTCF) while downregulating the paired box gene 6 (PAX6), mediating the migration and proliferation of corneal epithelial cells. Simultaneously, HDAC6 promotes the migration of corneal epithelial cells and contributes to injury repair (12, 13).

There are four EGF receptors, with EGFR1 showing relatively high expression in corneal epithelial cells and demonstrating a reparative effect on the cornea during epithelial injury (14). EGFR2/ErbB2 and EGFR3/ErbB3 have also been confirmed to be expressed in the corneal epithelium, sharing a distribution pattern similar to EGFR1. Among them, the EGFR2/ErbB2 receptor enhances the corneal epithelial wound healing process by activating the ERK and PI3K signaling pathways (15). While the role of EGFR3/ErbB3 has not been fully elucidated, the existence of specific antibody inhibitors for EGFR3/ErbB3 has been confirmed. Utilizing these inhibitors and genetic techniques, studies have demonstrated that EGFR3/ErbB3 signaling can assist in the migration of corneal epithelial cells (16, 17). It's worth noting that EGFR4/ErbB4 is not expressed in the corneal epithelium (18).

Presently, seven EGFR ligands have been identified. In addition to EGF, six other endogenous ligands capable of binding to EGFR have been recognized, including heparin-binding EGF-like growth factor (HB-EGF), transforming growth factor- α (TGF- α), betacellulin (BTC), epiregulin, amphiregulin, and epigen. HB-EGF, integral in promoting growth and development, plays a crucial role, as evidenced by the fact that knockout mice perish shortly after birth (19).

Functioning as a soluble transmembrane protein, HB-EGF binds to an additional domain of negatively charged polysaccharides, thereby enhancing in vitro cell adhesion and promoting corneal epithelial injury repair (20). A notable discovery in the study indicates that HB-EGF exhibits prolonged cell attachment compared to EGF, resulting in a sustained impact on wound healing following brief therapy (21). TGF- α , a member of the epidermal growth factor family, is produced by both epidermal cells and macrophages. It plays a crucial role in the repair of corneal epithelial injuries by initiating multiple signaling cascade reactions upon binding with the EGFR (18, 22). The bidirectional interaction facilitated by TGF- α between corneal epithelial cells and mesenchymal cells assumes a pivotal morphological role in both corneal development and tissue repair. Any disruption in this intricate interplay can result in ocular lesions. Notably, TGF-a knockout mice exhibit significant ocular abnormalities, characterized by corneal epithelial thinning, inflammation, and edema (23, 24). TGF- α also promotes the proliferation of corneal epithelial and stromal cells (25). Additionally, TGF-α stimulates EGFR, facilitating the internalization and recycling of ligand-receptor complexes (22). Conversely, overexpression of TGF- α has been observed to induce corneal damage by activating EGFR in both corneal epithelium and stroma. This pathological manifestation is evident through a reduction in the number of corneal epithelial cell layers, corneal epithelial degeneration, conjunctivalization of the cornea, inhibition of the expression of the corneal pigment protein Kera, and a marked decrease in fibrocollagen types I and V collagen. Simultaneously, TGF-α overexpression can lead to corneal opacity by upregulating α -SMA and Wnt5a, while downregulating Col1a1, Col1a2, and Col5a1 (25-28). Some in vitro analysis of BTC indicates that BTC can expedite corneal epithelial injury repair and may even possess advantages over EGF in promoting corneal epithelial injury repair (18). LSCs, primarily located at the corneal-scleral junction, possess lifelong self-renewal capabilities and can produce transient amplifying cells (TACs). During corneal epithelial injury repair, TACs migrate towards the corneal center, proliferate, and differentiate into corneal epithelial cells, thereby promoting the healing of corneal epithelial wounds (29). Research has shown that treating injured mouse eyes with BTC results in significant increases in the expression of putative stem cell markers, such as DNp63a, ABCB5 and CK14. This suggests that BTC accelerates corneal LSCs proliferation and enhances mouse corneal epithelial repair by phosphorylating erk1/2 (30, 31). Despite the efficacy of various EGFR ligands in in vitro settings, in vivo wound healing is uniquely facilitated by EGF among the seven mentioned ligands. EGF also stands out as the sole ligand in human tears with an EGFR concentration closely aligned with the ligand Kd (18). Furthermore, EGFR can be reactivated through various effectors, such as phospholipase D (PLD) and extracellular ATP, to foster the migration and proliferation of cells during the wound healing process (32). Transient receptor potential (TRP) non-selective cation channels constitute a superfamily, which contains 28 different genes, and widely distributed in corneal epithelial cells and endothelial cells, its expression in the corneal epithelial layer contributes to the maintenance of corneal transparency and barrier function of the corneal epithelium. Research has shown that TRPV1 stimulation also induces increases in the proliferation and migration of corneal epithelial cells and the release of IL-6 and IL-8, and reduces the formation and of corneal neovascularization (CNV) and scar through

transactivation of the EGFR. Meanwhile, TRPC4 stimulates corneal epithelial proliferation and migration by transactivation of the EGFR. TRPV in corneal epithelium can also promote homeostasis under thermal stimulation (33, 34).

While EGF holds the potential to stimulate the migration and proliferation of corneal epithelial cells, caution is warranted, as excessively increasing the intensity and duration of EGF may not yield positive effects. An experiment assessing EGF's effectiveness has revealed potential harm from continuous daily injections in rats (35). Furthermore, injecting recombinant EGF into the cornea post-corneal epithelial cell injury can lead to CNV (36). Elevated tear EGF levels are also associated with meibomian duct hypertrophy, contributing to meibomian gland hyperplasia (37). These findings underscore the need for caution when employing exogenous EGF in treating corneal injuries. Additionally, EGFR activity is a critical determinant in maintaining corneal epithelial homeostasis and plays a pivotal role in restoring damaged corneal epithelial cells. Despite the inherent regulatory mechanisms preventing sustained EGFR signal transduction, a deeper understanding of the molecular mechanisms governing EGFR signaling holds promise for developing new methods to overcome these regulatory barriers and enhance the efficacy of EGF (Figure 2 shows the EGF signaling pathway).

2.2 Hepatocyte growth factor

Hepatocyte growth factor (HGF) is a growth factor originating from fibroblasts, predominantly produced by mesenchymal cells, and expressed in various cell types, including corneal epithelial cells, keratocytes and endothelial cells. Its mode of action is paracrine, exerting its effects on adjacent cells (38). Structurally, HGF consists of α - and β -chains and serves as a mitogen and motility factor. In the cornea, HGF plays a significant role by binding to its receptor c-met and primarily participating in the proliferation, mitosis, and morphogenesis of corneal epithelial cells (39–41).

When the corneal epithelium undergoes damage, the expression of HGF in corneal epithelial cells and keratocytes is upregulated. This upregulation activates the signal mediators phosphatidylinositol of PI3K/Akt, phosphoprotein 70 ribosomal protein S6 kinase (p70s6K), and ERK. Consequently, it controls the cell cycle, promoting cell division and proliferation of corneal epithelial cells by triggering the activity of NF-KB. Simultaneously, it reverses the anti-proliferative effect of pro-inflammatory cytokines interleukin-1ß (IL-1ß) and TNF- α on these cells in the inflammatory environment, mediating corneal epithelial injury repair (42-46). HGF also exhibits wound repair effects by inhibiting the inflammatory response of corneal epithelial cells. Studies have demonstrated that HGF can inhibit the activation of immune cells and the expression of inflammatory factors. It further suppresses the expression of TNF- α , monocyte chemotactic protein-1 (MCP-1), and IL-6 in the macrophage system in vitro. Additionally, it promotes the production of the anti-inflammatory cytokine IL-10 in bone marrow-derived macrophages and dendritic cells stimulated by lipopolysaccharide (LPS) (43, 47-49). Evidence supports that HGF significantly inhibits cell apoptosis, temporarily downregulates the expression of cell cycle inhibitors in corneal epithelial cells, and upregulates cyclin and cyclin-dependent kinases. It also influences tumor suppressor proteins Rb and p53, which regulate cell cycle and apoptosis. Through these mechanisms, HGF



EGFR, stimulating pathways of EGF. This ingure intestrates the signal transduction mechanism of EGF in corneal epithetial injury repair. EGF binding activate EGFR, stimulating various signaling pathways like PLC γ , Ras-GAP, Grb2, and Shc. These pathways collectively contribute to the reparative effects on corneal epithetial injuries.

actively participates in corneal epithelial injury repair (50, 51). HGF also has the capability to penetrate through LPS-induced corneal opacity, promoting recovery, diminishing corneal fibrosis, restoring normal corneal tissue structure, and reestablishing immune quiescence after keratitis (52). In cases of diabetes-related corneal epithelial damage, HGF exhibits a reparative effect by restoring the level of c-met in the cornea of diabetic patients through downstream activation of p38 mitogen-activated protein kinase (MAPK) and the production of several putative stem cell markers. Importantly, this positive effect is observed in cultured corneas, regardless of whether gene therapy is applied to the entire corneal epithelial cells or only to the corneal edge area containing stem cells (53). Research indicates that silencing the HGF gene inhibits corneal epithelial proliferation and UVR-induced CNV. Additionally, HGF contributes to the upregulation of vascular endothelial growth factor (VEGF) and plays a role in angiogenesis regulation. These findings open up new avenues for exploring treatment strategies for CNV (54, 55).

2.3 Insulin-like growth factor

Insulin-like growth factor (IGF) belongs to the multifunctional cell proliferation regulatory factor, representing a group of peptide

substances capable of promoting growth. Its secretory cells are widely distributed in various tissues, including the liver, kidneys, heart and eyes of the human body. The IGF family comprises two peptide ligands (IGF-1 and IGF-2), three receptors, and six binding proteins, collectively maintaining tissue homeostasis by regulating metabolism and/or mitotic pathways at the level of all corneal cells (56).

IGF-1, a multifunctional cytokine with broad biological activity, holds considerable promise for applications in corneal epithelial injury repair. By binding to the insulin-like growth factor 1 receptor (IGF-1R), IGF-1 actively maintains and regulates corneal epithelial cell growth, proliferation, differentiation, maturation, migration, regeneration, and energy metabolism. It promotes corneal epithelial cell proliferation through the activation of the hybrid of IGF-1R and insulin receptor (INSR), leading to subsequent Akt phosphorylation. Additionally, IGF-1 mediates corneal epithelial cell migration through the PI3K/AKT pathway. Furthermore, IGF-1 promotes the expression of IGF receptors in corneal limbal cells, stimulating LSCs to differentiate into corneal epithelial cells (57, 58). Beyond its role in cell proliferation and migration, IGF-1 serves as a crucial neurotrophic factor facilitating the regeneration and restoration of nerves following peripheral nerve damage in the cornea (59). In combination with substance P, IGF-1 demonstrates a synergistic effect in promoting corneal epithelial injury repair. The co-application accelerates the ex

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vivo migration of corneal epithelial cells in the injured corneal stroma. Mediated by the interaction between substance P and tachykinin receptors, it enhances the adhesion of corneal epithelium to fibronectin (FN) and type IV collagen, thereby augmenting the protective role of the corneal epithelium through the stimulation of wound healing (60-64). The low levels of IGF-1 in tears, particularly the reduced proportion of IGF-1 and IGF binding protein 3 (IGFBP-3) in tears of diabetic patients, have been associated with decreased proliferation of corneal epithelial cells and delayed wound repair. This change inhibits the capacity of IGF-1 to induce IGF-1R or hybrid R phosphorylation (65, 66). Research has demonstrated that mRNA of adipose-derived stem cells (ADSCs) modified with IGF-1 exhibits stronger cell proliferation and migration abilities, promoting wound repair, morphological and functional recovery, corneal nerve regeneration, and maintenance of corneal homeostasis after acute alkali burns. Importantly, it can prevent the generation of CNV and corneal lymphatic vessels, highlighting the crucial role of IGF-1 in the repair of corneal epithelial injury (60). However, the application of IGF-1 protein to the cornea in the form of eye drops faces limitations, including a limited duration of effect, elevated attrition rates, and the need for repeated administration. Further research is expected to explore new carrier forms that can overcome these shortcomings and enhance the effectiveness of IGF-1 in corneal injury repair. The expression of IGF-2 and its receptors significantly increases after corneal epithelial cell injury, promoting the transformation of LSCs in the basal layer of the cornea into corneal epithelial cells and subsequently supporting corneal epithelial cell repair (67). Additionally, both IGF-1 and IGF-2 play roles in promoting the proliferation of keratocytes and collagen synthesis (68).

IGFBP primarily exists in the aqueous humor and vitreous body, exerting unique, cell- and tissue-dependent effects through interactions with the IGF family via binding (69-71). The primary function of IGFBP is to bind to IGF-1, extending its half-life in circulation and preventing IGF-1R activation induced by IGF-1 (72, 73). IGFBP-2 and IGFBP-3 play pivotal roles in corneal tissue homeostasis, particularly in regulating the growth of corneal epithelial cells and the localization of intracellular receptors (74, 75). The mutual regulation between IGFBP-3 and IGF-1R maintains corneal epithelial homeostasis. Previous studies have shown that IGFBP-3 is essential for inducing the transport of IGF-1R, and the absence of IGF-1R will downregulate IGFBP-3 in turn (76). During conditions such as hypoxia and hyperglycemia, the secretion of IGFBP-3 increases. For instance, the level of IGFBP-3 in the tears of diabetic patients rises, suggesting its potential role in regulating eye homeostasis in diabetic patients and indicating therapeutic potential in ocular surface diseases associated with diabetes (66).

Moreover, IGF and insulin share a close relationship, with the former mediating the action of insulin to promote the growth of corneal epithelial cells. This suggests a potential collaborative repair effect between the two, offering a promising avenue for future research exploration (77, 78).

2.4 Neurogenic growth factor

Neurogenic growth factor (NGF) belongs to the family of neurotrophic factors, exhibiting dual biological functions of neuronal nourishment and promoting synaptic growth (79). In the context of the cornea, signals mediated by NGF propagate through the highaffinity receptor tropomyosin receptor kinase A (TrkA) and the low-affinity non-selective transmembrane glycoprotein receptor p75NTR. When combined with NGF, TrkA activates Ras MAPK, ERK, phospholipase C- γ (PLC- γ), and PI3K. This activation includes stimulating D-type cell cycle regulatory proteins through PI3K/Akt and MAPK/ERK, subsequently promoting corneal epithelial cell cycle progression. Simultaneously, p75NTR activates the c-Jun kinase and NF-kB signaling pathway, exhibiting a protective effect on corneal epithelial cells by inhibiting the inflammatory signaling pathway of NF-kB (80–82).

Research has demonstrated that NGF participates in the repair process of corneal epithelial and stromal damage by upregulating matrix metalloproteinase-9 (MMP-9) and cleaving integrins β4 to stimulate the migration of corneal epithelial cells, promotes the differentiation of keratocytes into myofibroblasts, and reduces the formation of corneal haze (83-85). Moreover, NGF induces the differentiation of goblet cells and the production of mucin through receptors expressed in the lacrimal gland and neural reflexes, thereby contributing to the maintenance of corneal epithelial function (86). In addition to its role in cellular functions, NGF regulates immune function through Toll-like receptors (TLRs) in corneal physiology and pathology, playing a crucial role in maintaining corneal homeostasis both in vivo and in vitro settings (87). NGF has also been identified as a key promoter for the proliferation of LSCs, the formation of colonies in LSCs, and the maintenance of the LSC phenotype (79). For patients with corneal ulcers, local application of NGF eye drops has been shown to improve the speed of corneal epithelial repair and the sensitivity of the cornea. In cases of herpes simplex keratitis (HSK), endogenous NGF, akin to acyclovir, significantly improves the condition and inhibits recurrence. Clinical studies indicate that eye drops containing NGF can induce complete healing in HSK patients resistant to acyclovir (88, 89). Treatment with recombinant human NGF (rhNGF) has proven effective in enhancing corneal perception in patients with neurotrophic keratitis (NK) by increasing the density and number of nerve fibers in the basal layer of the corneal epithelium. It also promotes the healing of persistent corneal epithelial defects and ulcers. Furthermore, rhNGF provides lubrication and natural protection against pathogen damage to the corneal epithelium by promoting tear secretion from the lacrimal gland. RhNGF has received approval as a primary therapeutic drug for NK (87, 88). Additionally, NGF exhibits the ability to inhibit oxidative damage caused by hyperosmotic stress or high glucose levels. This finding suggests its potential therapeutic effect on conditions such as dry eye syndrome and diabetic keratopathy (DK) (90, 91).

2.5 TGF-β

TGF is a protein composed of amino acids in the cytoplasm, belonging to the family of peptide growth factors. It includes two main types: TGF- α and TGF- β (92). TGF- α has been described in the EGF section. TGF- β is a multifunctional growth factor, further divided into three subtypes: TGF- β 1, TGF- β 2 and TGF- β 3. All three subtypes and their receptors are expressed in corneal epithelium and keratocytes (93). TGF- β assumes a pivotal role in orchestrating and coordinating the response to corneal injury repair, exerting influence over various facets such as the proliferation, motility, and differentiation of corneal

epithelial cells. Moreover, TGF-β modulates the activity and apoptosis of keratocytes, as well as the development of myofibroblasts (94). By stimulating the migration of corneal epithelial cells through integrin β 1, TGF- β enhances the fluidity of these cells (95). Notably, the conditional ablation of its type II receptor has been found to impede the repair of corneal epithelial wounds and the activation of p38 MAPK, thereby hindering the migration of corneal epithelial cells (96). Furthermore, TGF-\u03b32 has been substantiated to expedite the repair of corneal epithelial wounds in rabbits, augmenting barrier integrity by promoting cell adhesion to substrates and enhancing the functionality of corneal endothelial cells (CECs) (97). Tgfbr-2 also plays a crucial role in maintaining corneal stromal homeostasis, as studies have demonstrated that Tgfbr-2 knockout mice display significant corneal thinning and a potential for corneal ectasia (98). TGF-β3 exhibits the capability to mitigate interstitial scars induced by the activity of TGF-\u00b31 and TGF-\u00b32. Moreover, it demonstrates potential therapeutic effects in addressing corneal and skin wounds in diabetic patients, acting through the PI3K-Akt and SMAD signaling pathways, along with their target genes (99). Additionally, if the Bowman layer is damaged, corneal cells are highly susceptible to exposure to TGF-β. In such cases, TGF-β promotes damage repair through various mechanisms (100). Despite its essential role in corneal epithelial injury repair, TGF-B also has negative effects on the cornea. For instance, it can promote the aging of corneal epithelial cells through the NF-kB signaling pathway. This aging process can be alleviated by inhibiting the NF- κ B signaling pathway (101). TGF- β is implicated in the pathogenesis of various eye diseases, including pterygium, vernal keratoconjunctivitis (VKC), atopic keratoconjunctivitis (AKC), and graft-versus-host disease (GVHD). Elevated levels of TGF- β are observed in the corneas of individuals with these diseases (102). Additionally, TGF- β regulates the transformation of corneal epithelial cells and corneal fibroblasts into myofibroblasts, and the high expression of α-SMA and F-actin in myofibroblasts can lead to the loss of corneal transparency and corresponding corneal haze (103). Moreover, TGF-β1 and TGF-β2 can prevent corneal epithelial cells from proliferating in vitro (104).

2.6 Platelet-derived growth factor

Platelet-derived growth factor (PDGF), secreted by epithelial cells, endothelial cells and inflammatory cells, serves as a potent mitogenic factor, existing in diverse isoforms, namely PDGF-AA, PDGF-BB, PDGF-CC, PDGF-DD, and PDGF-AB (105). Featuring both α and β types of receptors, PDGF exerts its cellular effects by inducing the complex formation of α -tyrosine kinase receptors and β -tyrosine kinase receptors. This induction, in turn, triggers processes such as cell growth, chemotaxis, actin recombination and protection against apoptosis. Analogous to TGF-β, PDGF assumes a pivotal role in regulating and coordinating the response to corneal wound repair. It influences the proliferation, motility and differentiation of corneal epithelial cells, while also modulating the activity and apoptosis of keratocytes and contributing to the development of myofibroblasts (94). Corneal epithelial cells express PDGF AA, PDGF BB and PDGF AB, which regulate the migration and proliferation of keratocytes. In the presence of FN, these isoforms can enhance the migration of corneal epithelial cells (106-108). Research indicates that PDGF-AB and PDGF-BB promote the migration of corneal fibroblasts in vitro, leading to an increase in the concentration of cytosolic free Ca²⁺. PDGF-BB also significantly stimulates DNA synthesis in bovine corneal endothelial cells (BCEC) and human corneal fibroblasts (HCF) in a dose-dependent manner (108–110). Moreover, under high fibroblast density, PDGF isomers act as mitogens for interstitial fibroblasts during wound healing, conversely, at low cell density, PDGF-AA and PDGF-AB can prevent cell loss during the corneal homeostasis process (111). The secretion of PDGF is activated during corneal trauma, infection, or inflammation, providing significant stimulation for tissue repair. However, hyperstimulation can have negative effects. For instance, PDGF- α hyperstimulation can promote the proliferation and migration of lens epithelial cells, leading to epithelial-mesenchymal transition (EMT) (112, 113).

2.7 Fibroblast growth factor

Fibroblast growth factor (FGF) is secreted by the hypothalamus and pituitary gland, serving as a broad-spectrum mitogen, currently, at least 23 FGF families have been identified, stimulating or maintaining specific cellular functions required for tissue metabolism, homeostasis and development through signaling axes mediated by their receptors (114). Corneal epithelial changes, accompanied by decreased vision and dry eye symptoms, have been observed after treatment with inhibitors of the FGF receptor (FGFR), indirectly indicating FGF's involvement in corneal epithelial homeostasis (115). Basic FGF (b-FGF/FGF2), approved for the treatment of corneal damage, accelerates the repair of corneal epithelial cell damage and reduces keratitis by promoting the proliferation, differentiation and migration of corneal epithelial cells (116, 117). In experiments involving FGFR2 knockout mice, observations reveal localized central corneal thinning, along with the loss of collagen fibers and apoptosis of keratocytes (118). In addition, the signal transduction of FGFR2b promotes corneal epithelial injury repair, studies have found that FGFR2b knockout mice exhibit reduced proliferation of corneal epithelial cells, as well as loss of lacrimal gland and meibomian gland. FGFR2b is also necessary for the development of submandibular glands (119). FGF-10 plays a crucial role in the development of the cornea, morphogenesis and growth of the lens and induction and branching of the lacrimal gland and meibomian gland. Research has shown that FGF-10 is essential for the development of lacrimal glands in humans and mice (120, 121). FGF-10 can upregulate the expression of mucin in conjunctival epithelial cells, protecting the ocular surface in a dry eye model of rabbits and controlling the migration of epithelial cells during the process of embryonic eyelid closure (122, 123). Additionally, FGF-10 is associated with adult tissue homeostasis and the function of stem cells (124). In an experimental study of DK, it has been found that rhFGF-21 can improve the vitality and migration of human corneal epithelial cells, promote the healing of corneal wounds and the production of tears, and improve corneal edema. RhFGF-21 significantly reduces the expression of pro-inflammatory cytokines such as TNF-α and MMPs in corneal epithelial cells, while increasing the level of anti-inflammatory molecules IL-10 and SOD-1. RhFGF-21 also inhibits excessive production of reactive oxygen species (ROS) and alleviates oxidative stress induced by hyperglycemia in corneal epithelial cells. Therefore, the application of drugs containing FGF-21 may be a potential treatment method for DK (125, 126).

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2.8 Keratinocyte growth factor

Keratinocyte growth factor (KGF) belongs to the FGF family, officially known as FGF-7. It is produced by mesenchymal cells and acts on adjacent cells in a paracrine manner. KGF promotes the repair of corneal epithelial wounds through the signaling cascade of MAPK and PI3K/p70 S6 in corneal epithelial cells. It also inhibits the destruction of the barrier function caused by hypoxia in corneal epithelial cells by activating ERK (46, 127, 128). In vitro experiments have shown that KGF can protect cells from apoptosis for an extended period, with the final percentage of apoptosis in cells treated with KGF being only 10% (129). Similar to HGF, KGF has the capacity to inhibit UVR-induced corneal epithelial proliferation. Through gene silencing, it downregulates the expression of VEGF and its receptors, consequently mitigating CNV (48, 49). Additionally, KGF-2 exhibits certain effects in re-epithelialization, accelerating migration, reducing scar formation and edema. It is considered superior to b-FGF and holds potential as a new drug for treating corneal injuries (116). KGF also can promote the migration of LSCs, thereby promoting the repair of corneal epithelial damage (36).

2.9 Opioid growth factor

Opioid growth factor (OGF) is an endogenous peptide found together with its receptor OGFr in or on the basal layer of many species' corneas, when combined, OGF regulates DNA synthesis in corneal epithelial cells and influences cell migration (130). Experimental evidence suggests that OGF inhibits cell overgrowth by upregulating cyclin-dependent inhibitory kinases p16 and p21, contributing to the maintenance of corneal epithelial homeostasis (131). Studies conducted on patients and rats with diabetes have indicated elevated levels of OGF and OGFr in serum and corneal epithelium. This elevation has been associated with ocular surface complications, including dry eyes, abnormal sensitivity of the corneal surface, and delayed corneal epithelial repair. In rats, an OGF-OGFr axis is present in the corneal limbus, and its dysregulation in hyperglycemia impacts the morphology of the corneal limbus, exacerbating diabetes-related complications on the corneal surface. Local application of the opioid antagonist naltrexone (NTX) has demonstrated improvement in this situation (132, 133). NTX disrupts the OGF-OGFr interaction, resulting in increased DNA synthesis in epithelial cells of the peripheral cornea and limbus corneae, as well as the proliferation of fibroblast cells. The latter plays a crucial role in corneal wound healing (134, 135). Additionally, the application of NTX significantly promotes corneal re-epithelialization and increases tear production (136-138). In summary, the use of eye drops containing opioid antagonists, such as NTX, holds promise as a novel therapy for treating wound repair disorders of the corneal epithelium.

2.10 VEGF

VEGF, recognized as a highly specific mitogen promoting endothelial cell growth, is also referred to as vascular permeability factor (VPF) due to its ability to significantly enhance vascular permeability (139). The VEGF family comprises seven subtypes: VEGF-A, VEGF-B, VEGF-C, VEGF-D, VEGF-E, VEGF-F, and placental growth factor (PLGF). Additionally, there are three receptors, VEGFR-1, VEGFR-2, and VEGFR-3. Among these, VEGF-A stands out for its profound ability to promote angiogenesis and is the most prevalent subtype in the eyes (140). The VEGF family and its receptors are expressed in the corneal epithelium (4). Under normal circumstances, a delicate balance is maintained between ocular angiogenic factors and anti-angiogenic factors to prevent pathological CNV production. However, factors such as wearing contact lenses, inflammation, and infection can disrupt this balance and lead to CNV (141). Research has highlighted the crucial role of VEGF in the pathogenesis of CNV, making the use of anti-VEGF drugs a feasible therapeutic approach (142, 143). Additionally, VEGF is implicated in non-angiogenic functions, such as neuroprotection and serving as a nutritional factor for corneal nerves (144). VEGF also plays a role in wound repair; studies have shown that VEGF accelerates corneal epithelial wound healing by stimulating corneal nerve regeneration (145). VEGF may be linked to the pathogenesis of pterygium, with higher expression detected in pterygium compared to normal tissue (146). Pigment epithelial-derived factors (PEDFs) are closely related to VEGF, sharing anti-angiogenic functions and a protective role for corneal nerves (147). Both factors exhibit synergistic therapeutic effects in certain diseases. However, in pterygium, a decrease in PEDF expression has been observed (146). Moreover, PEDF has been found to promote the self-renewal and migration of LSCs, thereby facilitating corneal epithelial repair (148). (The expression of growth factors in the corneal epithelium is depicted in Figure 3, and the biological effects of corneal epithelial growth factor receptors are summarized in Table 1).

3 Discussion and outlook

Eye injuries often involve damage to the corneal epithelial layer, leading to symptoms such as eye pain, bleeding, ulcers, and vision loss, significantly impacting quality of life (151, 152). The repair of corneal epithelial injuries has emerged as a prominent research focus, with growing recognition of the crucial role played by growth factors in this process. These growth factors contribute to the wound healing of corneal epithelium through intricate mechanisms.

Despite the significant role of growth factors in corneal epithelial injury repair, there are existing limitations. The current understanding of the signal transduction pathways of various growth factors is not yet comprehensive, and their full potential as a treatment method for corneal epithelial injury remains to be realized. Some growth factors, like TGF, have stringent usage and dosage guidelines in corneal epithelium treatment-only within a safe usage range can they effectively repair the corneal epithelium. This presents a crucial and challenging aspect in utilizing growth factors for corneal epithelium treatment. For targeted repair effects of growth factors, many studies focus on single targets or signal pathways. The comprehensive repair mechanisms involving different growth factors still require further exploration. Additionally, some growth factors are primarily limited to basic research, and their potential for improving corneal epithelial cells in clinical practice needs further validation. Growth factors can be categorized into endogenous and exogenous types, with exogenous growth factors often utilized in experimental studies involving mice. However, when it comes to the role of growth factors in corneal epithelial injury repair, there is a noticeable gap in research and discussion regarding whether the mechanisms differ between the two types. Furthermore, in the case of the growth factor VEGF, exploring more suitable drug carriers could potentially enhance its therapeutic



efficacy. This avenue of research could lead to the development of more effective delivery systems for VEGF, optimizing its impact on corneal epithelial repair.

Future research endeavors should leverage current multi-omics techniques to explore and study the signaling mechanisms and synergistic effects of different growth factors, aiming to enhance their roles in corneal epithelial injury repair. Simultaneously, through in-depth basic and clinical research, optimizing the dosage and understanding side effects of growth factors can contribute to revealing their basic mechanisms and optimal usage methods. This foundational work will pave the way for the development of new treatment methods involving growth factors. Further research and exploration are essential to determine potential differences in the roles of endogenous and exogenous growth factors in corneal epithelial repair, a facet not addressed in the current study. Additionally, investigating ways to enhance exogenous growth factors and identifying more suitable drug carriers are critical for optimizing the use of growth factors in the future.

Author contributions

JG: Writing – original draft. GD: Writing – review & editing. ZH: Writing – review & editing. YL: Writing – review & editing. AD: Writing – review & editing. CZ: Writing – review & editing.

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TABLE 1 Biological effects of corneal epithelial growth factor receptor.

Factor	Receptor	Effects on corneal wound healing	References
EGF	EGFR	 Promotes migration and proliferation of epithelial cells EGFR can be reactivated through various effectors 	(7, 8, 12, 13, 17, 32)
HB-EGF			(19)
IID-EGF			(19)
BTC		 Accelerates corneal epithelial injury repair Accelerates the proliferation of LSCs 	(18, 30)
		Promotes the proliferation of corneal epithelial and stromal cells	()
TGF-α		Facilitates the internalization and recycling of ligand-receptor complexes	(22, 25)
HGF		 Participates in the mitosis and morphogenesis of corneal epithelial cells Promotes cell division and proliferation of corneal epithelial cells by controlling the cell cycle Reverses the anti-proliferative effect of pro-inflammatory cytokines in the inflammatory environment 	
	c-Met	 Inhibits the activation of immune cells and the expression of inflammatory factors Promotes the production of anti-inflammatory cytokines Suppresses apoptosis of corneal epithelial cells Promotes corneal opacity recovery, reduces corneal fibrosis, normalize corneal tissue structure and 	(39–55)
		 reestablish immune quiescence post-keratitis Restores the level of c-met in the cornea of diabetic patients Inhibits UVR-induced corneal epithelial proliferation and CNV by HGF gene silencing Participates in angiogenesis 	
IGF/INS	IGF-1R、IGF- 2R、INSR、Hybrid-R	 Regulates metabolism and/or mitotic pathways Promotes the migration and proliferation of epithelial cells Facilitates the regeneration and restoration of corneal nerves Accelerates the <i>ex vivo</i> migration of corneal epithelial cells in the injured corneal stroma Enhances the adhesion of corneal epithelium to FN and type IV collagen Promotes the transformation of LSCs in the basal layer of the cornea into corneal epithelial cells Promotes the proliferation of keratocytes and collagen synthesis Plays pivotal roles in corneal tissue homeostasis 	(56–64, 67, 68, 74, 75)
NGF	TrkA, p75NTR	 Stimulates the migration of corneal epithelial cells, promotes the differentiation of keratocytes into myofibroblasts and reduces the formation of corneal haze Induces the differentiation of goblet cells and the production of mucin regulates immune function Be identified as a key promoter for the proliferation of LSCs, the formation of colonies in LSCs and the maintenance of the LSCs phenotype Improves the speed of corneal epithelial repair and the sensitivity of the cornea of patients with corneal ulcers, significantly improves the condition and inhibits recurrence of HSK Increases the density and number of nerve fibers in the basal layer of the corneal epithelium, promotes tear secretion Inhibits oxidative damage caused by hyperosmotic stress or high glucose levels 	(79, 83–91, 149, 150)
TGF-β	Tgfbr-1, Tgfbr-2, Tgfbr-3	 Modulates the activity and apoptosis of keratocytes, modulates the development of myofibroblasts Stimulates the migration of corneal epithelial cells Augments barrier integrity by promoting cell adhesion to substrates and enhancing the functionality of corneal endothelial cells (CECs) Plays an important role in corneal stromal homeostasis Reduces interstitial scars promotes damage repair of Bowman layer Promotes the aging of corneal epithelial cells Be implicated in the pathogenesis of various eye diseases, such as pterygium Regulates the transformation of corneal epithelial cells and corneal fibroblasts into myofibroblasts Prevents corneal epithelial cells from proliferating <i>in vitro</i> 	(94–103)
(PDGF)	PDGF-αR, PDGF-βR	 Promotes the proliferation, motility and differentiation of corneal epithelial cells Modulates the activity and apoptosis of keratocytes and contributes to the development of myofibroblasts Stimulates DNA synthesis in BCEC and HCF Acts as mitogens for interstitial fibroblasts during wound healing Prevents cell loss during the corneal homeostasis process 	(94, 105–110)

(Continued)

TABLE 1 (Continued)

Factor	Receptor	Effects on corneal wound healing	References
FGF	FGFR	 Participates in corneal epithelial homeostasis Promotes the proliferation, differentiation and migration of corneal epithelial cells Promotes the regeneration of the lacrimal gland, submandibular gland and meibomian gland Promotes the development of the cornea and morphogenesis and growth of the lens Be associated with adult tissue homeostasis and the function of stem cells Improves the vitality of human corneal epithelial cells Improves corneal edema Reduces the expression of pro-inflammatory cytokines and increases the level of anti-inflammatory molecules Suppresses excessive production of ROS and alleviates oxidative stress 	(114–120, 123–125)
(KGF)	KGFR	 Inhibits the destruction of the barrier function caused by hypoxia in corneal epithelial cells Suppresses apoptosis of corneal epithelial cells Inhibits UVR-induced corneal epithelial proliferation and CNV by HGF gene silencing Enhances re-epithelialization rates Promotes the migration and proliferation of epithelial cells Reduces corneal scars and edema Promotes the proliferation of LSCs 	(36, 48, 49, 115, 127, 128)
OGF	OGFr	 Regulates DNA synthesis in corneal epithelial cells and influences the migration of corneal cells Inhibits the overgrowth of corneal cells and maintains of corneal epithelial homeostasis 	(129, 130)
VEGF	VEGFR-1, VEGFR-2, VEGFR-3	 Promotes the growth of endothelial cell Promote corneal angiogenesis Protecting corneal nerves Be related to the pathogenesis of pterygium 	(138, 139, 143, 145)

EGF, epidermal growth factor; EGFR, epidermal growth factor receptor; HB-EGF, heparin-binding EGF-like growth factor; BTC, betacellulin; TGF-α, transforming growth factor-α; HGF, hepatocyte growth factor; c-Met, hepatocyte growth factor receptor; IGF, insulin-like growth factor; INSA, insulin-like growth factor: 1 receptor; IGF-2R, insulin-like growth factor-2 receptor; INSR, insulin receptor; Hybrid-R, IGF-1R and INSR hybrid; NGF, nerve growth factor; btt, tropomyosin receptor kinase A; p75NTR, low-affinity non-selective transmembrane glycoprotein receptor; TGF-β, transforming growth factor-β; Tgfbr-1, transforming growth factor; PDGF-α, platelet derived growth factor; PDGF-α, platelet derived growth factor; PDGF-α, platelet derived growth factor; GGF, β, platelet derived growth factor; CGF, β, FGF, fibroblast growth factor; FGFR, fibroblast growth factor receptor. (GFF, opioid growth factor; CGF, opioid growth factor; CGF, opioid growth factor; CGF, opioid growth factor; CGF, opioid growth factor receptor.

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Conflict of interest

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

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Glossary

EGFipidermal growth factorEGFN/EbBipidermal growth factor receptor 1EGFN/EbBipidermal growth factor receptor 1PI3KOrole Ikinase BEARNolcel racional spain s		
InterfactorFileFilePI3KFilePI3KType III phosphoinositid 3-kinaseAktProtein kinase BERKExtracellular signal-regulated kinaseNF-kBNuclear factor kappa-BHDAC6Histone deacetylase 6CTCFCCCTC binding factorPAX6Paired box gene 6EGR2/ErbB2Epidermal growth factor receptor 3EGR4/ErbB4Epidermal growth factor receptor 4HB-EGFEpidermal growth factor receptor 4HB-EGFEpidermal growth factor-αTGF-4Tansforming growth factor-αFGTEdtaellulinPLDOroneal neovascularizationFGFHepato-tip for biosponal protein S6 kinaseFL7Nonocyte chemotactic protein-1L1βHerelewin-1βMCP-1Niogen-activated protein kinaseFGFIssulin-like growth factorFGFScaluar endothelial growth factorIGFBGirbinding proteinIGFBIssulin-like growth factorIGFBGirbinding proteinIGFAScaluar endothelial growth factorIGFBGirbinding protein-3ADSCAAipose-derived stem cellsIGFBP-3Girbinding protein-3ADSCAAipose-derived stem cellsIGFBP-3Girbinding protein-3ADSCAAipose-derived stem cellsIGFBP-3Girbinding protein-3ADSCAAipose-derived stem cellsIGFBP-3Girbinding protein-3ADSCAGirbonding scaluar-3IG	EGF	Epidermal growth factor
InterfactPI3KType III phosphoinositid 3-kinaseAktProtein kinase BERKExtracellular signal-regulated kinaseNF-kBNuclear factor kappa-BHDAC6Histone deacetylase 6CTCFCCCTC binding factorPAX6Paired box gene 6EGFR4/ErbB2Epidermal growth factor receptor 2EGFR4/ErbB3Epidermal growth factor receptor 4HB-EGFHepaten-binding EGF-like growth factorTGF-aTansforming growth factor aTGF-aTeatsforming growth factorPTCBetacellulinPLDPhospholipase DCNVCorneal neovascularizationHGFHepatocyte growth factorP70s6KPhosphoprotein 70 ribosomal protein S6 kinaseIL-1βIterleukin-1βMCP-1Monocyte chemotactic protein-1MAFKMitogen-activated protein kinaseVEGFVacular endothelial growth factorIGFBPIpoineline growth factorIGFBPIpoineline growth factorIGFBPIpoineline growth factorIGFBPGF binding proteinIGSGCorneal LSCsFNHoropolysin receptor kinase AIGFBP-2Ipoineline inseIGFBP-3FibrinectinIGFBP-3GF binding protein-3ADSCAAipose-derived stem cellsIGFBP-3FibrinectinIGFBP-3FibrinectinIGFBP-3FibrinectinIGFBP-3GF binding protein-3IGFBP-3FibrinectinIGFBP-3Fibr	EGFR/ErbB	Epidermal growth factor receptor
AkProtein kinaseAkiProtein kinaseRKExtracellular signal-regulated kinaseRF-kBNuclear factor kappa-BHDAC6Histone deacetylase 6CTCFCCCTC binding factorPAX6Paired box gene 6EGFR2/ErbR2Epidermal growth factor receptor 2EGFR4/ErbB4Epidermal growth factor receptor 2EGFR4/ErbB4Epidermal growth factor receptor 4HB-EGFHeparin-binding EGF-like growth factorTGF-aTansforming growth factor-aBTCBetacellulinPD0Oroneal neovascularizationCNVCorneal neovascularizationHGFHepatocyte growth factorp70s6KPhospholpase DL1βInerleukin-1βMCP-1Monocyte chemotactic protein-1MCP1Monocyte chemotactic protein-1LFSLipopolysaccharideMAPKMitogen-activated protein kinaseGFF4Isulin-like growth factorIGFBPGF binding proteinIGFBPGF binding proteinIGFBPGF binding protein-3ADSCAAjnose-derived stem cellsIGFBP-3GF binding protein-3ADSCAAjnose-derived stem cellsIGFBP-3GF binding protein-3ADSCAGrowpolic Act, YMP-9Mirt metalloproteinase-9THAToponyosin receptor kinase -9IGFAGrowpolic Act, YMAP-9Mirt metalloproteinase-9THASiebeinkart kinaseTHANeurotophic keratitis <td< td=""><td>EGFR1/ErbB1</td><td>Epidermal growth factor receptor 1</td></td<>	EGFR1/ErbB1	Epidermal growth factor receptor 1
RRKExtracellular signal-regulated kinaseNF-kBNuclear factor kappa-BHDAC6Histone deacetylase 6CTCFCCCTC binding factorPAX6Paired box gene 6EGFR2/ErbB2Epidermal growth factor receptor 3EGFR3/ErbB3Epidermal growth factor receptor 4HB-EGFEpidermal growth factor receptor 4HB-EGFBetacellulinTGF-αTansforming growth factor-αTGF-αBetacellulinPLDBospholpase DCNVCorreal neovascularizationHGFHosphoprotein 70 ribosomal protein S6 kinaseIL-1βInterleukin-1βMCP-1Mitogen-activated protein factorVGFVascular endothelial growth factorGFBAPIsulin-like growth factorTGFASingen-activated protein factorICFBGirb inding protein 70 ribosomal protein S6 kinaseIL-1βMitogen-activated protein factorMCP-1Mitogen-activated protein factorIGFBGirb inding protein 2TGFASinglin erceptorIGFBGirb inding protein-3IGFBGirb inding protein-3IGSAGirb inding protein-2TGAJoponyosin receptor kinase AIGPS-2Girb inding protein-2IGFAToponyosin receptor kinase AIGPS-2Girb inding protein-2IGFAToponyosin receptor kinase AIGPS-2Girb inding protein-3IGFAGirb inding protein-3IGFAToponyosin receptor kinase AIGPS-2Gir	PI3K	Type III phosphoinositide 3-kinase
NF-kBNuclear factor kappa-BHDAC6Histone deacetylase 6CTCFCCCTC binding factorPAX6Paired box gene 6EGFR2/ErbB2Epidermal growth factor receptor 2EGFR3/ErbB3Epidermal growth factor receptor 3EGFR4/ErbB4Peipidermal growth factor receptor 4HB-EGFHeparin-binding EGF-like growth factorTGF-qTansforming growth factor-qTGFBetacellulinPTCBetacellulinPLDPhospholipase DCNVCorneal neovascularizationHGFHepatocyte growth factorp70s6KPhosphoprotein 70 ribosomal protein S6 kinaseIL-1βInterleukin-1βMCP-1Monocyte chemotactic protein-1LFSLipopolysaccharideMAPKMitogen-activated protein kinaseVEGFInsulin-like growth factorIGFBPInsulin-like growth factor 1 receptorIGFBPInsulin-like growth factor 1 receptorIGFBPGF binding proteinIGS2Corneal LSCsFNSibrionectinIGFBP-3GF binding protein-3ADSCsGF binding protein-2TrkATopomyosin receptor kinase AIGFP-2IGF binding protein-2TrkATopomyosin receptor kinase AIGFSGio-like receptorsMMP-9Matrix metalloproteinase-9TrkAForposinbleak reatitisrhNGFAccombinant human NGFTrkANeurotrophic keratitisThKANeurotrophic keratitisCKVi	Akt	Protein kinase B
H H HDAC6 Histone deacetylase 6 CTCF CCCTC binding factor PAX6 Paired box gene 6 EGFR2/ErbB2 Epidermal growth factor receptor 2 EGFR3/ErbB3 Epidermal growth factor receptor 3 EGFR4/ErbB4 Epidermal growth factor receptor 4 HB-EGF Heparin-binding EGF-like growth factor TGF-α Transforming growth factor-α BTC Betacellulin PLD Phospholipase D CNV Corneal neovascularization HGF Hepatocyte growth factor p70s6K Phosphoprotein 70 ribosomal protein S6 kinase IL-1β Interleukin-1β MCP-1 Monocyte chemotactic protein-1 LFS Lipopolysaccharide MAPK Mitogen-activated protein kinase GFGF Insulin-like growth factor 1 receptor ISF Insulin-like growth factor 1 receptor ISF Groreal LSCs FN GF binding protein-3 ADSCs Adipose-derived stem cells IGF binding protein-2 TrkA	ERK	Extracellular signal-regulated kinase
CTCF CCCTC binding factor PAX6 Paired box gene 6 EGFR2/ErbB2 Epidermal growth factor receptor 2 EGFR3/ErbB3 Epidermal growth factor receptor 3 EGFR4/ErbB4 Epidermal growth factor receptor 4 HB-EGF Epidermal growth factor -α TGF-α Tansforming growth factor-α BTC Betacellulin PLD Phospholipase D CNV Corneal neovascularization HGF Hepatocyte growth factor p70s6K Phosphoprotein 70 ribosmal protein S6 kinase IL-1β Interleukin-1β MCP-1 Monocyte chemotactic protein-1 LFS Lipopolysaccharide MAPK Mitogen-activated protein kinase IGF Insulin-like growth factor 1 receptor IGFS Groreal LSCs FN Eibronectin IGFBP-2 IGF binding protein-3 ADSCs Adipose-derived stem cells IGFBP-3 IGF binding protein-3 ADSCs Adipose-derived stem cells IGFBP-4 Foromyosin receptor kinase A <tr< td=""><td>NF-kB</td><td>Nuclear factor kappa-B</td></tr<>	NF-kB	Nuclear factor kappa-B
PAX6 Paired box gene 6 EGFR2/ErbB2 Epidermal growth factor receptor 2 EGFR3/ErbB3 Epidermal growth factor receptor 3 EGFR4/ErbB4 Epidermal growth factor receptor 4 HB-EGF Heparin-binding EGF-like growth factor TGF-α Transforming growth factor-α BTC Betacellulin PLD Phospholipase D CNV Corneal neovascularization HGF Hepatocyte growth factor p70s6K Hosphoprotein 70 ribosomal protein S6 kinase IL-1β Interleukin-1β MCP-1 Monocyte chemotactic protein-1 LPS Lipopolysaccharide MAPK Mitogen-activated protein kinase VEGF Issulin-like growth factor IGF Insulin-like growth factor IGF Insulin-like growth factor 1 receptor IGFs Gorneal LSCs IGFBP-3 IGF binding protein-3 ADScs Adipose-derived stem cells IGFBP-3 IGF binding protein-3 ADScs Gorpomysin receptor kinase A PLC-γ Phospholipase C-	HDAC6	Histone deacetylase 6
Bit of a set of	CTCF	CCCTC binding factor
Image: Procession of the section of the se	PAX6	Paired box gene 6
Image: Properties EGFR4/ErbB4 Epidermal growth factor receptor 4 HB-EGF Heparin-binding EGF-like growth factor TGF-α Transforming growth factor-α BTC Betacellulin PLD Phospholipase D CNV Corneal neovascularization HGF Hepatocyte growth factor p70s6K Hepatocyte growth factor p70s6K Hosphoprotein 70 ribosomal protein S6 kinase IL-1β Interleukin-1β MCP-1 Monocyte chemotactic protein-1 LPS Lipopolysaccharide MAPK Mitogen-activated protein kinase VEGF Vascular endothelial growth factor IGF Insulin-like growth factor 1 receptor IGF Insulin-like growth factor 1 receptor ISCs Corneal LSCs FN Eibronectin IGFBP-3 IGF binding protein-3 ADSCs Adipose-derived stem cells IGFBP-2 IGF binding proteins-2 TrkA Topomyosin receptor kinase A PLC-γ Matrix metalloproteinase-9 TLR	EGFR2/ErbB2	Epidermal growth factor receptor 2
Here Free Formula in the	EGFR3/ErbB3	Epidermal growth factor receptor 3
Transforming growth factor-α TGF-α Transforming growth factor-α BTC Betacellulin PLD Phospholipase D CNV Corneal neovascularization HGF Hepatocyte growth factor p70s6K Phosphoprotein 70 ribosomal protein S6 kinase IL-1β Interleukin-1β MCP-1 Monocyte chemotactic protein-1 LIS Lipopolysaccharide MAPK Mitogen-activated protein kinase VEGF Vascular endothelial growth factor IGF Insulin-like growth factor 1 receptor IGF Insulin-like growth factor 1 receptor ISR Insulin receptor ISS Corneal LSCs FN Fibronectin IGFBP-3 IGF binding protein-3 ADSCs Adipose-derived stem cells IGFBP-2 IGF binding protein-2 TRA Tropomyosin receptor kinase A PLC-γ Phospholipase C-γ MMP-9 Matrix metalloproteinase-9 TLRs Recombinant human NGF NK Neurotrophic keratitis <td>EGFR4/ ErbB4</td> <td>Epidermal growth factor receptor 4</td>	EGFR4/ ErbB4	Epidermal growth factor receptor 4
Bit of the second se	HB-EGF	Heparin-binding EGF-like growth factor
PLD Phospholipase D PLD Corneal neovascularization CNV Corneal neovascularization HGF Hepatocyte growth factor p70s6K Phosphoprotein 70 ribosomal protein S6 kinase IL-1β Interleukin-1β MCP-1 Monocyte chemotactic protein-1 LPS Lipopolysaccharide MAPK Mitogen-activated protein kinase VEGF Vascular endothelial growth factor IGF Insulin-like growth factor 1 receptor IGF Insulin-like growth factor 1 receptor INSR Insulin receptor ISCS Corneal LSCs FN Fibronectin IGFBP-3 IGF binding protein-3 ADSCs Adipose-derived stem cells IGFBP-2 IGF binding protein-2 TrkA Tropomyosin receptor kinase A PLC-γ Phospholipase C-γ MMP-9 Matrix metalloproteinase-9 TLRs Tol-like receptors NK Neurotrophic keratitis NK Neurotrophic keratitis NK Diabeti	TGF-α	Transforming growth factor-α
CNVCorneal neovascularizationHGFHepatocyte growth factorp70s6KPhosphoprotein 70 ribosomal protein 56 kinaseIL-1βInterleukin-1βMCP-1Monocyte chemotactic protein-1LPSLipopolysaccharideMAPKMitogen-activated protein kinaseVEGFVascular endothelial growth factorIGFInsulin-like growth factorIGFInsulin-like growth factor 1 receptorINSRInsulin receptorLSCsCorneal LSCsFNFibronectinIGFBP-3IGF binding protein-3ADSCsAdipose-derived stem cellsIGFBP-2IGF binding protein-3ADSCsAdipose-derived stem cellsIGFBP-3Topomyosin receptor kinase APIC-γPhospholipase C-γMMP-9Matrix metalloproteinase-9TLRsToll-like receptorsHSKHerpes simplex keratitishNGFRecombinant human NGFNKNeurotrophic keratitisDKJiabetic keratopathyTGFTransforming growth factorCECsCorneal endothelial cellsCKKVernal keratoonjunctivitis	BTC	
HGFHepatocyte growth factorp70s6KPhosphoprotein 70 ribosomal protein S6 kinasep70s6KPhosphoprotein 70 ribosomal protein S6 kinaseIL-1βInterleukin-1βMCP-1Monocyte chemotactic protein-1LPSLipopolysaccharideMAPKMitogen-activated protein kinaseVEGFVascular endothelial growth factorIGFInsulin-like growth factorIGFInsulin-like growth factor 1 receptorIGF-1RInsulin-like growth factor 1 receptorISSaCorneal LSCsFNFibronectinIGFBP-3IGF binding protein-3ADSCsAdipose-derived stem cellsIGFBP-2IGF binding protein-2TrkATropomyosin receptor kinase APLC-γPhospholipase C-γMMP-9Matrix metalloproteinase-9TLRsIol-like receptorsHSKHerpes simplex keratitisNKNeurotrophic keratitisDKJiabetic keratopathyTGFTransforming growth factorCECsCorneal endothelial cells	PLD	Phospholipase D
p70s6KPhosphoprotein 70 ribosomal protein S6 kinaseIL-1βInterleukin-1βMCP-1Monocyte chemotactic protein-1LPSLipopolysaccharideMAPKMitogen-activated protein kinaseVEGFVascular endothelial growth factorIGFInsulin-like growth factorIGFInsulin-like growth factor 1 receptorINSRInsulin-like growth factor 1 receptorISSRCorneal LSCsFNFibronectinIGFBP-3IGF binding protein-3ADSCsAdipose-derived stem cellsIGFBP-2IGF binding protein-2TrkATropomyosin receptor kinase APLC-γPhospholipase C-γMMP-9Matrix metalloproteinase-9TLRsToll-like receptorsHSKHerpes simplex keratitisrhNGFRecombinant human NGFNKDiabetic keratopathyTGFTransforming growth factorCECsCorneal endothelial cellsVKCVernal keratocnjuncivitis	CNV	Corneal neovascularization
IL-1βInterleukin-1βMCP-1Monocyte chemotactic protein-1LPSLipopolysaccharideMAPKMitogen-activated protein kinaseVEGFVascular endothelial growth factorIGFInsulin-like growth factorIGFInsulin-like growth factor 1 receptorIGF-1RInsulin-like growth factor 1 receptorINSRInsulin-like growth factor 1 receptorLSCsCorneal LSCsFNFibronectinIGFBP-3IGF binding protein-3ADSCsAdipose-derived stem cellsIGFBP-2IGF binding protein-2TrAATropomyosin receptor kinase APLC-γPhospholipase C-γMMP-9Matrix metalloproteinase-9TLRsToll-like receptorsHSKHerpes simplex keratitisrhNGFRecombinant human NGFNKDiabetic keratopathyTGFTransforming growth factorCECsCorneal endothelial cellsVKCVernal keratocnjunctivitis	HGF	Hepatocyte growth factor
MCP-1Monocyte chemotactic protein-1LPSLipopolysaccharideMAPKMitogen-activated protein kinaseVEGFVascular endothelial growth factorIGFInsulin-like growth factorIGFInsulin-like growth factor 1 receptorIGF-1RInsulin-like growth factor 1 receptorINSRInsulin-like growth factor 1 receptorISCsCorneal LSCsFNFibronectinIGFBP-3IGF binding protein-3ADSCsAdipose-derived stem cellsIGFBP-2IGF binding protein-2TrkATropomyosin receptor kinase APLC-γPhospholipase C-γMMP-9Matrix metalloproteinase-9TLRsToll-like receptorsHSKHerpes simplex keratitisrhNGFRecombinant human NGFNKDiabetic keratopathyTGFTransforming growth factorCECsCorneal endothelial cellsVKCVernal keratoconjunctivitis	p70s6K	Phosphoprotein 70 ribosomal protein S6 kinase
LPS Lipopolysaccharide MAPK Mitogen-activated protein kinase VEGF Vascular endothelial growth factor IGF Insulin-like growth factor IGFBP IGF binding protein IGF-1R Insulin-like growth factor 1 receptor INSR Insulin receptor LSCs Corneal LSCs FN Fibronectin IGFBP-3 IGF binding protein-3 ADSCs Adipose-derived stem cells IGFBP-2 IGF binding protein-2 TrkA Tropomyosin receptor kinase A PLC-γ Phospholipase C-γ MMP-9 Matrix metalloproteinase-9 TLRs Toll-like receptors HSK Herpes simplex keratitis rhNGF Recombinant human NGF NK Neurotrophic keratitis DK Diabetic keratopathy TGF Transforming growth factor CECs Corneal endothelial cells	IL-1β	Interleukin-1β
MAPKMitogen-activated protein kinaseWEGFVascular endothelial growth factorIGFInsulin-like growth factorIGF binding proteinIGF binding proteinIGF-1RInsulin-like growth factor 1 receptorINSRInsulin receptorLSCsCorneal LSCsFNFibronectinIGFBP-3IGF binding protein-3ADSCsAdipose-derived stem cellsIGFBP-2IGF binding protein-2TrkATropomyosin receptor kinase APLC-γPhospholipase C-γMMP-9Matrix metalloproteinase-9TLRsToll-like receptorsHSKHerpes simplex keratitishNGFRecombinant human NGFNKDiabetic keratopathyTGFTransforming growth factorCECsCorneal endothelial cellsVKCVernal keratoconjunctivitis	MCP-1	Monocyte chemotactic protein-1
VEGFVascular endothelial growth factorIGFInsulin-like growth factorIGFBPIGF binding proteinIGF-1RInsulin-like growth factor 1 receptorINSRInsulin receptorLSCsCorneal LSCsFNFibronectinIGFBP-3IGF binding protein-3ADSCsAdipose-derived stem cellsIGFBP-2IGF binding protein-2TrkATropomyosin receptor kinase APLC-γPhospholipase C-γMMP-9Matrix metalloproteinase-9TLRsToll-like receptorsHSKHerpes simplex keratitisrhNGFRecombinant human NGFNKDiabetic keratopathyTGFTransforming growth factorCECsCorneal endothelial cellsVKCVernal keratoconjunctivitis	LPS	Lipopolysaccharide
IGF Insulin-like growth factor IGFBP IGF binding protein IGF-1R Insulin-like growth factor 1 receptor INSR Insulin receptor ISCs Corneal LSCs FN Fibronectin IGFBP-3 IGF binding protein-3 ADSCs Adipose-derived stem cells IGFBP-2 IGF binding protein-2 TrkA Tropomyosin receptor kinase A PLC-γ Phospholipase C-γ MMP-9 Matrix metalloproteinase-9 TLRs Toll-like receptors HSK Herpes simplex keratitis NK Neurotrophic keratitis DK Diabetic keratopathy TGF Transforming growth factor CECs Corneal endothelial cells	MAPK	Mitogen-activated protein kinase
IGF BPIGF binding proteinIGF-1RInsulin-like growth factor 1 receptorINSRInsulin receptorLSCsCorneal LSCsFNFibronectinIGFBP-3IGF binding protein-3ADSCsAdipose-derived stem cellsIGFBP-2IGF binding protein-2TrkATropomyosin receptor kinase APLC-γPhospholipase C-γMMP-9Matrix metalloproteinase-9TLRsToll-like receptorsHSKHerpes simplex keratitisrhNGFRecombinant human NGFNKDiabetic keratopathyTGFTransforming growth factorCECsCorneal endothelial cellsVKCVernal keratoconjunctivitis	VEGF	Vascular endothelial growth factor
IGF-1RInsulin-like growth factor 1 receptorINSRInsulin receptorLSCsCorneal LSCsFNFibronectinIGFBP-3IGF binding protein-3ADSCsAdipose-derived stem cellsIGFBP-2IGF binding protein-2TrkATropomyosin receptor kinase APLC-γPhospholipase C-γMMP-9Matrix metalloproteinase-9TLRsToll-like receptorsHSKHerpes simplex keratitishNGFRecombinant human NGFNKNeurotrophic keratitisDKDiabetic keratopathyTGFTransforming growth factorCECsCorneal endothelial cellsVKCVernal keratoconjunctivitis	IGF	Insulin-like growth factor
INSRInsulin receptorISCsCorneal LSCsFNFibronectinIGFBP-3IGF binding protein-3ADSCsAdipose-derived stem cellsIGFBP-2IGF binding protein-2TrkATropomyosin receptor kinase APLC-γPhospholipase C-γMMP-9Matrix metalloproteinase-9TLRsToll-like receptorsHSKHerpes simplex keratitisrhNGFRecombinant human NGFNKDiabetic keratopathyTGFTransforming growth factorCECsCorneal endothelial cellsVKCVernal keratoconjunctivitis	IGFBP	IGF binding protein
LSCsCorneal LSCsFNFibronectinIGFBP-3IGF binding protein-3ADSCsAdipose-derived stem cellsIGFBP-2IGF binding protein-2TrkATropomyosin receptor kinase APLC-γPhospholipase C-γMMP-9Matrix metalloproteinase-9TLRsToll-like receptorsHSKHerpes simplex keratitisrhNGFRecombinant human NGFNKDiabetic keratopathyTGFTransforming growth factorCECsCorneal endothelial cellsVKCVernal keratoconjunctivitis	IGF-1R	Insulin-like growth factor 1 receptor
FNFibronectinIGFBP-3IGF binding protein-3ADSCsAdipose-derived stem cellsIGFBP-2IGF binding protein-2TrkATropomyosin receptor kinase APLC-γPhospholipase C-γMMP-9Matrix metalloproteinase-9TLRsToll-like receptorsHSKHerpes simplex keratitisNKNeurotrophic keratitisDKDiabetic keratopathyTGFTransforming growth factorCECsCorneal endothelial cellsVKCVernal keratoconjunctivitis	INSR	Insulin receptor
IGFBP-3IGF binding protein-3ADSCsAdipose-derived stem cellsIGFBP-2IGF binding protein-2TrkATropomyosin receptor kinase APLC-γPhospholipase C-γMMP-9Matrix metalloproteinase-9TLRsToll-like receptorsHSKHerpes simplex keratitisrhNGFRecombinant human NGFNKDiabetic keratopathyTGFTransforming growth factorCECsCorneal endothelial cellsVKCVernal keratoconjunctivitis	LSCs	Corneal LSCs
ADSCsAdipose-derived stem cellsIGFBP-2IGF binding protein-2TrkATropomyosin receptor kinase APLC-γPhospholipase C-γMMP-9Matrix metalloproteinase-9TLRsToll-like receptorsHSKHerpes simplex keratitisrhNGFRecombinant human NGFNKNeurotrophic keratitisDKDiabetic keratopathyTGFTransforming growth factorCECsCorneal endothelial cellsVKCVernal keratoconjunctivitis	FN	Fibronectin
IGFBP-2 IGF binding protein-2 TrkA Tropomyosin receptor kinase A PLC-γ Phospholipase C-γ MMP-9 Matrix metalloproteinase-9 TLRs Toll-like receptors HSK Herpes simplex keratitis rhNGF Recombinant human NGF NK Neurotrophic keratitis DK Diabetic keratopathy TGF Transforming growth factor CECs Corneal endothelial cells VKC Vernal keratoconjunctivitis	IGFBP-3	IGF binding protein-3
TrkA Tropomyosin receptor kinase A PLC-γ Phospholipase C-γ MMP-9 Matrix metalloproteinase-9 TLRs Toll-like receptors HSK Herpes simplex keratitis rhNGF Recombinant human NGF NK Neurotrophic keratitis DK Diabetic keratopathy TGF Transforming growth factor CECs Corneal endothelial cells VKC Vernal keratoconjunctivitis	ADSCs	Adipose-derived stem cells
PLC-γ Phospholipase C-γ MMP-9 Matrix metalloproteinase-9 TLRs Toll-like receptors HSK Herpes simplex keratitis rhNGF Recombinant human NGF NK Neurotrophic keratitis DK Diabetic keratopathy TGF Transforming growth factor CECs Corneal endothelial cells VKC Vernal keratoconjunctivitis	IGFBP-2	IGF binding protein-2
MMP-9Matrix metalloproteinase-9TLRsToll-like receptorsHSKHerpes simplex keratitisrhNGFRecombinant human NGFNKNeurotrophic keratitisDKDiabetic keratopathyTGFTransforming growth factorCECsCorneal endothelial cellsVKCVernal keratoconjunctivitis	TrkA	Tropomyosin receptor kinase A
TLRs Toll-like receptors HSK Herpes simplex keratitis rhNGF Recombinant human NGF NK Neurotrophic keratitis DK Diabetic keratopathy TGF Transforming growth factor CECs Corneal endothelial cells VKC Vernal keratoconjunctivitis	PLC-γ	Phospholipase C-y
HSK Herpes simplex keratitis rhNGF Recombinant human NGF NK Neurotrophic keratitis DK Diabetic keratopathy TGF Transforming growth factor CECs Corneal endothelial cells VKC Vernal keratoconjunctivitis	MMP-9	Matrix metalloproteinase-9
rhNGFRecombinant human NGFNKNeurotrophic keratitisDKDiabetic keratopathyTGFTransforming growth factorCECsCorneal endothelial cellsVKCVernal keratoconjunctivitis	TLRs	Toll-like receptors
NK Neurotrophic keratitis DK Diabetic keratopathy TGF Transforming growth factor CECs Corneal endothelial cells VKC Vernal keratoconjunctivitis	HSK	Herpes simplex keratitis
DK Diabetic keratopathy TGF Transforming growth factor CECs Corneal endothelial cells VKC Vernal keratoconjunctivitis	rhNGF	Recombinant human NGF
TGF Transforming growth factor CECs Corneal endothelial cells VKC Vernal keratoconjunctivitis	NK	Neurotrophic keratitis
CECs Corneal endothelial cells VKC Vernal keratoconjunctivitis	DK	Diabetic keratopathy
VKC Vernal keratoconjunctivitis	TGF	Transforming growth factor
, , , , , , , , , , , , , , , , , , ,	CECs	Corneal endothelial cells
(Continued	VKC	Vernal keratoconjunctivitis
		(Continued

GLOSSARY (Continued)

АКС	Atopic keratoconjunctivitis
GVHD	Graft-versus-host disease
PDGF	Platelet-derived growth factor
BCEC	Bovine corneal endothelial cells
HCF	Human corneal fibroblasts
EMT	Epithelial-mesenchymal transition
FGF	Fibroblast growth factor
FGFR	Fibroblast growth factor receptor
b-FGF/FGF2	Basic FGF
ROS	Reactive oxygen species
KGF	Keratinocyte growth factor
OGF	Opioid growth factor
OGFr	Opioid growth factor receptor
NTX	Naltrexone

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