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3	TAK1 signaling is a potential therapeutic target for pathological angiogenesis								
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26 Abstract

Angiogenesis plays a critical role in both physiological responses and disease pathogenesis. Excessive 27 angiogenesis can promote neoplastic diseases and retinopathies, while inadequate angiogenesis can 28 lead to aberrant perfusion and impaired wound healing. Transforming growth factor β activated kinase 29 1 (TAK1), a member of the mitogen-activated protein kinase kinase kinase (MAPKKK) family, is a 30 key modulator involved in a range of cellular functions including the immune responses, cell survival 31 32 and death. TAK1 is activated in response to various stimuli such as proinflammatory cytokines, hypoxia, and oxidative stress. Emerging evidence has recently suggested that TAK1 is intimately 33 34 involved in angiogenesis and mediates pathogenic processes related to angiogenesis. Several detailed mechanisms by which TAK1 regulates pathological angiogenesis have been clarified, and potential 35 therapeutics targeting TAK1 have emerged. In this review, we summarize recent studies of TAK1 in 36 angiogenesis and discuss the crosstalk between TAK1 and signaling pathways involved in pathological 37 angiogenesis. We also discuss the approaches for selectively targeting TAK1 and highlight the 38 39 rationales of therapeutic strategies based on TAK1 inhibition for the treatment of pathological angiogenesis. 40

42 **1. Introduction**

Angiogenesis is the process by which new capillaries grow from preexisting blood vessels. It is 43 fundamental in embryonic vascular development and reproduction, as well as wound healing and 44 repair in adults. During embryonic development, angiogenesis is the basis for the maturation of the 45 circulatory system. Angiogenesis starts with the proliferation of endothelial cells, followed by 46 endothelial tube formation, a process enriched by smooth muscle cells, and later facilitates the 47 48 formation of a specific vascular system [1]. Upon injury, severed vessels are elongated and anastomosed with each other, and the vessels then become tortuous with endothelial cell proliferation 49 50 and pericyte coverage, eventually normalizing through vessel regression over a few months [2]. 51 Angiogenesis is a highly regulated process that is activated under physiological stresses and inactivated 52 when those stresses are relieved [3].

53 Angiogenesis is also central to several pathological conditions, such as solid tumors and neovascular eye diseases. Under pathological conditions such as tumor growth, host blood vessels are 54 55 stimulated to grow into the vicinity of the tumor to maintain cell growth, and tumor vascularization is 56 characterized by dilated, tortuous and disorganized blood vessels [4,5]. Tumors constantly promote 57 the growth of new blood vessels to ensure an adequate supply of nutrients for expansion, and these 58 new vessels also provide potential routes for tumor metastasis [6]. In addition, pathological 59 angiogenesis, a process responsive to inadequate perfusion or ischemia, occurs in some eye diseases, 60 particularly in the retina. Such ocular neovascularization can result in severe impairment of vision. For instance, retinal neovascularization occurs in patients with advanced diabetic retinopathy, often 61 62 leading to fundus hemorrhage and severe vision impairment due to invasion and leakage of fluid from abnormal blood vessels into the retina and vitreous. 63

64 Transforming growth factor β (TGF- β) activated kinase 1 (TAK1) is a key regulator of immune and proinflammatory signaling pathways [7]. TAK1 activates nuclear factor-kappa B (NF- κ B) and 65 mitogen-activated protein kinase (MAPK) pathways in response to a diverse range of stimuli, 66 including inflammation, hypoxia and oxidative stress, the major causes of angiogenesis [8]. Activation 67 of the NF-kB and MAPK pathways regulated by TAK1 promotes the expression of various 68 69 inflammatory response proteins, including those encoding cytokines and chemokines, and participates 70 in inflammasome regulation, all of which in turn facilitate angiogenic processes [9]. Naito et al. found 71 that, in addition to its role in mediating inflammatory signals, activated TAK1 can prevent endothelial 72 apoptosis and maintain vascular integrity under inflammatory conditions [10]. In fact, TAK1 deficiency leads to embryonic lethality due to vascular destruction, which implies its crucial role in 73 74 maintaining vascular integrity during embryogenesis [11]. Therefore, a better understanding of the 75 mechanism that underlies TAK1-mediated signaling in angiogenesis is of great significance for 76 developing therapeutic strategies for the management of pathological angiogenesis.

77

78 **2. TAK1 and its activity**

79 TAK1 was discovered in 1995 as a member of the mitogen-activated protein kinase kinase kinase 80 (MAPKKK) family [12]. It is a critical signal transduction mediator that can be activated by cell membrane receptor interacting protein kinases or second messengers in cells after a variety of 81 82 stimulations, including proinflammatory cytokines or antigens such as tumor necrosis factor α (TNF α), 83 IL-1, or lipopolysaccharides (LPS). Normally, TAK1 binds to adaptor proteins such as TAK-binding protein 1 (TAB1) and its homologs TAB2 and TAB3 to form heterotrimeric complexes consisting of 84 85 either TAK1-TAB1-TAB2 or TAK1-TAB1-TAB3 [8]. TAB1 binds to the N-terminal kinase domain of TAK1, whereas the homologs TAB2 and TAB3 bind to the C-terminal region (Figure 1). Through 86 different signaling pathways, both TAB1 and TAB2 activate the TAK1 protein. TAB1 is essential for 87 osmotic stress-induced TAK1 activation, whereas TAB2 or TAB3 is required for TNF- or IL-1-88 induced TAK1 activation (Figure 2) [13]. 89

Although TAB1 constitutively binds to TAK1, it possesses no phosphatase or other enzymatic 90 91 activity. Pathak et al. recently reported that glycosylation with N-acetylglucosamine (O-GlcNAcylation) of a single residue (Ser395) on TAB1 can modulate the activation of TAK1 in 92 93 response to IL-1 stimulation or osmotic stress [14]. O-GlcNAcylation of TAB1 substantially increases the autophosphorylation of TAK1, phosphorylation of inhibitory kappa B kinase (IKK) and 94 translocation of NF-kB, which results in increased production of cytokines. Moreover, an E3 ubiquitin 95 96 ligase X-linked inhibitor of apoptosis protein (XIAP) has also been found to directly interact with 97 TAB1 and to further activate TAK1 as a downstream biological factor in TGF-β receptor (TGFBR) and bone morphogenetic protein (BMP) receptor (BMPR) activation through formation of the XIAP-98 99 TAK1-TAB1 complex. Activated TAK1 then upregulates the expression of NF-kB and transcription 100 factor activator protein-1 (AP-1) by activating the NF- κ B and MAPK (JNK and p38) pathways [15] 101 (Figure 2A). However, the detail mechanism by which TAK1 is activated by TAB1 remains unclear. 102 In contrast to TAB1, TAB2 and its analogous protein TAB3 have been extensively studied in TAK1-mediated signaling pathways in angiogenesis. When IL-1 and LPS bind to their receptors, 103 interleukin-1 receptor kinase 1 (IRAK1) and IRAK4 recruit TNF receptor (TNFR)-associated factor-104 6 (TRAF6) and its associated enzymes ubiquitin conjugating enzyme 13 (Ubc13) and ubiquitin E2 105 variant 1a (Uev1a). In a similar manner, when TNF- α binds to TNFR, receptor interacting protein 106 107 kinase (RIPK1) recruits TRAF2/5 with its associated enzymes Ubc13 and Uev1b. TRAF2/5 complexes generate lysine 63 (K63)-linked polyubiquitin chains on either TAB2 or TAB3, thus activating TAK1 108

[16-18]. K63-linked polyubiquitin activates TAK1 by inducing conformational changes that lead to
the autophosphorylation of Thr187, Thr178, Thr184, and Ser192 residues [19,20]. Activated TAK1
then phosphorylates IKK, MKK4/7 and MKK3/6 to activate NF-κB, JNK and p38 MAPK, respectively

112 (Figure 2B). Collectively, these involved signaling pathways result in inflammation and immune

113 responses, apoptosis and angiogenesis [21].

114 TAK1-TAB2 maintains vascular homeostasis under TNF- α stimulation by preventing endothelial apoptosis [22]. Morioka et al. reported that cell migration and tube formation were significantly 115 affected in TAK1- and TAB2-deficient endothelial cells but not in TAB1-deficient endothelial cells, 116 suggesting that TAB2 instead of TAB1 plays an important role in angiogenesis [11]. Furthermore, 117 118 TAB2 deficiency in mouse embryos led to the abnormal growth of capillary blood vessels due to reduced TAK1 activity, revealing that TAB2 is crucial for maintaining normal vascular homeostasis 119 [11]. Nonetheless, the activation of TAK1 is arguably regulated by both TAB1 and TAB2/3, and their 120 respective contributions are complex and dependent upon the tissue type and cellular context [23]. 121

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3. Molecular mechanisms of TAK1 involved in angiogenic activities

Inflammation is a physiological response to harmful stimuli such as pathogens, damaged cells and 124 125 toxic compounds with the overall aim of removing the source of injury and repairing damaged tissue to restore tissue architecture and maintain tissue homeostasis [24]. When inflammation lasts for an 126 127 extended period of time, endothelial cells proliferate and migrate to form new capillaries to restore nutrient supply, therefore facilitating the immune response [25]. Inadequate supply of vasculature and 128 129 the resultant reduction in oxygen level leads to angiogenesis to fulfill the oxygen needs of tissue [26]. Apart from tissue hypoxia and inflammation, oxidative stress plays a significant role in angiogenesis. 130 Oxidative stress is the excessive production of reactive oxygen species (ROS) and reactive nitrogen 131 species (RNS) in the tissue under harmful stimuli. However, short exposure to ROS or low levels of 132 ROS can also promote physiological angiogenesis and maintain healthy blood vessel homeostasis [27]. 133 There is substantial evidence that inflammation, hypoxia and oxidative stress are three important 134 135 inducers of angiogenesis, and each has a complicated molecular mechanism for promoting and inhibiting angiogenic activities [6,28]. Given that TAK1 is an important mediator in many pathways 136 involved in angiogenesis, its role and function in mediating inflammation, hypoxia and oxidative stress 137 in angiogenesis are discussed below. 138

139

140 **3.1 TAK1** activates the inflammatory response

141 There is increasing evidence that inflammation plays a central role in various pathophysiological

processes, such as angiogenesis. Inflammation has been shown to be involved in angiogenesis via 142 several physiological processes, such as embryonic development and tissue repair [29], as well as 143 144 angiogenic diseases, such as a variety of tumors and neovascular eye diseases. TAK1 has been identified as a key mediator in inflammation and defense immune signaling pathways [30]. TAK1 is 145 activated by several inflammatory signaling pathways, such as the IL-1 β , TNF- α , Toll-like receptor 146 (TLR), T-cell receptor (TCR) and B-cell receptor (BCR) signaling pathways, after TRAF6 and the 147 148 ubiquitin-binding enzyme complex (Ubc13 and Uev1a) catalyzes the polyubiquitination of the Lys63 residue on TAB2 or TAB3 [31]. Activated TAK1 phosphorylates NF-kB-inducing kinase (NIK) and 149 IKK or MAP kinase kinases (MKKs), which leads to the activation of NF-κB and AP-1 [32], ultimately 150 resulting in the expression of inflammatory cytokines (e.g., IL-1 and IL-6), chemokines (e.g., CXCL1 151 and IL-8) or adhesion molecules (e.g., intercellular adhesion molecule-1 (ICAM-1) and vascular cell 152 153 adhesion molecule-1 (VCAM-1)) that participate in tissue inflammatory and angiogenic responses (Figure 3). As such, these inflammatory proteins induce diverse physiological and pathological effects, 154 such as tissue repair and tumor progression [33-37]. Therefore, TAK1, which is involved in 155 156 inflammatory signaling pathways, has been found to play a vital role in the development of multiple 157 physiopathological conditions, especially in angiogenic processes.

A number of studies have attempted to determine the role of TAK1-induced inflammatory 158 159 signaling in angiogenic diseases. Singh et al. found that arginine transferase 1 (ATE1) gene knockout in mouse embryos can cause contractile dysfunction, cardiovascular dysplasia and impaired 160 angiogenesis due to blockage of the TAK1-dependent JNK1/2 signaling pathway [38]. Moreover, 161 blocking TAK1 inhibited NF- κ B by downregulating the phosphorylation of IKK α/β and NF- κ B p65, 162 163 resulting in the reduced expression of proinflammatory genes, such as IL-6, monocyte chemoattractant 164 protein-1 (MCP-1) and ICAM-1, in vascular smooth muscle cells, ultimately leading to attenuation of neointimal formation in wire-injured femoral arteries [39]. 165

TNF- α can induce endothelial cell death during inflammation via either caspase-dependent 166 167 apoptosis or RIP1 kinase-dependent necrosis [40]. Naito et al. found severe bleeding in the lung and hind limb muscles in endothelium-specific TAK1 knockout mice, an animal model of acute lung and 168 169 muscle inflammation, and this effect was not observed in wild-type mice. Immunostaining of knockout mouse tissues revealed high levels of TNF- α in CD11b+ and F4/80+ hematopoietic cells and vascular 170 171 deformation or massive loss of blood vessels, indicating that TAK1 is essential for endothelial cell survival through inhibition of inflammatory apoptosis induced by TNF- α during acute inflammation 172 [10]. In addition, inhibiting TAK1 alleviated joint inflammation and pannus caused by abnormal 173 174 neovascularization in a collagen-induced mouse model of rheumatoid arthritis [41], suggesting that TAK1 plays a crucial role in promoting inflammation and angiogenesis in rheumatoid arthritis. Chang *et al.* further demonstrated that TAK1 phosphorylation is enhanced upon adenosine monophosphateactivated protein kinase (AMPK) activation *in vivo* and *in vitro*, leading to a proinflammatory phenotype in endothelial cells that facilitates angiogenesis via a downstream p38 MAPK signaling cascade [42]. Moreover, overexpression of TAK1 and TAB1 also enhances the phosphorylation of AMPK in cervical cancer cells [43], suggesting that TAK1 and AMPK are more likely to act together rather than alone to regulate these processes under different circumstances.

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183 **3.2 TAK1 signaling in hypoxia**

184 Hypoxia is known as a reduction in oxygen supply that cannot meet cellular requirements. It is one of 185 the key mechanisms involved in both physiological and pathological angiogenesis. Among several 186 regulatory genes in hypoxia, hypoxia-inducible factor 1 (HIF-1) plays an important role in facilitating 187 the response of cells to changes in systemic oxygen levels. HIF-1 is a heterodimeric transcription factor that consists of a constitutively expressed β -subunit (HIF-1 β) and an oxygen-regulated α -subunit 188 189 (HIF-1 α) [44]. HIF-1 α contains an oxygen-dependent degradation (ODD) domain that is hydroxylated by prolyl hydroxylase 2 (PHD2) under normoxic conditions to degrade HIF-1 through the ubiquitin-190 proteasome pathway [45]. However, hypoxia can strongly induce HIF-1 accumulation by preventing 191 192 HIF-1 α ubiquitination, which activates anaerobic metabolism and inflammation-related signaling pathways, including the MAPK [46,47], NF-kB [48] and AMPK [49] [50] pathways, in cells. 193 194 Interestingly, TAK1 is closely related to these pathways [23], indicating that TAK1 may participate in 195 hypoxia-induced angiogenesis (Figure 4). Indeed, studies related to cancer progression and 196 cardiomyocyte hypertrophy have shown that hypoxia can activate TAK1 via a mechanism that is 197 dependent upon the activation of calcium calmodulin kinase (CaMK2) signaling and is mediated through the Ubc13-XIAP complex, resulting in the activation of NF-kB and the promotion of an 198 199 inflammatory state in cells [51-53]. Such processes are mediated by NF-kB and MAPK signaling, both 200 of which are downstream of TAK1 activation [54]. Unlike other signaling pathways, the role of TAK1 as a genuine upstream kinase of AMPK is still highly debated. In several studies, the potential role of 201 TAK1 as an upstream mediator of AMPK activation was verified using various knockdown strategies 202 [55-57]. Nagata et al. found that inhibition of AMPK signaling can inhibit endothelial cell migration 203 204 and tube formation under hypoxic conditions and suppress the growth of blood vessels in mice subcutaneously implanted with Matrigel [58]. Although a number of studies have shown that TAK1 is 205 closely related to hypoxia-induced angiogenesis, there is no clear evidence that TAK1 is causally 206 207 involved in angiogenesis under hypoxic conditions.

209 **3.3 TAK1 signaling in oxidative stress**

Angiogenesis can be affected by oxidative stress in different ways. When the degree of oxidation 210 exceeds the oxide clearance rate, the oxidation system and antioxidant system become unbalanced, 211 resulting in pathophysiological changes in tissue [28]. TAK1 also participates in redox regulation 212 through various cellular signaling pathways [59], which may be related to pathological angiogenesis 213 (Figure 5). Zippel et al. reported that TAK1 knockdown by siRNA results in a significant change in 214 the proteins that are involved in redox regulation in IL-1B-treated endothelial cells [60]. Kajino-215 216 Sakamoto et al. showed that ablation of TAK1 leads to the accumulation of ROS in the intestinal 217 epithelium by reducing the expression of nuclear factor-erythroid 2 (NF-E2)-related factor 2 (NRF2), a key antioxidant transcription factor, and related antioxidant-responsive molecules [61]. ROS 218 219 accumulation results in epithelial cell death, causing intestinal hemorrhage. NRF2 is known to promote 220 angiogenesis by regulating NADPH oxidase 2 (Nox2) in several physical and pathological conditions, 221 such as corneal neovascularization, ischemia-induced retinopathy, and tissue repair [62-65]. Reasonably, TAK1 is able to protect epithelial or endothelial cells from ROS-induced death by 222 223 regulating NRF2 and NOX-related signals, promoting blood vessel formation. Indeed, Menden et al. 224 reported that silencing Nox2 can suppress LPS-induced ICAM-1 expression through inhibition of 225 TAK1 phosphorylation (Thr184/187) in human pulmonary microvascular endothelial cells, which 226 limits macrophage-endothelial cell interactions and lung microvascular remodeling [66].

Superoxide dismutase (SOD), another endogenous antioxidant, also plays an important role in the 227 oxidative stress response and is involved in TAK1-related angiogenesis. Zippel et al. found that TAK1 228 is an AMPK mediator that regulates angiogenesis by modulating SOD2 and redox signaling in 229 230 endothelial cells [57]. Specifically, the dysregulated endothelial germination processes of ring and tube formation are normalized in the presence of polyethylene glycol-SOD under the condition of 231 endothelial cell-specific TAK1 knockout in the aortic ring model. Similar rescue of angiogenesis was 232 also observed in polyethylene glycol-SOD-treated aortic rings from AMPKa1 knockout mice [60]. 233 Since AMPK can be activated under oxidative stress, can facilitate angiogenesis [67,68], and closely 234 235 interacts with TAK1, it can be implied that TAK1 plays a role in oxidative stress-induced angiogenesis. 236 However, there is still a lack of evidence that redox signaling directly activates TAK1, and the crosstalk between TAK1 and oxidative stress signals in angiogenesis remains unclear. 237

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4. Translational potentials in diseases associated with pathological angiogenesis

Angiogenesis is an important event in a variety of physiological settings, and it is also central to the pathogenesis of several pathological conditions. Activated TAK1 participates in crucial signaling pathways of inflammation, hypoxia and oxidative stress, which could lead to pathological angiogenesis under these conditions. We therefore discuss below the crosstalk between TAK1 and the signaling pathways involved in pathological angiogenesis processes, such as tumor angiogenesis and retinal neovascularization.

246

247 **4.1 Tumor angiogenesis**

Tumor growth and metastasis depend upon angiogenesis, which is usually stimulated by chemical 248 signals from the tumor cells themselves [69]. Tumor cells may become necrotic or apoptotic without 249 250 vascular support [70]. Therefore, tumors need to be supported by the rapid development of a new 251 vascular network in order to progress [6]. TAK1 acts as a key mediator of angiogenic signaling in tumor environments. TAK1 regulates MAPK signaling through P38 MAPK and JNK activation, which 252 253 promotes the expression of VEGF, plasminogen activator inhibitor-1 (PAI-1) and MMPs, which are 254 involved in vascular remodeling, angiogenesis and extracellular matrix degradation in tumors such as glioma [71,72]. In addition to angiogenesis, TAK1 plays a significant role in preventing TNF- α -255 induced endothelial cell death. Knocking out or inhibiting TAK1 can induce the apoptosis of 256 endothelial cells and destroy tumor vasculature, resulting in tumor regression [10]. Safina et al. showed 257 258 that deletion of TAK1 can reduce the activity of NF-kB and the expression of MMP-9, thereby suppressing TGF-B-mediated tumor angiogenesis and metastasis [73]. Furthermore, studies also 259 revealed that the inhibition of TAK1 with natural or artificial compounds such as cyclopeptide RA-V 260 261 and triterpene celastrol suppressed angiogenesis and tumorigenesis.

262 In addition to endogenous angiogenic genes, hypoxia-related genes in the tumor 263 microenvironment play a critical role in activating TAK1 and ultimately promoting tumor angiogenesis [74]. HIF-1 α , a master regulator of the hypoxia response, can induce NF- κ B activation 264 in a TAK1-dependent manner [53]. Activation of this inflammatory pathway can in turn promote the 265 266 expression of HIF-1 α itself, forming a positive feedback loop. Such crosstalk between hypoxia and inflammation, which is centrally regulated by TAK1, further enhances tumor cell proliferation and 267 angiogenesis [51]. Although there is still a lack of evidence that TAK1 directly leads to HIF-1 268 269 accumulation, it is reasonable to postulate that NF-kB might be a potential node between TAK1 and 270 HIF-1.

271

272 4.2 Retinal neovascularization

Retinal neovascularization refers to abnormal vascular growth with increased permeability of blood
vessels in the retina resulting in severe retinal hemorrhage and even blindness. Numerous studies have

shown that VEGF signaling certainly plays a key role in retinal neovascularization [75,27,76,77].
Hence, anti-VEGF drugs have been extensively studied and have been demonstrated to be effective in
suppressing retinal neovascularization, thus ameliorating vision impairment. However, recent clinical
trials showed that anti-VEGF therapy is not effective for all patients and that patients who benefit from
treatment exhibit a high recurrence rate [78-80]. Therefore, it is important to look for other therapeutic
target genes for retinal neovascularization.

Studies have shown that hypoxia and ischemia in the retina contribute to the progression of retinal 281 neovascularization through HIF-1 α - and NF- κ B-related signaling involving TAK1 signaling [81]. 282 TAK1 was found to be activated under hypoxic conditions, which stimulates the expression of 283 proinflammatory and proangiogenic cytokines, including ICAM-1, IL-8 and TNF-α, through NF-κB 284 [52,53,82]. Our recent study provided the first piece of evidence that TAK1 inhibition can significantly 285 attenuate retinal neovascularization in a rat model of ischemia-induced retinopathy [83]. The data 286 further suggest that selective inhibition of TAK1 by 5Z-7-oxozeaenol ameliorates the inflammatory 287 response, which contributes to the promotion of aberrant retinal angiogenesis [83]. Furthermore, 288 hypoxia and ischemia in the retina are accompanied by the production of ROS, including H₂O₂, and 289 the induction of inducible nitric oxide 2 (NOS2). Hypoxia and ischemia promote retinal angiogenesis 290 291 by upregulating the antioxidant transcription factor NRF2 and SOD and enhance the expression of epidermal growth factor (EGF), IL-8, platelet-derived growth factor (PDGF) and adhesion molecules 292 under oxidative stress [84,65,85]. Interestingly, TAK1 has been found to be involved in compensatory 293 cellular antioxidant responses, including NFR2- and SOD-related signaling pathways [60,61]. TAK1 294 295 therefore appears to be crucial in pathological angiogenesis, suggesting that TAK1 could be a potential therapeutic target for retinal neovascularization. 296

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5. Approaches for therapeutic targeting of TAK1 signaling

299 5.1 Inhibition of TAK1 activity with small molecule drugs

The role of TAK1 in multiple cellular pathways suggests that it might be a potential target for small molecule interventions against diseases, including cancer and inflammation- and angiogenesis-related diseases (**Table 1**).

303

304 5.1.1 (5Z)-7-oxozeaenol

(5Z)-7-Oxozeaenol, a kind of macrolide compound, is the 7-Oxo derivative of zeaenol (the 5Z
 stereoisomer) [86]. It is a natural product of fungal origin that functions as a TAK1-specific inhibitor
 through covalent interactions with TAK1 [87]. The therapeutic effects of (5Z)-7-oxozeaenol have been

observed in a number of studies. For example, (5Z)-7-oxozeaenol was found to inhibit TAK1 activity 308 and downregulate downstream signaling pathways, including p38 MAPK, IKK and JNK, and reduce 309 chemokine receptor 7 (CCR7) expression, ultimately suppressing the lymphatic invasion and lung 310 metastasis of breast cancer [88,89]. In addition, other studies have shown that treatment with (5Z)-7-311 312 oxozeaenol effectively inhibits TAK1 and NF-κB activation and induces caspase-3 and -7 in colon and cervical cancer, resulting in enhanced apoptosis of cancer cells [90,91]. In neurological diseases, such 313 314 as cerebral ischemia and subarachnoid hemorrhage, studies have shown that inhibiting TAK1 can downregulate p38 MAPK-JNK and NF-kB-related inflammatory pathways, thereby reducing cerebral 315 316 inflammation and brain damage [92-94]. Similarly, treatment with (5Z)-7-oxozeaenol can also reduce 317 the production of inflammatory cytokines and the formation of abnormal blood vessels in the cavum 318 articulare by suppressing synovial fibroblast activation [140] and attenuating neointimal formation in 319 wire-injured femoral arteries of mice [50]. Although (5Z)-7-oxozeaenol is widely used to study the biological functions of TAK1 in diseases, it also effectively inhibits a panel of at least 50 other kinases 320 321 and forms a covalent bond with reactive cysteines in the activation loop of its targets, producing several undesired side effects. Such nonspecific binding creates off-target effects, which likely limits its 322 323 potential use in clinical settings [95,87].

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325 5.1.2 NG25

326 Other small molecules that target TAK1 have also been investigated. Tan et al. found that NG25 is a potent dual inhibitor that targets TAK1 and MAP4K2 kinases, with weak inhibition of 11 other kinases 327 [96]. Wang et al. also reported that targeting TAK1 with NG25 can partially block doxorubicin (Dox)-328 induced p38 MAPK phosphorylation and IkBa degradation and enhance Dox-induced cytotoxic 329 effects and apoptosis in breast cancer cells by targeting TAK1 [97]. Wang et al. further confirmed that 330 injection of NG25 prior to insult significantly inhibited TAK1/JNK activity and dramatically 331 attenuated acute hypoxic and ischemic cerebral injury and abnormal angiogenesis by regulating cell 332 survival and behavior in perinatal rats [98]. Therefore, NG25 may also be a potential candidate drug 333 that can be applied to target TAK1 by inhibiting TAK1-related inflammation and angiogenesis. 334

335

336 5.1.3 Takinib

A recently developed compound named takinib has proven to bind more specifically to TAK1 than (5Z)-7-oxozeaenol. Totzke *et al.* found that takinib is more selective than other TAK1 inhibitors since it targets germinal center kinase (GCK), an important kinase that participates in both the determination of cell fate and the regulation of cell functions, with a 45-fold lower potency than TAK1 [95]. Takinib is an aminobenzimidazole-based competitive inhibitor of TAK1 that was previously identified as a Src

kinase family inhibitor. However, the initial kinome profiling study showed that takinib only weakly 342 inhibited Src and Yes1 [99]. In contrast, takinib shows significant inhibitory activity against six other 343 kinases, including TAK1, IRAK4, IRAK1, GCK, CDC-like kinase 2 (CLK2), and misshapen like 344 kinase 1 (MINK1); of these targets, TAK1 is most potently inhibited by takinib [100]. Compared to 345 (5Z)-7-oxozeaenol, takinib does not inhibit any members of the MAP2K or MAP3K family and shows 346 no efficacy on TAK1-related MAP3K5/apoptosis signal-regulating kinase 1 (ASK1). Additionally, 347 p38 MAPK is completely insensitive to takinib [95]. Due to its higher specificity for TAK1 and its 348 capability to phosphorylate IKK, MAPK 8/9 and c-Jun upon TNF-α stimulation, takinib induces 349 350 apoptosis upon TNFa stimulation in cell models of breast cancer and rheumatoid arthritis [95]. 351 Furthermore, takinib treatment was found to inhibit proinflammatory cytokines in a mouse model of 352 type II collagen-induced arthritis and in NRAS-mutated melanoma cells through TAK1 inhibition 353 [101], suggesting that it may be useful in progressive malignant diseases and inflammatory diseases.

354

355 **5.1.4 LYTAK1**

Other orally active TAK1 inhibitors, such as LYTAK1, have been described; LYTAK1 attenuates the 356 357 chemoresistance of pancreatic cancer by inhibiting TAK1 but has cytotoxic activity in vitro [102]. LYTAK1 was reported to significantly suppress LPS-induced TAK1-NFkB and MAPK (ERK, JNK 358 359 and p38 MAPK) activation in vitro and in vivo [103]. Oral administration of LYTAK1 can significantly 360 inhibit the growth of colorectal cancer cell xenografts in nude mice [104]. Moreover, LYTAK1 attenuates proliferation and epithelial-mesenchymal transition in retinal pigment epithelial cells 361 through the TAK1-mediated Smad and ERK/AKT signaling pathways, which may further attenuate 362 retinal neovascularization [105,106]. 363

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365 5.1.5 Other TAK1 inhibitors

Some uncommon TAK1 inhibitors, including fisetin, gamma-tocotrienol and tanshinone IIA, are 366 currently being developed [107-109]. In addition, there are some new discoveries of plant extracts that 367 may be useful for inhibiting TAK1. Rubiaceae-type cyclopeptides, a type of plant cyclopeptide from 368 369 Rubia, can inhibit the NF-kB signaling pathway by disrupting the TAK1-TAB2 interaction and 370 targeting TAK1, ultimately suppressing the inflammatory response and angiogenesis [110]. Other molecules, such as sesamin, pinitol, gambogic acid and celastrol, can inhibit the NF-kB signaling 371 pathway and related genes involved in apoptosis (cIAP-1/2, Bcl-2, Bcl-xL, XIAP, survivin, and 372 TRAF1), proliferation (cyclin D1, c-Myc, COX2), metastasis (ICAM-1 and MMP-9), and 373 angiogenesis (VEGF) by targeting TAK1, thus enhancing apoptosis and attenuating proliferation, 374 invasion and angiogenesis in cancer [111-114]. However, even though there are a number of studies 375

on the development of different kinds of TAK1 inhibitors and the characterization of their specific mechanisms and functions, there is still a lack of evidence regarding their clinical effects. In this case, more preclinical studies are needed to determine whether there is potential to develop TAK1 inhibitors as potential antiangiogenic therapies for various diseases.

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5.2 Genetic approaches for TAK1 gene targeting

Although the aforementioned TAK1 inhibitors can inhibit the activation of TAK1 at the protein level, 382 their off-target effects may need to be considered. Even though more selective TAK1 inhibitors have 383 384 been reported, most of them can also target a wide type of kinases other than TAK1 [95,86,96]. 385 Therefore, the side effects induced by many TAK1 inhibitors remain largely unclear. Thus, 386 pharmaceutical inhibitors of TAK1 may not be ideal candidates for specifically inhibiting TAK1. As 387 a result, emerging approaches such as microRNA (miRNA)-based targeting strategies and clustered regularly interspaced short palindromic repeat (CRISPR)-based gene editing have been increasingly 388 389 applied in research.

390

391 5.2.1 TAK1 regulation by miRNA

miRNAs are small noncoding RNA molecules (22 to 25 nucleotides long) found in all eukaryotes and 392 393 some viruses. miRNA silence gene expression at the posttranscriptional level through base pairing 394 with complementary sequences at the 3' untranslated region (3'-UTR) of mRNA [115-117]. The role 395 of various miRNAs in TAK1 inhibition has been extensively studied. Jiang et al. revealed that when 396 overexpressed, miR-892b can attenuate NF-κB signaling by directly targeting and suppressing TAK1 in breast cancer, resulting in significantly decreased tumor growth, metastatic capacity and 397 398 angiogenesis [118]. Likewise, miR-26b can also inhibit the expression of TAK1 and TAB3 by binding to their 3'-UTRs, thus blocking the activation of NF-kB signaling and sensitizing cells to apoptosis 399 [119]. 400

TAK1 silencing by miR-143 has been shown in pancreatic ductal adenocarcinoma cells and 401 hepatocytes. miR-143 can directly target TAK1 and inactivate MAPK/NF-KB signaling, therefore 402 403 inhibiting cell proliferation, cell migration, inflammation and fibrosis, which are important cellular 404 activities related to angiogenesis [120]. TAK1 can also be targeted by miR-10a in endothelial cells; 405 miR-10a is expressed at lower levels in the atherosusceptible regions of the inner aortic arch and aortorenal branches than in other regions. Interestingly, the TAK1 gene contains a highly conserved 406 miR-10a binding site in the 3'-UTR by which miR-10a can negatively regulate TAK1 expression. Such 407 regulation by miR-10a directly mediates the expression of NF-kB p65 and contributes to the regulation 408 409 of proinflammatory endothelial phenotypes in atherosusceptible regions in vivo [121]. It is worth noting that a single miRNA can target multiple mRNAs, suggesting that miRNAs that regulate TAK1
may target other genes, causing off-target effects.

412

413 5.2.2 CRISPR/Cas-mediated gene modification

414 Clustered regularly interspaced short palindromic repeats (CRISPR) is a repetitive DNA sequence in the genome of prokaryotic organisms that is derived from DNA fragments of bacteriophages that have 415 previously infected prokaryotes. It can detect and destroy DNA from similar bacteriophages during 416 subsequent infections, generating a unique immune response to protect against foreign invasion [122]. 417 418 CRISPR-associated protein (Cas) is an enzyme that uses guide RNA to recognize and cleave target 419 strands of DNA that are complementary to the guide RNA [123]. As CRISPR/Cas-based gene editing 420 technology has become more established, it is being widely used to knock out genes completely and 421 permanently by targeting gene loci, thus achieving stable and persistent gene editing. These engineered 422 nucleases generate a double-strand DNA break at the targeted genome locus. The break activates repair 423 through error-prone nonhomologous end joining (NHEJ) or homology-directed repair (HDR). In the absence of a template, NHEJ is activated, resulting in insertions and/or deletions that disrupt the target 424 425 loci. In the presence of a donor template with homology to the targeted locus, the HDR pathway is 426 initiated, allowing for precise mutations to be made [124].

427 Although CRISPR/Cas-based gene editing has not been used extensively as a therapeutic measure 428 for the treatment of pathological angiogenesis, it has been increasingly used in studies to understand 429 the role of TAK1 in various disease contexts. In a study of the role of TAK1 in pneumoconiosis, CRISPR technology was used to generate TAK1 knockout mice via lentiviral vectors expressing 430 CRISPR/Cas9 components. Li et al. confirmed that TAK1 knockout in mice significantly reduced 431 fibrotic nodule formation in the lung tissues after silica exposure [125]. Morioka et al. also showed 432 that the endothelial-specific deletion of TAK1 by CRISPR/Cas9 editing caused increased cell death 433 and vessel regression at embryonic day 10.5 (E10.5), eventually leading to embryo death, which made 434 it difficult to breed endothelial-specific TAK1 knockout mice [11]. CRISPR/Cas-based gene editing 435 has been increasingly studied in the context of manipulating the expression of specific genes in 436 437 pathological angiogenesis. Huang et al. used AAV1-mediated CRISPR/Cas9 editing to target the 438 genomic VEGFR2 locus, resulting in abrogation of angiogenesis in a mouse model of oxygen-induced 439 retinopathy and laser-induced choroidal neovascularization [126]. Moreover, depletion of ONECUT homeobox 2, a highly expressed gene in ovarian cancer tissues, by CRISPR/Cas9 editing remarkably 440 suppressed the expression of several proangiogenic growth factors, such as VEGFA, HGF, and HIF-441 1α, and the activation of Akt/ERK pathways, thus attenuating ovarian cancer progression [127]. With 442 the great advantages of CRISPR/Cas-based gene editing, research has rapidly moved to clinical study. 443

In fact, the latest clinical study using CRISPR/Cas9 editing to design immune cells with enhanced abilities to seek and attack tumors has shown promise in treating some cancers without causing any significant side effects [128]. It is worth noting that the long-term efficacy and safety of CRISPR/Casbased therapy remains unclear. Nevertheless, the rapid developments in modified CRISPR technology have validated its efficacy and safety, providing a new path for the clinical study of gene editing to treat pathological angiogenesis.

450

451 5.3 Potential adverse effects on TAK1 inhibition

Given the pleiotropic nature of TAK1 gene, we can observe diverse roles of TAK1 in multiple 452 physiological activities such as inflammation, immune responses, neural and vascular development. 453 However, this also brings additional risks of undesired side effects when targeted to inhibit its kinase 454 activity. So far, such undesired side effects of TAK1 inhibition either by gene knockout or 455 pharmaceutical inhibitors have not been clinically studied. Nevertheless, a number of studies have 456 suggested that such adverse effects have been observed in various in vitro and in vivo models. For 457 instance, a conditional TAK1 knockout in parenchymal cells of mice liver caused hepatocyte dysplasia 458 and liver carcinogenesis with spontaneous hepatocyte apoptosis and cholangiocytes fibrosis [129,130]. 459 460 Moreover, a study showed that 5Z-7-oxozeaenol can attenuate inflammation and fibrosis in experimental rats with silica-induced pneumoconiosis. However, cytotoxicity in primary lung 461 462 fibroblasts of healthy rats was detected, suggesting that 5Z-7-oxozeaenol may be toxic during the 463 treatment of pneumoconiosis [131]. Similar cytotoxic effects of 5Z-7-oxozeaenol were observed on SK-N-AS and IMR-32 cells at a relatively high dose during the treatment of neuroblastoma [132]. In 464 retinal pigment epithelial cells, TAK1 inhibition led to accelerated cellular senescence, decreased cell 465 proliferation and increased senescence-associated β-galactosidase expression [133]. Selective TAK1 466 467 inhibitor such as Takinib has also demonstrated a significant amount of synoviocyte death at 48 hours when used for the treatment of arthritis in type II collagen induced arthritis mice [134]. These findings 468 469 unarguably suggest that more work is needed on comprehending potential adverse effects of TAK1 470 inhibition. Regardless, TAK1 is still an immensely attractive molecular target for small molecule interventions against diseases, including cancer and inflammation- and angiogenesis-related diseases. 471

472 473

6. Conclusions and future perspectives

TAK1 is an important mediator of multiple signaling pathways that is involved in a variety of 474 pathophysiological processes, including inflammation and the responses to hypoxia and oxidative 475 stress. Increasing evidence indicates that these TAK1-mediated processes clearly participate in 476 angiogenesis-related disorders, such as tumor angiogenesis and retinal neovascularization. 477 Pharmacological inhibitors and genetic approaches for targeting TAK1 have been widely studied in 478 various cancers, such as breast, colon and cervical cancers. Inhibition of TAK1 and its downstream 479 signaling are also effective strategies for inducing the apoptosis of cancer cells and enhancing the 480 chemotherapeutic efficacy of TAK1 inhibitors by regulating the inflammatory and angiogenic 481 482 processes in tumors. However, precisely how TAK1 is involved in regulating angiogenesis and related diseases and the crosstalk between TAK1 and downstream signaling pathways under different conditions remain to be clarified. Nevertheless, TAK1 is a potential therapeutic target that needs to be further studied to provide an alternative to current treatment for pathological angiogenesis.

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- 492

493 **Competing Interests**

- 494 The authors have declared that no competing interest exists.
- 495

496 Author contributions

- 497 Conceptualization- L.Z., S.L., J-H.W., G-S.L. Writing (Original Draft)- L.Z., S.L., J-H.W., G-S.L.
- 498 Writing (Review & Editing)- L.T., G.J.D. Project Administration- J-H.W., G-S.L. Funding
- 499 Acquisition- L.T., G-S.L. Project Supervision- G-S.L.
- 500

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Table

Table 1 Summary of pharmacological inhibition of TAK1

	Inhibitor	Structuro	М.	Origin	Solubil	Commonts	Pre-clinical	Reference
	IIIIIDIUUI	Structure	W.	Origin	ity	Comments	research	S
	5Z-7- Oxozeaenol	H ₃ CO	362. 37	Natural product of fungal/ resorcylic acid lactones	DMSO: >10 mg/mL	 Competitive inhibitor of ATP binding to TAK1 in irreversible manner. Inhibits the catalytic activity of TAK1. TAK-Inhibitory concentration (IC50) = 8.1nM Potently inhibits 50 other kinases. Commercially available for research purposes. 	Tumor suppression (triple negative breast cancer, melanoma), Rheumatoid arthritis	[135- 139,86]
Common chemical inhibitors	NG25 Trihydrochlo ride	F ₃ C N H ₃ C H ₃	646. 96	Synthetic compound	H ₂ O: 5 mg/mL	 Inhibits by binding to ATP binding pocket of TAK1 when TAK1 is in DFG motif "out" or inactive conformation. Also inhibits MAP4K2. 	Tumor suppression (breast cancer, colorectal cancer), cerebral injury	[140,141,1 10,96]
	Takinib	H H H H H H H H H H	322. 36	Amino- benzimidazole	DMSO: 2 mg/mL	 Competitive inhibitor of TAK1. Binds to DFG "in" conformation of ATP binding site of TAK1 Prolongs the rate limiting step of TAK1 activation i.e., prolongs time for TAK1 	Tumor suppression (melanoma), arthritis	[142,95]

						 autophosphorylation. TAK1(IC50=9.5nM) Also inhibits other 5 other kinases.
	LYTAK1	$\begin{array}{c} 0 \\ H_{3}C \\ H$	729. 61	N/A	H ₂ O: 1 mg/mL	 Only known orally active TAK1 inhibitor. Blocks TAK1 phosphorylation at Thr-184/187. Great cytotoxic activity
	Fisetin	НО О ОН О ОН О ОН	286. 24	Natural flavonol	DMSO: ≥50 mg/mL	 Under research Attenuates TAK1 and TAB1 interaction by Dose-dependent phosphorylation inhibition.
Uncommo n chemical inhibitors	Gamma- tocotrienol	HO H ₃ C H ₃ C CH ₃ C H ₃ C CH	410. 63	Lipid-soluble isomers of the essential micronutrient vitamin E	Neat	• Under research Tumor [149]
	Tanshinone IIA	H ₃ C CH ₃	294. 34	Root of Salvia miltiorrhiza Bunge (Chinese Traditional Medicine)	Methan ol: 5 mg/mL	 Reduce TAK1 phosphorylation Under research Lipopolysacchari de induced inflammatory modulation (Atherosclerosis) [150]
Chinese Natural inhibitors	Rubiaceae- type cyclopeptides (RAs)	N/A	N/A	Rubia plant cyclopeptides	N/A	Binds to ATP binding pocket to interrupt TAK1-TAB2 interaction. Inflammatory modulation, tumor suppression, [151]

						angiogenesis inhibition	
Sesamin	N/A	N/A	Sesame Seed	N/A	• Under research	Tumor suppression, prevention of heart failure	[152]
Pinitol	N/A	N/A	3-O-methyl- chiroinositol	N/A	• Under research	Inflammatory modulation, prevention of diabetic complication	[153]
Gambogic acid	N/A	N/A	A Xanthone derived from the resin of the <i>Garcinia</i> hanburyi	N/A	• Under research	Tumor suppression	[154]
Celastrol	N/A	N/A	A Quinone ethide triterpene from <i>Tripterygium</i> <i>wilfordii</i> (Thunder god vine plant)	N/A	• Under research	Tumor suppression, Gastric cancer	[155,156]



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982 Figure 1 Schematic illustration of the domain structures of human TAK1 and TABs. The kinase activity of TAK1 is mediated by binding

983 interactions with TAB1 and its homologs TAB2/3. TAB1 binds to the N-terminal kinase domain of TAK1, whereas the homologs TAB2 and
984 TAB3 bind to the C-terminal region, resulting in the activation of TAK1 catalytic activity.



Figure 2 Interaction between TAK1 and TABs. (A) Proinflammatory ligands bind to IL-1R, TGF- β receptor (TGFBR) and bone morphogenetic protein receptor (BMPR) to trigger interaction with TAK-binding protein 1 (TAB1) and further activate TAK1. Activated TAK1 phosphorylates IKK and MKK/p38 MAPK/JNK, which further activates NF- κ B and AP-1. (B) Proinflammatory ligands bind to IL-1R, Toll-like receptor (TLR) and TNF receptor (TNFR). All these interactions trigger the strong interaction of TAB2/3 with K63-linked polyubiquitin chains to activate TAK1, which subsequently phosphorylates IKK and MKK/p38 MAPK/JNK to activate NF- κ B and AP-1, ultimately regulating inflammation, proliferation and angiogenesis processes. String of beads: polyubiquitination. Created with BioRender.com.



Figure 3 Activation of TAK1 by injury and inflammation. Engagement of agonist with TNF receptor (TNFR) during inflammation and injury. The ubiquitin complex containing TRADD and TRAF activates TAK1, which phosphorylates IKK and MAPKK. IKK phosphorylates NF- κ B. MAPKK phosphorylates JNK and p38 MAPK. Both IKK and MAPKK increase the expression of IL-6, MCP-1, ICAM-1, MMP-9, COX2 and several other chemokines. The binding of an agonist to TNFR during inflammation and injury also activates caspase 8, which leads to cellular apoptosis. Activation of TAK1 via the RIP1/RIP2 complex leads to cellular necroptosis. Created with BioRender.com.



Figure 4 Activation of TAK1 by hypoxia. Hypoxia activates TAK1 via the stimulation of CaMK2, AMPK and Ubc13-XIAP. Activated TAK1
 phosphorylates IKK and MAPKs, which further triggers the transcriptional activation of NF-κB and AP-1, leading to increased expression of
 various cytokines that contribute to cellular survival and angiogenesis. Created with BioRender.com.



Figure 5 TAK1 participates in redox balance. Activation of TAK1 prevents ROS accumulation, protects against ROS-induced apoptosis and enhances angiogenesis. TAK1 maintains ROS at levels that promote angiogenesis by activating Nox2 and upregulating endogenous antioxidants (such as NRF2 and SOD2). When TAK1 is active and the ROS level is low, trigger NF-κB transcriptional activation is triggered, leading to increased expression of angiogenic and antiapoptotic proteins, thereby promoting angiogenesis and inhibiting ROS-induced apoptosis. Created with BioRender.com.