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NF- κ B involvement in Glaucoma-associated neuroinflammation: Focus on glial cells

Francesca Veglianti^{1#}, Mauro Di Vito Nolfi^{1#}, Irene Flati^{1,2}, Francesca Dall'Aglia¹, Fiorenza Di Giovanni¹, Daniela Verzella¹, Daria Capece¹, Adriano Angelucci¹, Edoardo Alesse¹, Francesca Zazzeroni¹, Davide Vecchiotti^{1,*}

¹Department of biotechnological and applied clinical sciences, University of L'Aquila, L'Aquila, Italy

² Department of Molecular Medicine, Sapienza University of Rome, Rome, Italy

*Correspondence: davide.vecchiotti@univaq.it (Davide Vecchiotti)

These authors contributed equally.

Abstract

Glaucoma is a complex neurodegenerative disease characterized by the progressive loss of retinal ganglion cells (RGCs) and optic nerve damage. Both mechanical and vascular factors are believed to contribute to the etiology of glaucoma. However, the underlying pathogenic mechanisms are not yet fully understood. In this article, although it's a single component of a multifactorial condition, we argue that neuroinflammation is a significant factor in glaucoma pathogenesis. Glaucoma, at present, is recognized as a neurodegenerative disorder sharing common neuroinflammatory mechanisms with classical neurodegenerative diseases. The involvement of classical immune signaling pathways, such as TLRs and NF- κ B, as well as proinflammatory cytokines like TNF- α , aligns glaucoma with other neurodegenerative diseases where inflammation is pivotal (e.g. Parkinson's and Alzheimer's diseases). As such, glaucoma should be considered not only an ocular pressure disorder but also a neurodegenerative condition with a strong immune component. This perspective opens new avenues for novel therapeutic intervention, including the target of glial cells or modulators of inflammatory signaling.

However, the complexity of microglial phenotypes and the timing of their activation relative to astrocytes remain areas that require further clarification. The current M1/M2 paradigm is



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acknowledged as overly simplistic, highlighting the need for more refined and nuanced models. Although oxidative stress and other inter-connected signaling such as STAT3 are involved in the pathogenesis of glaucoma, here, we focus on the role of NF- κ B signaling pathway within the glaucomatous condition with a special focus on the main characters fostering the neuroinflammation.

Keywords

Glaucoma 1; Neuroinflammation 2; Microglia 3; NF- κ B 4; Müller Cells 5; Astroglia 6;

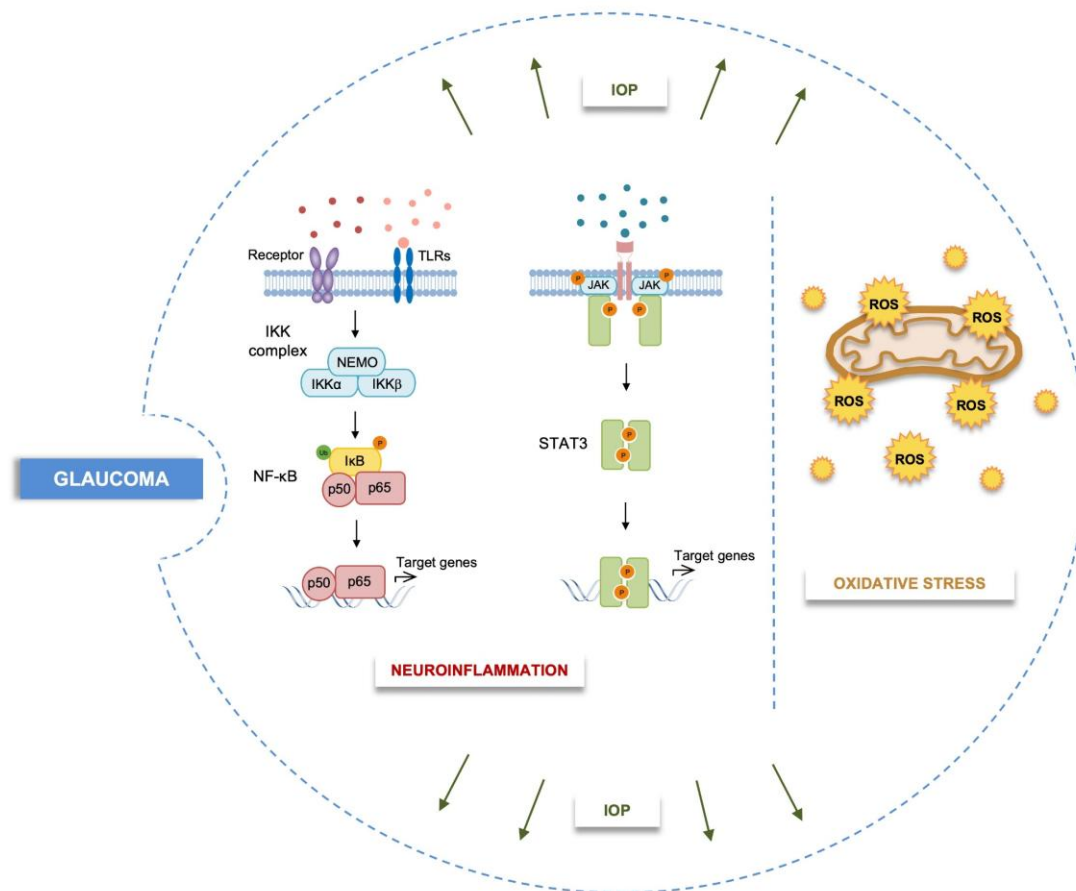
1. Glaucomatous disease and neuroinflammation

Glaucoma is a chronic neurodegenerative disease characterized by the progressive degeneration of retinal ganglion cells (RGCs), loss of synapses, optic nerve damage, and corresponding visual field defects (1). Elevated intraocular pressure (IOP) is a well-known risk factor; however glaucomatous neurodegeneration also occurs in patients with normal IOP, a condition commonly referred to as normal tension glaucoma (NTG). The evidence that NTG patients can still develop glaucoma, points to mechanisms beyond mechanical pressure, with neuroinflammatory pathways emerging as central contributors (Figure 1).

Genetic factors have been identified as prodromal in the development of Glaucomatous dysfunction (e.g. MYOC, OPTN CYP1B1). Furthermore, it has been demonstrated that key genes contribute to the onset of the neuroinflammatory state associated with the disease. Gain-of-function mutations in the pro-inflammatory gene *TBK1* play a crucial role in driving neuroinflammation in glaucoma pathogenesis, independent of IOP elevation (2–4). Moreover, genome-wide association studies (GWAS) have implicated the ATP-binding cassette (ABC), subfamily A member 1 (ABCA1) gene in primary open angle glaucoma (POAG), the most commonly diagnosed form of glaucomatous

dysfunction (5, 6). ABCA1 is expressed in glaucoma-relevant ocular tissues such as the iris, ciliary body, retina, optic nerve head, optic nerve, and trabecular meshwork. It has been demonstrated that ABCA1 may regulate neuroinflammation and neurodegeneration in several murine models, and some studies have shown its involvement in retinal pathogenesis and POAG (7, 8). ABCA1 has been implicated in retinal inflammation and RGC apoptosis, and has also been shown to play a role in IOP regulation through modulation of aqueous humor dynamics (9, 10).

The inflammatory process involves the activation of resident glial cells, the production of proinflammatory cytokines and chemokines, and the infiltration of peripheral cells into the central nervous system due to the disruption of the blood-brain barrier. Initially, this response serves as a protective mechanism against various pathogens (11). Glial cells continuously monitor the retina microenvironment and immediately respond to even minor changes to maintain tissue homeostasis. Macroglia (astrocytes and Müller cells), and microglia are resident immune cells that perform innate immune functions within the retina and optic nerve (12). In the healthy undamaged retina, these cells provide nutritional and structural support, participate in metabolism, and regulate homeostasis. They also coordinate with each other to regulate neuronal activity through phagocytosis and the secretion of inflammatory cytokines and neurotrophic factors (13). Functional dysfunction and neuronal damage occur when this homeostasis is perturbed and glial cells are unable to return to their resting state, as demonstrated by morphological, functional and molecular deregulations. This process is commonly referred as “reactive gliosis” and is marked by specific morphological features such as hypertrophic cell bodies and thickened processes (14–16). The dynamic interplay between microglia and macroglia further complicates the inflammatory milieu, potentially creating feedback loops that can either exacerbate or resolve damage (17). Overall this findings strongly suggest that glial cells are a driving force in the retina inflammatory process and a deeper understanding of its role within the glaucomatous disfunction could potentially support the identification of new therapeutic approaches.



80 Figure 1. Schematic representation of multifactorial pathogenic processes involved in the glaucoma. Neuroinflammation, oxidative
81 stress and biomechanical dysfunction are key factors in the disease onset and progression.

82 2. Macroglia and microglia: the main characters

83 Recent findings indicate that glaucomatous optic neuropathy is a neurodegenerative disorder that
84 shares neuroinflammatory mechanisms with classical neurodegenerative diseases, highlighting the role
85 of glial cell-mediated neuroinflammation in the glaucoma's pathophysiology (18, 19).

86 Glial activation is crucial for restoring tissue homeostasis, facilitating repair, and providing
87 neuroprotection in the central nervous system (CNS), where astrocytes and microglia mediate the
88 innate immune response (20). The strong interplay between neurons and glial cells suggests
89 that a dysfunction in either can result in harmful processes that may disrupt neuron-glia

90 communication or jeopardize neuronal health (21, 22). The retina is an immune-privileged site where
91 the impact of the systemic immune system is strictly regulate (23). This immune privilege helps
92 maintain retinal homeostasis and protects it from persistent and detrimental inflammation.

93 At least three types of glial cells exist in the mammalian retina: Müller cells and astrocytes,
94 collectively referred to as macroglia, and microglia. Each of these cell types play distinct roles in
95 supporting retinal function and responding to injury or disease by carefully modulating immune
96 responses. Astrocytes and microglia are now recognized to display various activation phenotypes, with
97 their shift from protective to potentially harmful roles closely associated with the severity and
98 progression of neurodegeneration (24).

99 Microglia are the resident immune cells of the CNS including the retina where they are found in
100 several layers such as nerve fiber layer (NFL), ganglion cell layer (GCL), inner plexiform layer (IPL)
101 and outer plexiform layer (OPL), acting as neuropathological sensors and as a first line of defense
102 against injury. Microglia originate from primitive erythromyeloid progenitors that enter the CNS via
103 the primitive bloodstream during early embryonic development.

104 Microglial cells can defend neural tissue from harmful stimuli like pathogen-associated molecular
105 patterns (PAMPs) and damage-associated molecular patterns (DAMPs) by detecting them through
106 their specialized receptors, including TLRs (25, 26). In response to neural damage, microglia undergo
107 morphological changes, proliferate, migrate, and release inflammatory cytokines such as IL-1 β , IL-6,
108 IL-12, IL-18, IL-23, TNF- α , CCL2, CXCL10, reactive oxygen species (ROS) and matrix
109 metalloproteinases (MMP9 and MMP3), which can exacerbate neuronal loss and dysfunction (27, 28).
110 In the context of glaucoma, reactive microglial cells initially engage in the phagocytosis of debris from
111 damaged RGCs, thereby contributing to maintain of a toxin-free microenvironment. Additionally,

these cells release neurotrophic factors, such as brain-derived neurotrophic factor (BDNF) and ciliary neurotrophic factor (CNTF), which provide neuroprotection and promote tissue regeneration (29).

A rapid increase of IOP is able to induce changes in microglial morphology, such as retraction of processes, enlargement of the soma, and increased expression of activation markers like CD68, in just 60 minutes (30). Activated microglia produce pro-inflammatory and cytotoxic molecules, including complement factors, nitric oxide (NO), TNF- α , IL-6, and increased MHC I and II expression, all of which contribute to neurotoxicity and propagation of neuroinflammation in glaucoma. TNF- α is reported to increase in glaucoma patients and are associated with microglial activation and neuronal damage (31). Gregory et colleagues observed that in glaucomatous DBA/2J mice, high levels of TNF- α and an increased expression of membrane-bound form of Fas ligand (mFasL) on microglia trigger apoptosis of RGCs via their Fas receptor. Pharmacological inhibition or genetic deletion of TNF- α or Fas ligand reduces microglial activation and protects RGCs (32, 33).

Indeed, microglial activation is one of the earliest events in glaucoma, preceding RGC loss, as demonstrated in DBA/2J animal model, where early microglial alterations correlate with the extent of neurodegeneration. Treatments that inhibit microglial activation, like minocycline, have been shown to reduce RGC death in this model, underscoring the critical role of activated microglia in the pathogenesis of glaucoma (16, 34, 35). Overall, these findings highlight the role of microglial activation in glaucomatous neurodegeneration. Within this context, it is now widely accepted that this microglial-activation status into the CNS can shift among a plethora of phenotypes (M0-M1-M2a-M2b-M2c-M2d) that may exacerbate the neurodegeneration or rescue the stress-induced damage with a strong neuroprotective effect. These microglia subsets are conventionally divided into M0 (resting), M1-like (Classical activated or Pro-inflammatory) induced by LPS (lipopolysaccharide), IFN γ

134 (interferon-gamma), and TNF- α , and M2-like (Alternative activated or Anti-inflammatory) induced by
135 IL-4, IL-10, IL-13 (36, 37). Other classifications take into account the morphology and activation
136 states of microglia (38, 39) and transcriptomics-based approach (40, 41) thus showing a more complex
137 but more specific description of the different phenotypes.

138 Retinal astrocytes mainly reside in the NFL and GCL layers and are classically identified by the
139 expression of Glial fibrillary acidic protein (GFAP), vimentin, and nestin. Paired box 2 (Pax2) and
140 SRY-Box Transcription Factor 2 (Sox2) instead are used as nuclear markers of astrocytes which are
141 present as well in the optic nerve (13, 42–48).

142 In order to counteract glaucomatous dysfunction, the morphological structure of astrocytes,
143 characterized by long processes that envelop RGCs, play a structural role in counteracting IOP-related
144 biomechanical stress and maintaining retina homeostasis. This is achieved through their ability to
145 remove cellular debris and increased phagocytic activity in glaucoma onset and progression (49).

146 Although astrocytes continue to remain enigmatic in term of retinal functions, they share common
147 features with microglia such as plasticity and ability to change their phenotype toward either
148 neurodegenerative or neuroprotective states under pathological conditions, as well as reactivity and
149 ability to migrates to injury sites as demonstrated by several studies (50–53). Morphological changes
150 in astroglia populations are classically recognized as key markers of reactive astrogliosis, that is also
151 characterized by a strong upregulation of GFAP (54, 55). Among the first to link glaucomatous
152 conditions to the inflammatory insults, Stevens et colleague demonstrated that retinal astrocytes
153 become reactive and increase C1q expression in RGCs contributing to the glaucoma inflammatory-
154 related degeneration and vision loss (54).

155 Astrocytes and microglia do not function independently in healthy or pathological tissue; rather, they

156 team up to drive and intensify neuroinflammation (20, 56). A mutual communication between
157 microglia and astrocytes has been well demonstrated during onset, progression and amplification of
158 the glaucomatous neuroinflammation (20, 56–58). This teamwork arise from both direct molecular
159 interactions and their shared responses to environmental cues. In glaucoma, stress from increased IOP
160 and damage to RGCs seem to be the key triggers that synchronize the astrocyte-microglia partnership.
161 In addition to their close relationship, the molecules produced by these glial cells can also recruit
162 immune cells from the bloodstream, further fueling neuroinflammation (15).

163 Astrocytes and microglia communicate with each other through different mechanisms. First, both cell
164 types can sense and respond to the same signals at the same time. They have similar immune receptors
165 that enable them to communicate and work together during inflammation. These cells can detect stress
166 in other cells by recognizing damage signals. Both cells respond to ATP, which is released from
167 stressed or dying RGCs. This ATP triggers inflammation through purinergic signaling. Besides
168 purinergic receptors (e.g. ionotropic P2X7R), astrocytes and microglia also express pattern recognition
169 receptors (PRR) like toll-like receptors (TLRs) and nucleotide-binding oligomerization domain-like
170 receptors (NLRs). When these receptors are activated by damage signals, they can quickly trigger an
171 inflammatory response (59).

172 Astrocytes and microglia crosstalk can either increase or decrease their responses. They do this by
173 releasing molecules like cytokines, chemokines, or complement molecules. Astrocytes mainly produce
174 chemokines, which bind to receptors on microglia and help microglia migration. Cytokines from
175 astrocytes can also assist microglia in removing synapses. Complement components, like C3 from
176 astrocytes, can support microglia with their eating function. Astrocytes and microglia also keep their
177 numbers balanced. Microglia can control astrocyte numbers by phagocytosing them in the retina.

178 Communication between astrocytes and microglia can also affect their behavior. Congruently,
179 microglia can induce astrocytes to become harmful; thus promoting inflammation (27). In glaucoma,
180 microglia can shift astrocytes from a protective anti-inflammatory A2 state to a harmful pro-
181 inflammatory A1 state (27, 60). Interleukin-1alpha (IL-1 α), tumor necrosis factor alpha (TNF- α), and
182 C1q released by microglia support this phenotypic change. Accordingly, the genetic knockout of IL-1,
183 TNF- α and C1q in mice model with IOP show a reduction of harmful change in retinal astrocytes (60).
184 Müller cells are a type of glial cell found only in the retina, stretching throughout its full thickness.
185 Müller cells are strategically positioned transversely across all nuclear and plexiform layers, with their
186 endfeet making contact with the vitreous cavity and their microvilli extending into the subretinal space.
187 Within the IPL and OPL, these processes envelop synapses and form connections with blood vessels
188 (61). These cells play a crucial role in directing light to photoreceptors and in mitigating mechanical
189 stress within the retinal tissue. They are classically identified by the expression of Glutamine
190 Synthetase (GS), cellular retinaldehyde acid-binding Protein (CRALBP), glutamate-aspartate
191 transporter (GLAST), S100b, and vimentin. Müller cell's soma is located in the inner nuclear layer
192 (INL), with its processes extending in both directions towards the inner and outer retina (62). Although
193 classically used as a marker of astrocytes, GFAP has been shown to be highly expressed by reactive
194 Müller cells in glaucoma patients (63).

195 In detail, Müller cells play a crucial role in the rapid clearance of excess glutamate from the
196 extracellular environment via amino acid transporters, particularly excitatory amino acid transporters.
197 This process is essential for maintaining low glutamate levels, thereby preventing excitotoxicity. The
198 conversion of glutamate to glutamine is facilitated by the enzyme glutamine synthetase (GS), which is
199 specifically expressed by Müller cells. Glutamine then acts as a precursor for the synthesis of glutamate

200 in neurons. In glaucoma, both in human patients and in animal models, an increase in the expression
201 of glutamine within Müller glia has been observed, suggesting a heightened activation of the
202 glutamate–glutamine cycle. It is proposed that the rise in glutamine concentrations within Müller glia
203 could be attributed to a diminished need for glutamine in injured RGCs. Beyond their function in
204 clearing excess glutamate from the synaptic cleft, Müller glia are also capable of metabolizing
205 glutamate as a substrate (64).

206 This support is vital for the survival of photoreceptors and neurons, as it enables the efficient uptake
207 and recycling of neurotransmitters and their precursors, thereby ensuring precise neuronal
208 communication (65).

209 Under stress conditions, Müller cells show cellular hypertrophy, which involves the thickening of both
210 their somas and processes. This condition is also associated with increased cellular proliferation and
211 an upregulation of cytoskeletal proteins, such as vimentin and a reduction in the expression of GS (66).

212 Similar to microglia and astrocytes, Müller cells initially display neuroprotective phenotypes through
213 the release of factors such as pigment epithelium-derived factor (PEDF) or VEGF. However, they can
214 also fuel neuroinflammation by releasing IL-1 β , TNF- α , IL-6, Complement protein C1q and NO
215 expression that induce RGC loss when overstimulated. In glaucoma, Müller cells, can play a role in
216 neuroinflammation by expressing monocyte chemoattractant protein-1 (MCP-1), a chemokine that
217 recruits monocytes and macrophages. If chronic and not promptly resolved, this recruitment leads to a
218 sustained inflammatory response that contributes to RGCs death (11, 67). Müller glia cells activated
219 in response to glaucoma play a crucial role in preserving the integrity and homeostasis of neuronal
220 tissue. This process involves the upregulation of receptors that are crucial for neuronal growth and
221 survival, enhanced uptake of extracellular glutamate to shield RGCs from excitotoxic damage, and the

release of nerve growth factor (NGF) (68).

Although Müller glia are more numerous than astrocytes in the retinal environment, both macroglial subtypes exhibit a range of shared functions alongside distinct attributes, many of which remain to be elucidated. Moreover, timing and site of macroglial activation are crucial to induce different activation states and to determinate the beneficial or detrimental effect thus making it more difficult to choose an appropriate therapeutic approach (69, 70). It is imperative for forthcoming research to explore both the unique and common roles of astrocytes and Müller glia. Furthermore, understanding their potential collaborative interactions is vital for gaining a comprehensive insight into their functions.

Generally, in the glaucomatous retina, proinflammatory cytokines plays a crucial role in both microglia and macroglia. Although microglia act as primary responder to neural damage, initiating an inflammatory response that leads to neuronal loss, macroglia provide essential structural and metabolic support to neurons but can become reactive under chronic stress, contributing to neuroinflammation and to the RGC degeneration. Within this context the involvement of NF- κ B signaling pathway attracts attention being considered a central hub for neuroinflammation as well as for neuroprotection.

3. NF- κ B Pathway in the glial retina cells

The nuclear factor kappa-light-chain-enhancer of activated B cells (NF- κ B) transcription factor proteins are pivotal regulators of cell growth, differentiation, apoptosis (71, 72), immune responses, inflammation and tumor initiation and progression, both in vertebrates and invertebrates (73, 74). NF- κ B was initially identified as a nuclear factor in B cells that binds the enhancer element controlling immunoglobulin κ light chain expression (75). Later, NF- κ B binding sites were identified in the promoter site of genes that encode inflammatory cytokines and chemokines, major histocompatibility

245 complex molecules, cell adhesion molecules, growth factors and regulators of apoptosis in various cell
246 types (76). Various stimuli, like bacterial or viral antigens, cytokines and growth factors (e.g. members
247 of the TNF receptor superfamily or toll-like receptors), can activate NF- κ B, leading to the
248 transcriptional upregulation of inflammatory cytokines. Structurally, Rel/NF- κ B transcription factor
249 family consists of several homo and hetero- dimeric complexes that bind to 9–10 base pair DNA
250 sequences known as κ B sites (77). The NF- κ B family includes five transcription factor proteins - NF-
251 κ B1 (p50/p105), NF- κ B2 (p52/p100), RelA (p65), RelB and c-Rel - which can dimerize to form all
252 possible combination of homo and heterodimers. The prototypical form of NF- κ B is the heterodimeric
253 p50/p65 complex (78). All NF- κ B proteins are characterized by the presence of an N-terminal Rel
254 homology domain (RHD), which is involved in DNA binding, dimerization process, nuclear
255 localization and I κ B inhibitor binding. RelA (p65), RelB, and c-Rel contain a C-terminal
256 transactivation domain (TAD) that enables them to activate target genes. p50 (p105 precursor) and p52
257 (p100precursor) possess a long ankyrin repeat-containing domain (ARD) at their C-terminus instead
258 of the TAD domain, and therefore cannot activate target gene expression as a homodimer (79). The
259 activity of NF- κ B is tightly regulated by its interaction with inhibitory I κ B proteins. In unstimulated
260 cells, NF- κ B dimers are retained in an inactive form in the cytosol through their binding to I κ B proteins
261 All I κ Bs recognize NF- κ B via their ARD. Family members of I κ B can be divided into three subgroups
262 based on their function and structure: 1) classical I κ B proteins (I κ B α , I κ B β , and I κ B ϵ); 2) NF- κ B
263 precursor proteins (p100 and p105); and 3) the nuclear I κ Bs (I κ B ζ , Bcl-3, and I κ BNS) (80, 81). I κ B α ,
264 I κ B β , and I κ B ϵ are the main inhibitors of NF- κ B dimers (p50/p65 and p65/p65) activation. The ARD
265 of these proteins includes six repetitions of ankyrin; the SRD region contains conserved serine-rich
266 sites that are phosphorylated by the I κ B kinase complex (IKK), and lysine residues that are
267 polyubiquitinated before proteolytic degradation (82). p105 and p100 play a dual role in cells, are both
268 precursors required for the maturation of p50 and p52 subunits of NF- κ B and are both nonspecific
269 inhibitors of NF- κ B. I κ B ζ , Bcl-3 and I κ BNS have nuclear localization signal (NLS) and are therefore

270 in the nucleus. Bcl-3 interacts with homodimers of NF- κ B p50 and p52, and can act either as an
271 activator or as a repressor of transcription (83).

272 Various exogenous signals, such as proinflammatory cytokines (e.g. IL-1 β and TNF- α),
273 constituents of cellular pathogens (e.g. LPS) and radiation can induce activation of NF- κ B. These
274 signals induce the activation of a specific I κ B kinase termed IKK. There are two pathways leading to
275 the activation of NF- κ B: the canonical and noncanonical pathways or the classical and alternative
276 pathways, respectively. Pro-inflammatory cytokines (IL-1 and TNF α), components of bacterial cell
277 walls (LPS), viruses and agents that damage DNA induce the canonical pathway through activation of
278 the IKK $\alpha/\beta/\gamma$ complex present into the cytoplasm. The IKK complex consists of two catalytic subunits,
279 IKK α and IKK β , as well as a regulatory subunit, NEMO (IKK γ). The IKK complex consists of a homo-
280 ($\alpha\cdot\alpha$ or $\beta\cdot\beta$) or heterodimer ($\alpha\cdot\beta$) associated stoichiometrically with NEMO (84). The catalytic subunits
281 IKK α and IKK β share about 50% sequence identity; show an N-terminal kinase domain and "leucine
282 zipper" and "basic helix-loop-helix" regions, while IKK γ has a helical structure interrupted by "coiled-
283 coil" structures and C-terminal Leucine-zipper (LZ) motif (85). IKK activation lead to the
284 phosphorylation of two serine residues (Ser-32 and Ser- 36) of I κ B inhibitor. Phosphorylation of I κ B,
285 in turn, leads to Lys48-linked polyubiquitination which targets I κ B for degradation by the 26S
286 proteasome. At this point NF- κ B is free and can translocate into the nucleus (86), where they activate
287 specific target genes expression involved in inflammatory response, cell survival, immune response
288 and apoptosis. The alternative pathway is induced by LT- β and BAFF. Ligand binding to their
289 respective receptors leads to activation of the NF- κ B-inducing kinase (NIK), which phosphorylates
290 and activates an IKK α complex. This complex, in turn, phosphorylates two serine residues adjacent to
291 the ankyrin repeat C-terminal I κ B domain of p100, leading to its partial proteolysis and liberation of
292 the p52/RelB complex; this alternative pathway is primarily involved in the activation of p100/RelB
293 complexes during B-and T-cell organ development (87). IKK activation process is coupled with a
294 negative feedback control mechanism that quickly modulate NF- κ B activity. The genes coding for

some I κ B isoforms show binding-sites for NF- κ B in their promoter, allowing NF- κ B to self-regulated its own activity through transcriptional induction of its inhibitors (88). Genetic alterations targeting components of the NF- κ B family as well as dysfunction within the NF- κ B signaling pathway have been associated with neurodegenerative diseases such as Parkinson disease (PD) and Alzheimer disease (AD) as well as in glaucoma (89). In silico and in vivo studies demonstrated that NF- κ B activation is triggered by α -sinuclein and amyloid- β in PD, AD and glaucoma. In addition, several studies demonstrated that microglia activation and neuroinflammatory process are involved in these neurodegenerative diseases (16, 90). In detail, recent Li et collaborators using transcriptomic approach, identified IGFBPL1 (Insulin-like growth factor-binding protein like protein 1) able to switch molecular signatures of activated microglia via the NF- κ B pathway in mice model of neuropathies (91). These findings suggest that AD, PD and glaucoma could share common mechanistic pathogenic process that could be considered potentially as an attractive target for neurodegenerative disorders (91–95). Although NF- κ B is also involved in IOP regulation through toll-like receptor 4 (TLR4) at level of trabecular meshwork (96), this signaling plays a crucial role in response to IOP and subsequent mechanical stress. Wang *et al.* showed as suppression of TLR4/NF- κ B in activated rat microglia following mechanical injury leads to a reduction of pro-inflammatory cytokines expression (97). Again, astrocytes sense IOP through mechanosensitive ion channels and release ATP (98) which, in turn, mediate microglia inflammatory response via P2X7R/Ca²⁺/NFAT/NF- κ B pathway (18). NF- κ B also regulates the response to S100B protein which is expressed by both astrocytes and microglia. Higher doses of S100B are able to activate astrocytes and microglia promoting the expression of iNOS and pro-inflammatory cytokines via NF- κ B (99, 100). Moreover, rats immunization with S100B activates NF- κ B and cytokines production leading to a glaucomatous condition in autoimmune glaucoma models (101). As previously discussed, microglia and macroglia polarization could shape the neuroinflammatory status of the glaucomatous retina. The existing cross-talks between these glial cells suggests that NF- κ B-regulated pro-inflammatory cytokines produced in astrocytes may also play a role

in modulating the inflammatory responses of microglia (102). Furthermore, a variety of pro-inflammatory cytokines and chemokines, which are associated with glial upregulation in glaucoma, including TNF- α , IFN γ , IL-1, IL-2, IL-12, IL-17, FasL, and iNOS, have been identified as target genes of NF- κ B (Figure 2) (Table 1) (59, 103).

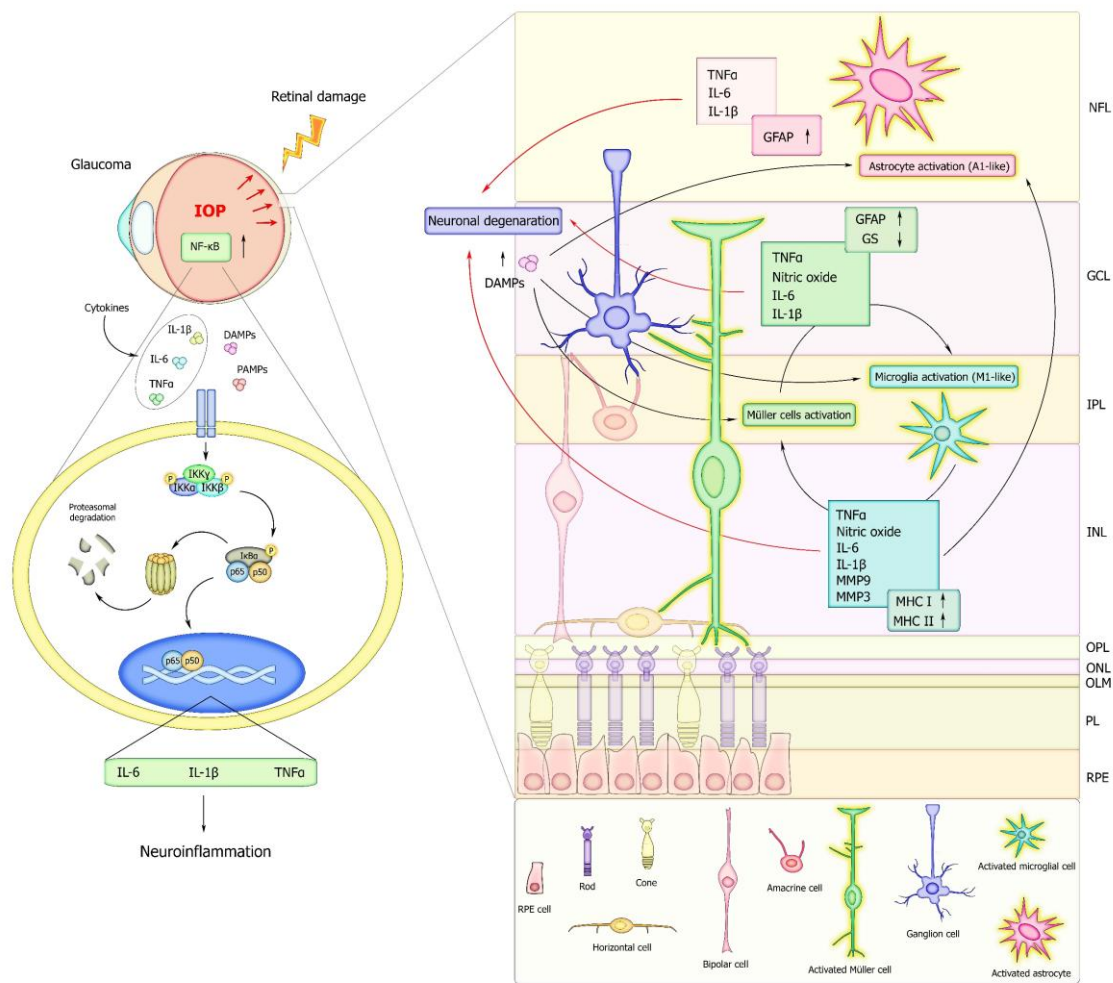


Figure 2. Schematic representation of the NF- κ B canonical pathway involvement in the glaucomatous glial cells. NFL (Nerve Fiber Layer); GCL (Ganglion cell layer); IPL (Inner Plexiform Layer); INL (Inner nuclear layer) OPL (Outer plexiform layer); ONL (Outer nuclear layer); OLM (Outer limiting membrane); PL (Photoreceptor layer); RPE (Retinal pigment epithelium).

Inhibition of NF- κ B activation through the cre/lox-based conditional deletion of astroglial p50 led to a reduction in the production of pro-inflammatory cytokines in experimental mouse models of

glaucoma (102). Notably, in addition to the diminished inflammatory activity of astroglia, alterations in microglial responses were also observed following p65 deletion in GFAP-expressing astroglia. Pro-inflammatory response of microglia was significantly reduced in ocular hypertensive GFAP/p65 mice compared to ocular hypertensive p65^{f/f} controls. These preliminary findings suggest that, similar to the modulation of astrocyte responses by microglia, astrocytes - through NF-κB-regulated mediators - can influence microglial responses, thereby exacerbating neuroinflammation (27).

Recent research focusing on the transcriptomic and proteomic profiling of human donor eyes and animal models affected by glaucoma has identified an early increase in the expression of numerous molecules and pathways involved in inflammatory signaling, such as pattern recognition receptors, TLRs, and NLRs, MyD88, MAPKs, and NF-κB (40, 104, 105). Furthermore, in glaucoma, it is widely accepted that neuroinflammation driven by glial cells involves multiple pathways, such as TNFR signaling, TLR signaling, and the inflammasome, all of which are predominantly regulated by NF-κB. Upon encountering environmental signals, TNFR and TLR signaling pathways initiate the transcriptional priming of the inflammasome through the activation of NF-κB. This activation enhances the expression of inflammasome components and the precursors of cytokines activated by the inflammasome, including IL-1β (106).

In an experimental glaucoma model, the conditional deletion of IκKβ in astroglia expressing GFAP (GFAP/IκKβ) using the cre/lox system led to a reduction in the production of pro-inflammatory cytokines which are transcriptional targets of NF-κB, including TNF-α, IFNγ, IL-1, and IL-2. By inhibiting the NF-κB-regulated inflammatory functions of astroglia, structural and functional protection was afforded to RGCs in eyes with ocular hypertension (102).

Lebrun-Julien and colleagues demonstrated that the activation of the NF-κB signaling pathway, along with the upregulation of NGF expression, leads to the induction of TNF-α expression by Müller cells, thereby contributing to neurotoxicity in vivo (107).

380 Targeting the NF- κ B pathway presents promising therapeutic strategies for mitigating glaucoma-
381 related damage. However, due mainly to the systemic dysfunction related to NF- κ B inhibition, few
382 attempts have been made. Initial approaches aiming at inhibiting the NF- κ B signaling pathway using
383 specific inhibitors could reduce inflammation and subsequently the progression of glaucoma. NF- κ B
384 inhibitors (e.g. Resveratrol, Pycnogenol) are already being explored for their benefits in various other
385 inflammatory and neoplastic diseases, due to their ability to suppress pro-inflammatory cytokines and
386 protect against tissue damage (108, 109). Ouyang and colleagues employed a multifunctional
387 nanosystem that co-delivers NF- κ B inhibitors with other therapeutic agents to enhance treatment
388 efficacy. For instance, a system combining a NF- κ B inhibitor like thalidomide with an activator of the
389 Nrf2 pathway, has shown promise in reducing inflammatory responses and promoting recovery in
390 inflammatory diseases (110). Compounds like parthenolide have demonstrated a strong NF- κ B
391 inhibition along with anti-inflammatory and anti-cancer activities. Such natural compounds may offer
392 a less toxic alternative for the long-term management of inflammatory responses in glaucoma (109).
393 Implementing chronic therapeutic schemes using specific NF- κ B inhibitors could be beneficial in
394 maintaining long-term therapeutic effects without significant adverse reactions. This strategy could
395 potentially halt or slow down disease progression, preserving vision (111). However, although
396 inhibition of NF- κ B represents a promising therapeutical approach, the use of core pathway NF- κ B
397 inhibitors could be detrimental in some contexts. As example, although S100B protein support
398 inflammation in astrocytes and microglia via NF- κ B, the inhibition of this signaling could have
399 negative effect on neurons in which S100B promotes NF- κ B-dependent neuronal survival (112).
400 Overall, these aspects imposing the identification of more specific downstream NF- κ B effectors and
401 the development of highly specific inhibitor molecules. The activation of NF- κ B signaling in Müller
402 cells occurs rapidly following retinal injury, suggesting its importance as an early response mechanism.
403 Moreover, Müller cells respond to IOP through pressure-sensitive transient receptor potential vanilloid
404 4 (TRPV4) that increases TNF- α expression and induces gliosis via JAK2/STAT3/NF- κ B signaling

(113) and statins are able to mitigate gliosis by inhibiting NF- κ B activation. Interestingly, this activation is not autonomous but depends on signals from reactive microglia. This intercellular communication underscores the intricate network of cellular interactions that orchestrate the retinal response to injury. Researchers have explored various approaches to modulate NF- κ B signaling in Müller cells, aiming to enhance the regenerative potential of the retina. Inhibition of NF- κ B signaling, achieved through either pharmacological interventions or genetic knockout techniques, has shown promising results. When combined with the overexpression of *Ascl1*, a transcription factor known to promote neuronal differentiation, NF- κ B inhibition significantly enhances the ability of Müller cells to reprogram into neurons. This finding suggests that NF- κ B signaling acts as a barrier to neurogenic reprogramming and that its suppression can unlock the regenerative potential of Müller cells. The role of NF- κ B in maintaining glial identity and suppressing neurogenic potential is not isolated but involves coordination with other signaling pathways and transcription factors. For instance, NF- κ B interacts with the TGF β /Smad3 pathway, known for its involvement in cell growth and differentiation (114). Additionally, NF- κ B works in concert with transcription factors such as Id proteins, which are important regulators of cell fate decisions. This complex network of interactions ensures the maintenance of glial identity and functions under normal conditions. Following retinal injury, NF- κ B signaling in Müller cells promotes the expression of inflammatory gene networks. This pro-inflammatory response contributes to the overall glial reactivity observed in the damaged retina. By driving these inflammatory and reactive processes, NF- κ B signaling acts as a key regulator that pushes Müller cells towards restoring quiescence rather than undergoing neurogenic reprogramming. This tendency to return to a quiescent state, rather than initiating a regenerative program, is a characteristic feature of mammalian retinal response to injury, contrasting with the robust regenerative capabilities observed in some non-mammalian vertebrates (115). Understanding the role of NF- κ B signaling in Müller cells provides valuable insights into the limitations of mammalian retinal regeneration and offers potential targets for therapeutic interventions. By modulating NF- κ B activity and its associated

pathways, researchers may be able to enhance the regenerative capacity of the mammalian retina, potentially leading to novel treatments for retinal injuries and degenerative diseases. These strategies focusing on NF- κ B may provide a multifaceted approach to managing glaucoma by emphasizing inflammation reduction, which is critical factor in protecting the optic nerve and maintaining vision. The inflammatory cascade initiated by NF- κ B activation in glial cells can have far-reaching effects on the retinal microenvironment. For instance, the release of MMPs by activated glial cells can lead to the remodeling of the extracellular matrix, potentially altering the structural integrity of the retina and optic nerve head. This remodeling process may contribute to the characteristic cupping of the optic disc observed in glaucoma patients. However, NF- κ B activation in glial cells also has potential neuroprotective effects. It induces the upregulation of antioxidant enzymes (superoxide dismutase (SOD) and catalase (CAT)), which help neutralize reactive oxygen species and protect against oxidative damage - and anti-apoptotic factors (Bcl-2, Bcl-xL and GADD45 β), which may help prevent retinal ganglion cell death in glaucoma. NF- κ B activation can also stimulate the production of neurotrophic factors, such as BDNF and CNTF, which support the survival and function of retinal neurons, and modulate glutamate uptake and metabolism, helping to maintain proper neurotransmitter homeostasis and protect against excitotoxicity. This dual nature of NF- κ B signaling in glial cells highlights the complex interplay between neuroprotection and neurodegeneration in glaucoma. The balance between these opposing effects may depend on various factors, including the severity and duration of the glaucomatous insult, the specific glial cell types involved, and the overall retinal microenvironment (Figure 3).

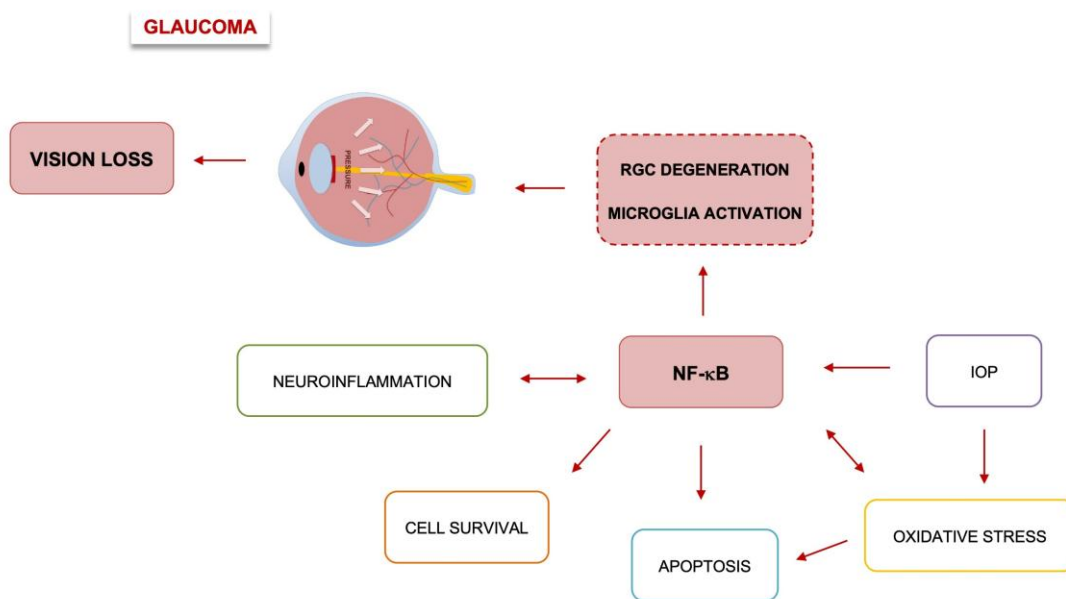


Figure 3. Role of NF- κ B signaling pathway in glaucoma. IOP (Intra ocular pressure), RCG (Retinal ganglion cell).

NF-κB PATHWAY IN GLIAL CELLS						
Functions	Cell type	Cytokines	Receptors	Diseases	Model	Refs
Neuroinflammation	Müller glia, Microglia, Astrocytes	TNF-α, IL-16, IL-6, IL-4, IL-10, CCL2, IL-1β, TGF-β, IFN-γ	CD44, TLR, TNFR1, P2X7R, TLR4, IL-1R	Glaucoma, Retinitis pigmentosa	In vitro, In vivo, Ex vivo	18, 61, 97, 105
Neuroinflammation, Neurodegeneration	Müller glia, Microglia, Astrocytes	TNF-α, IL-1β, HMGB1, CNTF, IL-6, IL-1α, IL-2, IL-10, IL- 12, IL-13, IL-17, IFN-γ	TLR, CNTFR, CD44, TRPV4, TNFR1	Glaucoma, Hypertensive Retinopathy	In vitro, In vivo, Ex vivo	102, 113, 115
Glial reactivity, Neurodegeneration	Müller glia, Microglia	TNF-α, IL-1β, IL-6, OPN	TNFR, IL-1R1, TLR, CD44, RAGE	Glaucoma, Neurodegenerative disease	In vitro, In vivo, Ex vivo	100, 114
Oxidative and Neurotoxic response	Astrocytes	TNF-α, IL-1β	RAGE	Glaucoma, Neurodegenerative disease	In vitro	99, 102
Cell Survival	Microglia, Astrocytes	IL-1α	p75	Glaucoma	In vitro	102

NF-κB PATHWAY IN OTHER RETINAL CELLS						
Functions	Cell type	Cytokines	Receptors	Diseases	Model	Refs
Neuroinflammation, Neurodegeneration	RGC	TNF-α, IL-1β, IL-6	TRPV4, TNFR1	Glaucoma	In vitro, In vivo	113
Neurodegeneration	RGC, Optic nerve	IL-1β	TLR4	Glaucoma	In vitro, Ex vivo	101



Conclusions

Reported evidence suggest that NF- κ B could play a relevant role also in the context of glaucoma pathogenesis, and in particular regulating inflammatory response and cell survival pathways within retinal glial cells. The current prevalent pathologic interpretation of glaucoma supposes the etiologic role of elevated intraocular pressure leading to mechanical stress on retinal tissues. However, the reactive phenotype of resident cells in response to mechanical stress, and in particular of glial cells, could contribute in a clinical relevant manner to the pathogenesis of this disease. Indeed, glial cells - including astrocytes, Müller cells, and microglia - are essential for maintaining retinal homeostasis and supporting neuronal function. These cells form a complex network that provides structural, metabolic, and functional support to retinal neurons, including retinal RGCs.

A clear pathologic description centered on NF- κ B in the context of ocular diseases is not yet completely achievable. One hurdle is represented by the fact that specific genes targeted by NF- κ B can vary depending on the cell type and the nature of the activating stimulus. For example, in Müller cells, NF- κ B activation promotes the release of neuroprotective factors such as BDNF and CNTF, which support the survival and function of retinal neurons, particularly in response to stress or injury. Conversely, in microglia, NF- κ B activation stimulates the production of pro-inflammatory mediators, including cytokines TNF- α and IL-1 β , which while beneficial for pathogen or damaged cells clearance, may promote chronic inflammation and tissue damage if sustained. In addition to cytokine production, activated microglia can also release reactive oxygen species, proteases, and other factors that can modulate the retinal microenvironment and influence the behavior of neighboring cells. Astrocytes also respond to NF- κ B activation. In these cells, NF- κ B signaling modulates the expression of genes involved in antioxidant defense, glutamate uptake, and the production of extracellular matrix



481 components, all critical processes for maintaining the blood-retinal barrier and regulating
482 neurotransmitter levels in the retinal microenvironment. Astrocytes play also a key role in supporting
483 retinal vasculature and regulating blood flow, both essential for maintaining proper oxygenation and
484 nutrient supply to retinal neurons.

485 Understanding these molecular dynamics is crucial for developing targeted therapies aimed at
486 mitigating glaucomatous damage and preserving vision. Research in this area may lead to the
487 development of novel therapeutic approaches that modulate NF- κ B activity in glial cells to promote
488 neuroprotection while minimizing harmful inflammatory responses. Such strategies could include the
489 use of selective NF- κ B inhibitors, targeted delivery of anti-inflammatory agents to retinal glial cells,
490 or the development of therapies that enhance the neuroprotective aspects of NF- κ B signaling while
491 suppressing its pro-inflammatory effects. In particular, reliable methods for cell-type-specific NF- κ B
492 inhibition, which would allow targeting only the disease-involved cells while sparing healthy ones, are
493 particularly needed, enabling in the same time a mitigation of adverse effects, and enhancing
494 therapeutic potential. These approaches could complement existing glaucoma treatments, which
495 primarily focus on reducing intraocular pressure, by addressing the underlying cellular and molecular
496 mechanisms of the disease. Future research directions in this field may also include the development
497 of advanced imaging techniques to detect NF- κ B activation in specific glial cell populations in vivo,
498 allowing for real-time monitoring of glial responses in glaucoma patients. Additionally, the use of
499 single-cell transcriptomics and proteomics could provide valuable insights into the heterogeneity of
500 glial cell responses and help to identify specific cellular subpopulations that may be particularly
501 important in mediating neuroprotection or neurodegeneration.

502 In summary, the role of NF- κ B signaling in retinal glial cells represents a complex and

503 multifaceted aspect of glaucoma pathophysiology. By unraveling the intricacies of this signaling
504 pathway and its effects on glial cell function, researchers and clinicians may be able to develop more
505 effective and targeted treatments for glaucoma, ultimately improving the outcomes of patients affected
506 by this sight-threatening disease.

507

508 **Author Contributions**

509 FV and MDVN drafting the manuscript prepared the figure and performed bibliographic search. IF, FD, FDG and
510 DVer performed bibliographic search and formatting; DC, AA, EA reviewed the manuscript critically given crucial
511 intellectual content. DVec and FZ conceived the project and reviewed the final version of the manuscript. All authors
512 read and approved the final manuscript.

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520 **Search Methodology**

521 We performed an extensive MEDLINE research on PubMed focusing our attention on high research paper and review
522 manuscript using the below keywords: “glaucoma”, “NF-κB”, “NF-κB and glaucoma pathophysiology”, “Retina
523 glial cells”, “NF-κB and retina glial cells”. We mainly selected publications from the past 10 years, but also include
524 highly referenced and highly regarded publications outside of this window.

525

526 **Conflict of Interest**

527 The authors declare no conflict of interest

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530 **References and Notes**

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