

Pathogenetic mechanisms and treatment options for ophthalmic pterygium: Trends and perspectives (Review)

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Abstract. Pterygium is a fibrovascular lesion of the ocular surface that can display an aggressive clinical behavior and, occasionally, threaten vision. Although there is no consensus on its pathogenesis, recent evidence suggests that it is a proliferative, rather than degenerative condition strongly correlated with exposure to ultraviolet radiation (UVR) of solar light. Molecular genetic alterations reported in association with pterygium include loss of heterozygosity (LOH), point mutations of proto-oncogenes, such as K-ras and alterations in the expression of tumor suppressor genes, such as p53 or p63. Other findings in pterygium include the frequent detection of HPV DNA, ocular surface changes such as the overexpression of various proteins, including defensins and phospholipases D, as well as the up-regulation of growth factors, such as bFGF or VEGF. Although pterygium management has traditionally involved surgery, often enhanced by the use of antimetabolites, recent advancements in the understanding of molecular and biochemical events underlying pterygium pathogenesis may enable the use of less invasive treatment methods.

Contents

1. Introduction
2. Epidemiology, clinical and histochemical features
3. Pathogenetic theories
4. Treatment options
5. Conclusions and future trends

1. Introduction

Despite the fact that its original description dates back to the time of Hippocrates, the ophthalmic pterygium still manages

to keep the secrets of its pathogenesis and, more importantly, may still occasionally threaten vision (1). Apparently, the condition has not changed since the ancient times and can be described as a fibrovascular membrane advancing on the corneal surface, often triangular in shape (with the apex towards the corneal centre), sometimes bilateral and usually originating from the nasal bulbar conjunctiva, but also occasionally from the temporal conjunctiva (Fig. 1) (1,2). Numerous theories have so far been proposed to describe its pathogenesis, although none has gained unanimous acceptance (2,3). Pterygium was originally considered an 'ophthalmoheliosis', i.e., a condition closely related to the exposure of the eyes to increased amounts of sunshine which could, through poorly understood mechanisms, result in fibrovascular growth on the ocular surface (2,4). However, during the last decades, several potential mechanisms for pterygium development have been examined, mostly by employing molecular genetic analysis techniques of tissue samples (5-10). Consequently, insights into the pathogenesis of pterygium were gained through the improved understanding of molecular genetic pathways (5-10). The advent of commercially available pharmaceutical agents which could, acting on a molecular level, affect important aspects of pterygium clinical behaviour, such as the angiogenesis or cellular proliferation, opens a promising new therapeutic avenue (11,12). Therefore, future treatment options may be less invasive and more focused on molecular pathogenetic pathways.

2. Epidemiology, clinical and histochemical features

The condition is common in the so-called 'pterygium zone' which is defined by a geographical latitude of 40° north and south of the equator (4). In countries within this area prevalence of up to 22% in the general population has been reported (2). In countries outside this area (in either northern or southern latitudes) reported prevalence rates usually do not exceed 2% of the general population and the lesion affects mostly patients with an increased exposure to solar light, such as people working outdoors (13,14). A slightly higher incidence in males is reported which may be attributed to different life-styles between genders (with males spending more time out-doors) in many countries (13,14).

Apart from the triangular shape (often described as having a head, body and tail), pterygium is characterized by its location along the inter-palpebral fissure (along the horizontal

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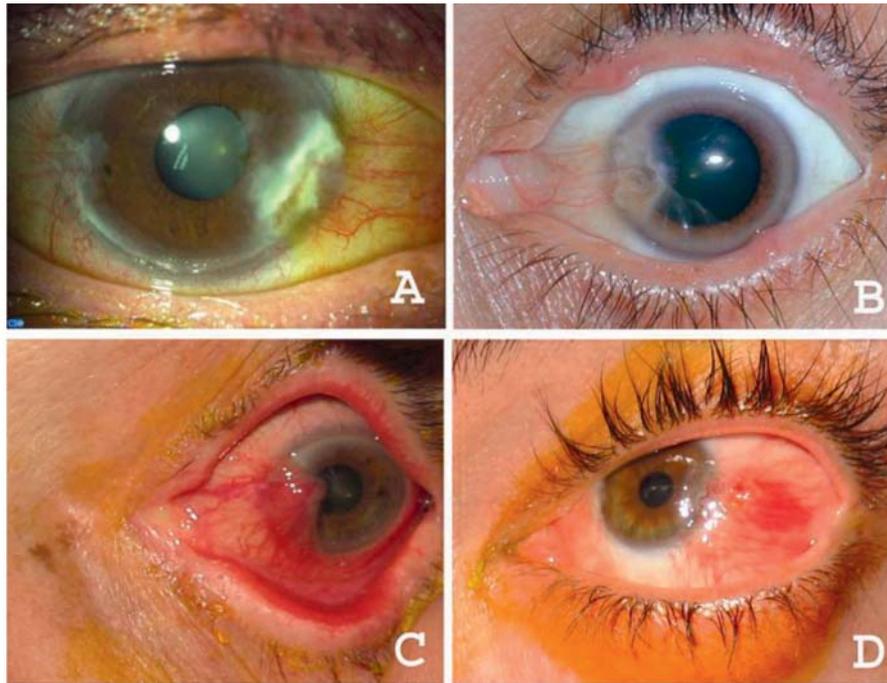


Figure 1. Various forms of pterygium, including simultaneous nasal and temporal pterygium (A), quiescent nasal pterygium (B), inflammatory nasal pterygium (C) and inflammatory temporal pterygium (D).

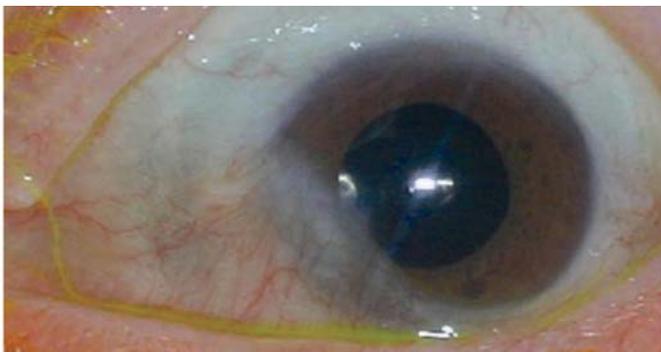


Figure 2. Pseudopterygium developing at the inferior-nasal quadrant of the bulbar conjunctiva following a local acid burn.

axis of cornea, usually nasally but occasionally also temporally or both nasally and temporally, Fig. 1), which was considered an important indication of the role of environmental factors, such as solar light, in its development. In fact, the development of pterygium-like lesions in axes other than the horizontal (pseudo-ptyerygium) is secondary to traumatic, inflammatory or vascular conditions and, therefore, not true pterygium (Fig. 2). Symptoms associated with pterygium development include chronic ocular surface inflammation, tearing, astigmatism and blurred vision due to optical axis involvement on the corneal surface (6). The natural history of the condition is variable, sometimes characterized by prolonged static periods, often associated with corneal linear iron deposition (Stocker's line) along pterygium head, originating from tear film lactoferrin (Fig. 3B) (15). However, progressive growth characterizes the majority of pterygia, especially those affecting younger individuals, often necessitating surgical removal (16,17). Pterygium surgery can result in further clinical

manifestations, including symblepharon formation (often affecting ocular motility), corneal clouding and corneal or scleral dellen (Fig. 3) (17,18). Importantly, postoperative recurrence (Fig. 4) is common (described in up to 50% of cases, depending on the technique employed) and sometimes results in even more aggressive clinical behaviour (17,18).

Histologically, pterygium is characterized by elastotic degeneration of conjunctival substantia propria, with associated eosinophilic or basophilic deposits (19). Degenerated type I and type IV collagen fibers as well as immature elastin fibers (elastin dysplasia) or degenerated elastin fibers (elastin dystrophy) have also been described (19,20). Epithelial changes are variable and include hyperkeratosis, parakeratosis or akantosis (21). Fibroblasts lying at the anterior corneal stroma (underneath Bowman's membrane) may become activated by ultraviolet radiation and may cause Bowman's membrane rupture resulting in the firm adherence of pterygium on underlying corneal stroma (22). On the contrary, the presence of Tenon's capsule prevents the adherence of the episcleral portion of pterygium on the underlying sclera (21,22). From a histochemical standpoint, abnormal forms of mucopolysaccharides, often with increased amounts of neutral sugars and sialic acid, are reported in pterygia, together with higher levels of specific amino acids, such as praline, compared with normal conjunctiva (23-25).

3. Pathogenetic theories

Early pathogenetic concepts. The association between pterygium development and specific lifestyles, such as outdoors working or exposure to sunshine or dust led to the idea that chronic ocular surface irritation by such environmental factors might be the cause of the condition (2,4). It was also proposed that pterygium arises from other sunshine-related conditions,

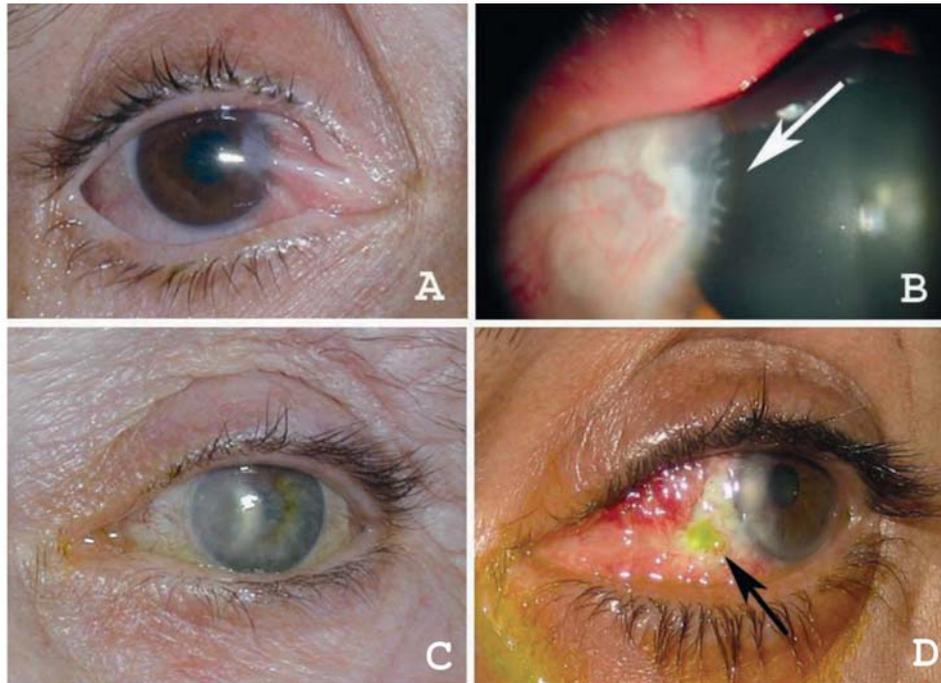


Figure 3. Aspects of pterygium clinical behaviour, including the postoperative symblepharon formation (A), Stocker's line along pterygium head (B), postoperative corneal opacity at the site of pterygium bed (C) and scleral dellen with impending perforation following pterygium removal with adjuvant MMC application (D).

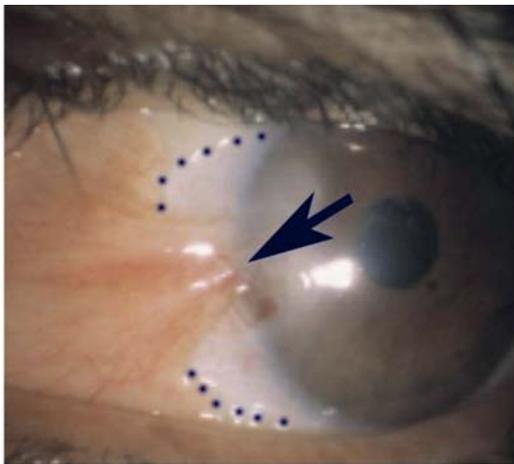


Figure 4. Pterygium postoperative recurrence as a fibrovascular band (arrow) crossing the epithelialized area of bare sclera created at the initial pterygium removal (blue dots) and advancing on the corneal surface.

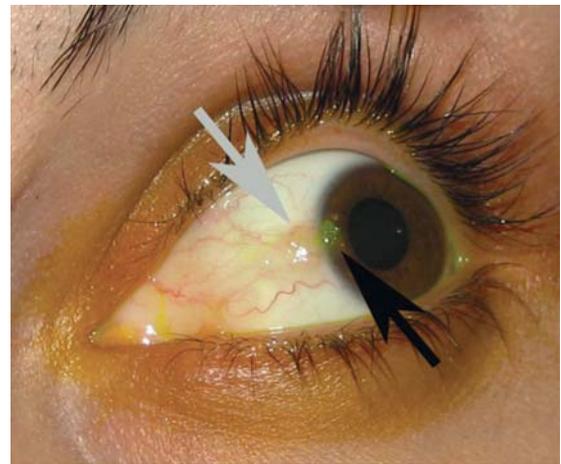


Figure 5. Pingueculum (grey arrow) is a lesion of the sclerocorneal limbus that differs from pterygium in that it does not grow on the corneal surface. However, it may become inflamed or be associated with local corneal dellen (black arrow) and, according to one theory, may be a precursor to pterygium development.

such as the pingueculum (Fig. 5), a hyalinized nodule appearing at the sclerocorneal limbus (26). Pingueculum has no growth potential *per se* but may become inflamed and can evolve into a true pterygium (26). The frequently observed conjunctival vascular congestion in the area of pterygium body has led to the suggestion that medial rectus activity, which underlies the area of pterygium growth nasally, can cause blood flow disturbance potentially associated with pterygium growth (2,4). Early reports have even involved alterations in lacrimal composition or flow (pooling of tears) or the unspecified local effects of lactic acid secreted by periocular sweat glands (27-29).

Concepts of inheritance. Early reports mentioned the possibility of an autosomal dominant pattern of inheritance, based on

cases with familial occurrence of pterygium (2). However, it could not be verified whether the pterygium itself is inherited, as an independent trait, or if the affected individuals shared an increased susceptibility to the oculodermal effects of solar light (6,7,30). The detection of potential tumor suppressor gene involvement in pterygium has raised the possibility of a 'two hit' mechanism (Knudson's theory) in its pathogenesis (7,30). According to this hypothesis, the 'first hit' in the process of a tumor suppressor gene deactivation may be inherited whereas the 'second hit' may be inflicted by environmental factors, such as a viral infection or UVR (30). However, the fact that no specific genetic locus has so far been associated

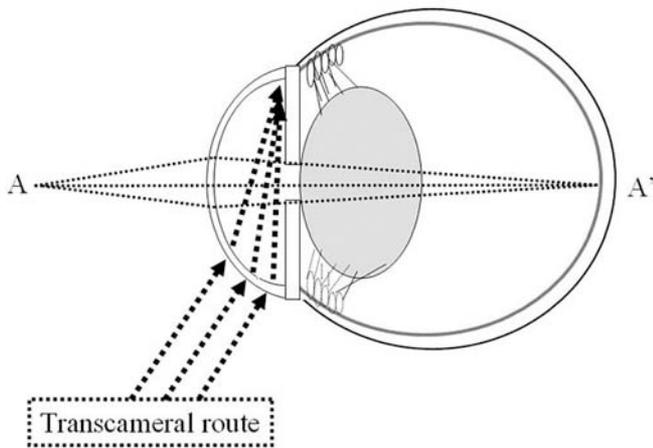


Figure 6. Transcameral light focusing on the nasal limbus may expose limbal basal stem cells to increased amounts of UVR and be associated with molecular genetic alterations to these cells, eventually leading to pterygium formation.

with pterygium development implies that further research in the area of genetic susceptibility for pterygium is required before safer conclusions on this issue are reached.

The role of UVR. The strong epidemiological correlation between pterygium development and exposure to sunshine has led to the assumption that some parts of the solar radiation may have a direct pathogenetic role (2,4). Peri-orbital architecture, including the presence of the superior orbital rim, the presence of eyebrows and the nasal prominence imply that the eyes are relatively protected from direct light exposure from above (2,4). On the other hand, the eyes are relatively unprotected from light scattered from below or laterally (albedo) (4). Early reports raised the possibility that solar light exposure acted in combination with exposure to dust or sand, thus leading to chronic ocular surface inflammation (2). However, the detection of high pterygium prevalence in sailors or fishermen, who lived in environments devoid of dust but instead were exposed to increased amounts of scattered light from reflective surfaces such as the sea surface, pointed towards an albedo involvement in pterygium formation (31). In fact, it was suggested that scattered light might follow alternative (transcameral) optical paths when entering the eye, thus hitting limbal stem cells from their inner surface (Fig. 6) (4). Some studies have even suggested that the nasal location of pterygium is an indication of the pathogenetic role of solar light, since light may be reflected onto the nasal sclerocorneal limbus from the lateral nasal wall whereas the nasal bulbar conjunctiva is more exposed to sunshine since brow hair is shorter nasally than temporally (Fig. 7) (4). The effects of UVR-A and UVR-B (290-400 nm) are considered particularly detrimental (32). At a molecular level, UVR is associated with the creation of active free radicals which attack and deactivate various macromolecules (33). Free radicals may in turn be deactivated by tear film proteins, including lactoferrin (34). The presence of Stocker's line along the head of pterygia lacking growth potential, as previously mentioned, may represent abnormal local iron metabolism along the advancing head of pterygium. Iron

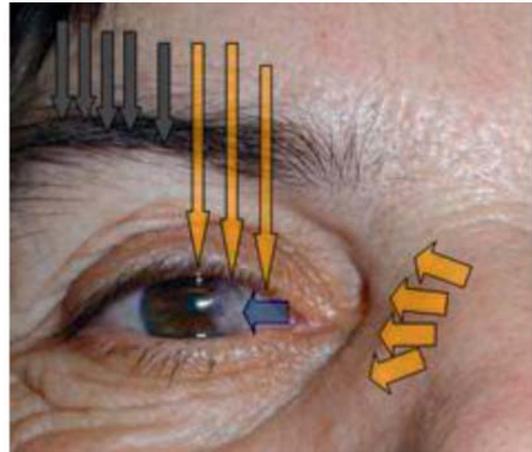


Figure 7. Light reflected from the lateral aspect of the nose on the corneal limbus as well as incident light falling on nasal limbus due to the short length of brow hair nasally (yellow arrows) may contribute to the formation of pterygium nasally (blue arrow). On the other hand, the temporal limbus is relatively protected from incident light due to the longer length of brow hair temporally (grey arrows).

itself may be associated with increased free radical formation through biochemical (Fenton) reactions (35).

Ocular surface changes and pterygium. Previous studies have reported an association between pterygium and dry eye changes, such as reduced tear-film break-up time (BUT) (28,36,37). Such findings indicated that pterygium may be a manifestation of a generalized ocular surface dysfunction, including a chronic inflammatory reaction (38). In accordance with these reports, an up-regulation of the expression of human α -defensins and S100 A8 and A9 in tear fluids of pterygium patients provides additional indications for a connection between pterygium and chronic ocular surface inflammation (38). Another study has reported the up-regulation of phospholipase D (PLD) types 2, 3 and 4 in pterygium, compared with normal conjunctiva (39). PLD are involved in various processes, including inflammation, cell differentiation, apoptosis and wound healing and the detection of PLD alterations in pterygium supports their potential role in pterygium pathogenesis (39).

Oxidative stress. Increased UVR-associated oxidative stress has been reported in pterygium, compared with normal conjunctiva, leading to the induction of proteins, such as survivin (40). The latter has been correlated with DNA oxidation and down-regulation of p53 (40). Interestingly, it has been suggested that the presence of iron deposits along the corneal head of pterygium may indicate oxidative stress affecting local epithelial cells and resulting in disturbed iron homeostasis (35). The topical use of antioxidants as an adjuvant treatment option for pterygium would thus seem logical and should be investigated in future research projects.

Molecular genetic alterations. Previous studies have detected both loss of heterozygosity (LOH) and microsatellite instability (MI) in pterygium (5,6). However, the association of LOH with pterygium was found much stronger, compared with the

respective association with MI implying that the involvement of the latter in pterygium etiopathogenesis is less important (6). On the contrary, the close association of LOH with pterygium points towards an involvement of tumor suppressor genes in its development (6,30). Indeed, *p53* levels have been reported increased in pterygia by several previous studies although other studies have not confirmed such findings (3,41,42). The up-regulation of *p53* expression in pterygia may merely reflect increased exposure to UVR, since the wild-type of *p53* is known to increase in normal tissues in response to DNA damaging agents, such as UVR (3,43). The generally employed methodology to measure *p53* tissue levels (antibodies against *p53* protein) cannot distinguish between wild-type and mutant forms (3). Wild-type *p53* is critical in causing cell-cycle arrest in rapidly proliferating cells (3). However, its action depends on various factors, such as the levels of acetylation or phosphorylation (44,45). Interestingly, the levels of *p53* expression in pterygia have been found to differ between epithelial layers being higher in basal cells, compared to more superficial layers (3,46,47). This finding could reflect increased exposure to UVR according to the proposed theory of transcameral exposure of limbal basal (stem) cells to solar light. Recurrent pterygia also display reduced amounts or total lack of *p53* expression which may be an indication that the epithelial cells escape the *p53*-dependent cell cycle checkpoint (3). Alterations in proteins with functional and structural homology to *p53*, such as *p63* and *p73*, have also been detected in pterygium (48,49). These proteins are involved in cellular differentiation and alterations in their genes have been linked to various systemic and ocular abnormalities (3,48,49). Interestingly, *p63* expression in pterygium resembles the patterns seen in normal limbus (expressed mainly at the basal epithelial layers) and differs from the pattern seen in normal cornea (almost total absence of *p63* expression), a finding supporting the possibility of limbal origin of pterygium (50).

Apart from TSGs, proto-oncogene (POG) sequences may also be affected by UVR (51,52). Mutations in genes of the *ras* family POGs, including Harvey-Ras (*H-ras*), Kirsten-Ras (*K-ras*) and Neuroblastoma-ras (*N-ras*), have been detected in cultured cells of mouse skin tumors following exposure to near-UVR (51) and are also frequently detected in human neoplastic conditions of sun-exposed body areas, such as cutaneous melanomas (52). Such mutations may convert these genes into active oncogenes (53). Restriction fragment length polymorphism (RFLP) and sequencing analysis of primary and recurrent pterygia has revealed *K-ras* mutations at codon 12 in 10% of pterygia (a transition from GGT-glycine to GTT-valine, in all cases) and an absence of such mutation in respective phenotypically normal conjunctival specimens, as well as peripheral blood samples (8). A lack of mutations at codons 12 and 13 for *H-ras* and *N-ras* was also reported (8). Importantly, mutations at codon 12 of *K-ras* was correlated with post-operative recurrence and young age, implying that such alterations have a direct link with the clinical profile of pterygium (8). The involvement of TSG and POG in pterygium implies that pterygium shares some similarities with neoplastic conditions (8,30). Many neoplasias and immortalized cell lines also display up-regulation of telomerase, a ribonucleoprotein participating in cell division by blocking telomere

shortening (3,54,55). Telomerase is not expressed in most adult human tissues, including normal conjunctiva (3,54,55). Telomerase is however expressed in pterygium, a finding attributed to neoplastic features of the lesion or, alternatively, to induction by UVR (3,54,55).

Another interesting aspect of the molecular genetic profile of pterygium is the altered apoptotic potential of its epithelial cells (3). Previous studies have found that UVR induces apoptosis to normal corneal epithelium (56,57). However, epithelium in pterygium is possibly resistant to UVR-mediated apoptosis (3). It is likely that this finding is due to a disturbed balance between pro-apoptotic proteins (such as *bax*) and anti-apoptotic proteins (such as *bcl-2*) in the epithelium of pterygium, in favour of the latter (58). Again, differences in the apoptotic status between different epithelial cell layers have been detected with apoptosis remaining active at the basal epithelium of pterygium but not at more superficial epithelial layers (3,59).

The role of genetically altered limbal stem cells. In their 1994 study, Kwok and Coroneo proposed that the initial biologic event in pterygium pathogenesis may be a genetic alteration of limbal stem cells, due to chronic UVR exposure (60). They postulated that a breakdown of the corneoscleral limbal barrier results in subsequent conjunctivalization of the cornea resulting in pterygium (60). They also suggested that the triangular (wing-like) shape of pterygium may in fact be explained by the limbal origin theory since a disrupted balance between the populations of epithelial cells in cornea and conjunctiva could, according to data from animal models, result in advancement of conjunctival epithelium on the corneal surface (60). Further studies in this area, employing immunohistochemistry, found that, in contrast to normal conjunctival, limbal, and corneal cells which immunostain primarily for matrix metalloproteinase-I (MMP-I), limbal basal epithelial cells (pterygium cells) immunostain for multiple types of MMPs (MMP-1, MMP-2, MMP-3, MMP-9, membrane type 1-MMP, and membrane type 2-MMP) (3). According to these results, the altered MMP expression of limbal basal epithelial cells (pterygium cells) enables them to invade and dissolve Bowman's layer leading to firm adhesion of the lesion on the corneal surface (3,61).

Viral involvement. Although early reports mentioned the possibility of an infectious origin of pterygium, evidence on this issue was not conclusive (4). However, the advent of PCR as a research tool enabled the detailed investigation for viral DNA in samples of both pterygia and normal conjunctiva (7). The presence of viruses known to cause oculodermal infections, such as herpes simplex virus (HSV), and human papilloma virus (HPV), was examined (7,62,63). Results from several studies point towards the involvement of HPV in pterygium although large regional and racial differences have been reported (62,63).

Growth factors and cytokines. UVR-mediated genetic trauma may affect the expression of various cytokines, growth factors and growth factor receptors (3). The presence or altered expression of such factors in pterygium has been evaluated by previous studies using various methods including immuno-

histochemistry, or ELISA (9). Many such factors also participate in normal corneal healing and their altered expression in pterygium may indicate a response to ocular surface damage inflicted by the lesion (3). UVR-inducible cytokines include the interleukin-1 (IL-1) system, acting in concert with tumor necrosis factor α (TNF α) it leads corneal keratocytes to adopt a repair phenotype (64,65), the IL-6 (it promotes epithelial cell migration through the induction of integrin receptors) (66,67) and IL-8 (displays mitogenic and angiogenic activity) (68,69). Growth factors involved in pterygium according to previous reports include the epidermal growth factor (EGF) and heparin-binding EGF (HB-EGF), vascular endothelial growth factor (VEGF), basic fibroblast growth factor (bFGF), platelet-derived growth factor (PDGF), transforming growth factor- β (TGF- β) and insulin-like growth factor binding proteins (IGF-BP) (3). VEGF is involved in angiogenesis and is produced by corneal fibroblasts in response to inflammation or other noxious stimuli, including UVR (70). VEGF has been detected in increased amounts in pterygium epithelium, compared with normal conjunctiva by studies employing immunohistochemistry (71). These results were confirmed by RT-PCR assays also revealing a correlation between VEGF expression and post-operative recurrence (own unpublished data). bFGF is involved in the corneal healing process as well as neovascularization and is also overexpressed in cultured fibroblasts from pterygia (3,9,72). Again, RT-PCR assays showed increased bFGF expression in pterygium, compared with normal conjunctiva, a finding possibly reflecting UVR-mediated induction (own unpublished data).

4. Treatment options

Surgical excision techniques. Early pterygium removal included simple excision (detachment) of pterygium head from the anterior corneal surface (16,73). However, the realization of the potential for recurrence, often more aggressive clinically than the original lesion, soon lead to modifications of simple excision (16,74). The 'bare sclera' technique involved the removal of the lesion and peri-limbal conjunctiva with suturing of the remaining conjunctival rim onto the bare scleral surface at a variable distance from sclerocorneal limbus (Fig. 8A) (75). The rationale for the 'bare sclera' technique was that the area left uncovered would be epithelialized from epithelial cells from the anchored conjunctival rim which could then act as a barrier against pterygium re-growth from pathological tissue remnants inevitably left *in situ* (16,75). However, up to 50% of recurrence rate following a successful 'bare sclera' pterygium removal have been described (16).

The understanding of the important role of normal conjunctiva in blocking pterygium re-growth on the corneal surface was the basis for further modifications of the initial technique which in fact departed from the 'bare sclera' idea (16,74). These included conjunctival flap transposition, i.e., the formation of a mobile conjunctival flap, either superiorly or inferiorly to the pterygium bed, which could then be transposed onto the area previously occupied by pterygium, thus providing the normal tissue blockage against pterygium recurrence from more peripheral sites (Fig. 8B and C) (76,77). Instead of the conjunctival flap, a free conjunctival

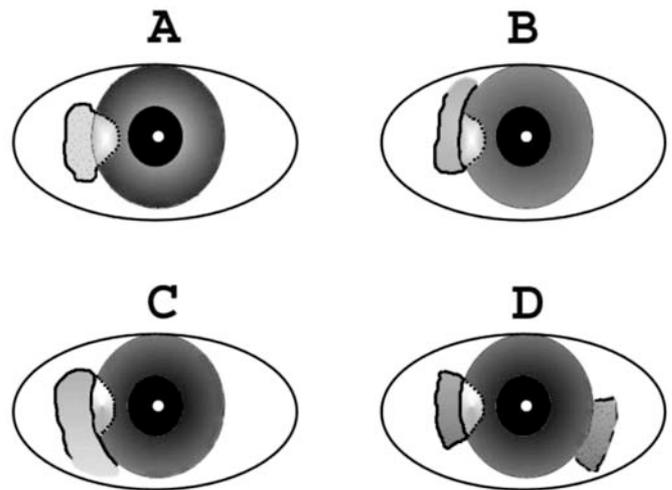


Figure 8. Techniques of pterygium excision include bare sclera removal (A), the creation of advancement conjunctival flaps from superior (B) or inferior (C), bulbar conjunctiva to cover the pterygium bed as well as the creation of a free conjunctival graft, often from the inferior-temporal bulbar conjunctiva, which also contains normal limbal stem cells (D).

graft may be even more advantageous towards this goal because it not only covers the pterygium bed with normal tissue but also can potentially provide a population of normal stem cells positioned onto the sclerocorneal limbal area originally associated with pterygium (Fig. 8D) (78,79). In cases with very large conjunctival defects created following pterygium excision, or if a proper autologous conjunctival graft cannot be harvested, an alternative technique is the use of preserved amniotic membrane, which is readily commercially available and provides an excellent substrate for epithelial re-growth (80). The use of more sophisticated conjunctival surgery is associated with considerably lower recurrence rates, often less than 10% (16,78). Other modalities for pterygium removal, such as ablation with excimer laser, have not gained widespread acceptance, although excimer laser may have a role in smoothing corneal surface irregularities caused by pterygium (81,82). However, lamellar keratoplasty may also be required, especially in cases of recurrent pterygia with firm adhesion to the corneal stroma (83).

The role of antimetabolites. The fear of an aggressive recurrence always follows pterygium excision, no matter how elegantly primary surgery has been performed. This is why surgery is often combined with adjuvant therapies such as antimetabolites *e.g.* mitomycin-C (MMC) (16,78). The latter is an antibiotic, a product of *streptomyces caesiopsis*, capable of alkylating DNA double helix and blocking both transcription and translation (84,85). Thus MMC is also a potent antimetabolite used in suppressing tumor cells (84,85). In the case of pterygium surgery MMC is used in concentrations of 0.2-0.4 mg/ml applied episclerally for various intervals (usually not more than 2 min) (84). The recurrence rate with the adjunctive intraoperative use of MMC is reported to be <10% (16,78,86). Apart from the intraoperative use, MMC has also been used as postoperative eye drops (16,87). Other agents include the alkylating agent Thiotepea (N,N,N'-triethylenethiophosphoramidate), used in concentrations of

1:2000, either postoperatively or as eye drops (16) and 5-fluorouracil (5-FU), a pyrimidine analogue used either intraoperatively or as postoperative subconjunctival injections (16,88).

Although very effective in reducing recurrence rates, anti-metabolite use is, nevertheless, associated with serious and potentially sight-threatening complications, such as delayed healing or even scleral melt (Fig. 2), sometimes threatening vision or requiring further surgery for their management (16). The thin or necrotic sclera resulting from antimetabolites may also be treated with the use of hyperbaric oxygen which induces hyperoxia, angiogenesis and episcleral fibroblast proliferation (89).

β-irradiation. Historically, irradiation has been one of the first attempts of modern surgery to suppress the potential for recurrence in pterygium management (90,91). β-irradiation, delivered through strontium-/yttrium-90 sources, effectively reduces cellular populations responsible for pterygium recurrence (92). However, its use is not innocuous and may be associated with serious complications, including vision-threatening endophthalmitis (16).

Growth factor inhibitors. Anti-VEGF monoclonal antibodies have become widely available in ophthalmic practice mainly because of their success in suppressing various forms of intraocular neovascular growth, such as exudative age-related macular degeneration and sub-retinal neovascular membranes, proliferative diabetic retinopathy, or neovascular glaucoma (93). Such factors include Pegaptanib (Macugen, Pfizer), an oligonucleotide aptamer that binds exclusively to the 165 amino acid isoform of VEGF, and recombinant monoclonal antibody Bevacizumab (Avastin, Genentech) as well as its fragment Ranibizumab (Lucentis, Genentech), both directed against VEGF (93). Previous studies have already evaluated the potential use of Bevacizumab in pterygium management (11,12). However, Bevacizumab is also associated with potential serious side effects, including significant cardiovascular toxicity (93). The study of VEGF expression in individual lesions may therefore allow for selective Bevacizumab or other anti-VEGF administration, potentially reducing the risk of recurrence or aggressive clinical behaviour without taking unnecessary systemic risks.

5. Conclusions and future trends

Ophthalmic pterygium continues to puzzle doctors, affecting large numbers of patients in many parts of the world. Nevertheless, research with modern molecular techniques has led to more accurate and detailed information on its origin and growth mechanisms. It is likely that UVR induces ocular surface changes especially affecting stem cells at the corneo-scleral limbus. Such changes may be potentially enhanced by the action of other environmental factors, such as HPV infection or oxidative stress. At a molecular level, findings include alterations in the expression of tumor suppressor genes, such as p53, the induction of MMPs as well as alterations in the expression of various growth factors, including VEGFA and bFGF. At a histological level, Bowman's membrane

damage, fibrovascular growth as well as elastotic degeneration of various connective tissue elements predominate. Until recently, surgery was the only feasible option of pterygium treatment, in the form of various conjunctival plastic procedures usually enhanced with adjunctive modalities, such as antimetabolites. However, apart from being invasive and associated with a variety of potentially serious complications, surgery often results in a more aggressive clinical behaviour of pterygium. The advent of commercially available anti-VEGF has already been explored by preliminary studies. Apparently however, pterygia differ significantly in the profile of cytokine/growth factor alterations they display. The individualized examination of each lesion may enable a customized treatment strategy, according to its molecular profile, thus reducing risks from invasive treatment methods and at the same time increasing success rates.

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