Elevated aqueous humour tissue inhibitor of matrix metalloproteinase-1 and connective tissue growth factor in pseudoexfoliation syndrome

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Background/aims: Pseudoexfoliation syndrome (PXF) was recently found to be associated with increased expression of transforming growth factor β_1 (TGF β_1) in the aqueous humour. As concern has been raised regarding anti-TGF β therapy, which can potentially disrupt the maintenance of anterior chamber associated immune deviation, the authors explored the levels of tissue inhibitor of matrix metalloproteinase-1 (TIMP-1), matrix metalloproteinase-9 (MMP-9), and connective tissue growth factor (CTGF) in aqueous humour to determine if these may represent alternative therapeutic targets.

Methods: Aqueous humour samples were collected from patients who underwent routine cataract surgery. All patients were categorised into three main groups—PXF, uveitis, and control. The PXF group was further subcategorised into three grades based on the density of the exfoliative material observed on biomicroscopy, as well as the presence or absence of glaucoma. TIMP-1, MMP-9, and CTGF levels were measured using specific enzyme immunoassays (ELISA).

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Accepted for publication 8 July 2004 **Results:** Eyes with PXF had significantly higher aqueous humour TIMP-1 concentration (n = 56, mean (SE), 9.76 (1.10) ng/ml) compared with controls (n = 112, 5.73 (0.43) ng/ml, p<0.01). Similarly, the CTGF level in PXF eyes (n = 36, 4.38 (0.65) ng/ml) was higher than controls (n = 29, 2.35 (0.46) ng/ml, p<0.05). Further, the CTGF concentration in the PXF glaucoma group is significantly higher compared with PXF eyes without glaucoma (6.03 (1.09) ng/ml v 2.73 (0.45) ng/ml, p<0.01). The MMP-9 levels were low and below detection limit in all PXF and control samples with no statistical difference between groups.

Conclusion: A raised TIMP-1 level and a low MMP-9 level in aqueous humour of PXF eyes may imply a downregulation in proteolytic activity. The increased CTGF concentration supports the proposed fibrotic pathology of PXF. Regulation of MMP/TIMP expression and anti-CTGF therapy may offer potential therapeutic avenues for controlling PXF associated ocular morbidity.

There has been an increasing awareness of the role of growth factors in aqueous humour homeostasis leading to the development of raised intraocular pressure and glaucoma.^{1 2} As the aqueous humour bathes the anterior segment, its compounds affect the metabolic functioning of the cells lining the conventional outflow routes—that is, the trabecular meshwork (TM). In humans, the TM is made up of collagen beams covered by endothelial-like cells. The spaces between the beams are filled with extracellular matrix (ECM). The ECM in the subendothelial region of Schlemm's canal plays an important part in the pathogenesis of glaucoma.³ Elevated intraocular pressure may be attributed to abnormal deposition of ECM resulting in a reduced filtration capacity.^{3 4}

Characterised by progressive accumulation of a fibrillar extracellular material most notable on the pupillary border and anterior lens capsule, pseudoexfoliation syndrome (PXF) encompasses more than half of the cases of open angle glaucoma in Norway, Ireland, Greece, and Saudi Arabia.⁵ ⁶ The exact biochemical nature of the exfoliative material, the mechanism of its production, and pathophysiology leading to elevated intraocular pressure in PXF remain to be unravelled.

Recent studies have found PXF to be associated with increased levels of transforming growth factor β_1 (TGF β_1),^{7 8} basic fibroblast growth factor (bFGF),⁹ and hepatocyte growth factor (HGF),¹⁰ and 8-isoprostaglandin F2a (8-IPGF)¹¹ in the aqueous humour. It has been speculated that these findings are indicative of oxidative stress as well as an

abnormality in fibrogenesis, matrix degradation, and cellular response to injury in the anterior segment of PXF eyes.

TGF β_1 , a fibrogenic cytokine, has a pivotal role in the ocular wound healing processes. It is also essential in the maintenance of the anterior chamber associated immune derivation (ACAID).² It promotes deposition of ECM by downregulating the production of proteases and stimulating the synthesis of protease inhibitors.¹²⁻¹⁴ Huang et al demonstrated the presence of matrix metalloproteinases (MMP) and tissue inhibitors of matrix metalloproteinases (TIMP) in normal aqueous humour. They suggested that an imbalance in the MMP/TIMP ratio may contribute to anterior eye segment ECM alteration associated with ocular neovascularisation and glaucoma.15 Connective tissue growth factor (CTGF), also plays a part in the ECM deposition remodelling and wound healing. Studies have shown that CTGF is an essential downstream mediator of many of the effects of TGF β_1 , particularly those associated with fibrosis,^{16 17} and CTGF has been detected in human aqueous humour.18

As concern has been raised regarding anti-TGF β_1 therapy which can potentially disrupt the maintenance of ACAID,¹ we

Abbreviations: ACAID, anterior chamber associated immune derivation; bFGF, basic fibroblast growth factor; CTGF, connective tissue growth factor; ECM, extracellular matrix; HGF, hepatocyte growth factor; 8-IPGF, 8-isoprostaglandin F2a; PXF, pseudoexfoliation syndrome; MMP-9, matrix metalloproteinase-9; TGFβ, transforming growth factor β; TIMP-1, tissue inhibitor of matrix metalloproteinase-1; TM, trabecular meshwork

MATERIALS AND METHODS

Aqueous humour was obtained from patients who underwent routine cataract surgery. All patients had slit lamp examination on the day before or on the morning before surgery by the same investigator (SLH) and they were categorised into three main groups:

- Pseudoexfoliation syndrome (PXF)
- Past history of intraocular surgery or ocular inflammatory diseases
- Control (no signs of intraocular pathology apart from cataract).

The pseudoexfoliation group was further subcategorised into three grades based on the biomicroscopy findings.

- PXF grade I (A1): PXF material only visible after pupillary dilatation
- PXF grade II (A2): PXF material visible scattered discontinuously along pupillary margin and anterior lens capsule before pupillary dilatation without iridodonesis or phacodonesis
- PXF grade III (A3): Dense PXF material visible continuously along the circumference of pupillary margin before dilatation with or without iridodonesis or phacodonesis.

In addition, the PXF group was divided into two subcategories—that is, PXF without glaucoma and PXF with glaucoma based on case notes history.

Patients with glaucoma were maintained on their preoperative antiglaucoma medication (ranging from timolol (MSD), betaxolol (Alcon), levobunolol (Allergen), dorzolamide/timolol(MSD), or latanoprost (Pfizer) monotherapy in combinations of the above with brimonidine (Allergen), brinzolamide (Alcon), dorzolamide (MSD), or apraclonidine (Alcon)). All patients who underwent cataract surgery were given topical ocular premedication of 2.5% phenylephrine hydrochloride, 1% cyclopentolate hydrochloride, and 0.5% chloramphenicol preservative free drops before surgery. Data on medical history, medications, and demographic information were obtained from the case notes.

Aqueous humour sampling

Samples of aqueous humour were obtained in the first intraocular manoeuvre (paracentesis site for phacoemulsification cases or first full thickness entry wound from a stepped corneolimbal incision for extracapsular extraction). Aqueous humour was aspirated into a 1 ml insulin syringe with a Rycroft cannula (Maersk Medical Ltd, Redditch, UK). Meticulous care was taken to avoid touching the iris, lens, and corneal endothelium and to prevent contamination of the aqueous sample from irrigation fluid and blood from the conjunctiva. A 100–150 µl sample of aqueous humour was collected from each patient. The aqueous humour sample was immediately stored at 4°C in the eye theatre fridge and was transferred to the surgical laboratory within the next 2 hours to be stored at -80°C until used.

Informed consent to the aqueous humour contribution was obtained from patients, and the research protocol was approved by the clinical research ethics committee of the Cork Teaching Hospitals and Mater Misericordiae Hospital.

A total of 202 aqueous humour samples were collected; 121 samples were from female subjects and 81 were from male. Out of these, we had 69 samples which came from patients with PXF (mean (SE) age, 79.9 (0.65) years; range,

69–91 years), 17 from patient with recent ocular surgery or uveitis (65.8 (3.85) years; range 27–86 years), and 116 samples from control subjects (75.5 (0.78) years; range 48–92 years).

Enzyme immunoassays

TIMP-1 and MMP-9

The concentration of the aqueous humour TIMP-1 and total MMP-9 (active and pro-MMP-9) were quantified using commercially available sandwich enzyme immunoassay kits (Quantikine, R&D systems, Abingdon, UK) with a minimum detectable concentration of 0.08 ng/ml and 0.156 ng/ml respectively. A 25-fold dilution was applied for the sample preparation during TIMP-1 assay and a 10-fold dilution for MMP-9 assay. Each ELISA kit has a specificity of detecting native human as well as recombinant TIMP-1 or MMP-9. A standard calibration curve was generated using the standard TIMP-1 or MMP-9 provided by the manufacturer: with seven standard solution within a range of 0.156-10 ng/ml for TIMP-1 and with six standard solution ranging from 0.312 to 10 ng/ml for MMP-9. Each TIMP-1 and MMP-9 assay was performed in duplicate according to the manufacturer's advice.

CTGF

CTGF sandwich ELISAs were performed by FibroGen Inc (South San Francisco, CA, USA) to determine the aqueous humour content of whole CTGF and N-terminal half fragments of CTGF. These sandwich ELISAs use pairs of CTGF specific monoclonal antibodies that allow capture and detection of: (1) whole CTGF; and (2) whole CTGF+Nterminal CTGF fragments.¹⁹ Content of N-terminal fragment of CTGF was determined by subtracting the content of whole CTGF from the content of whole CTGF+N-terminal CTGF fragments (that is, (2) minus (1)). CTGF can undergo proteolysis and its bioactivity has been reported from resulting fragments.²⁰⁻²² The N-terminal fragments appear more stable than whole CTGF.¹⁹ Therefore, determining levels of both whole CTGF as well as N-terminal CTGF fragments provides an accurate measurement of total CTGF production and processing.

A range finding experiment was first conducted to determine the most suitable sample dilution. Spike recovery experiments using recombinant human CTGF (rhCTGF) have demonstrated that the CTGF sandwich ELISAs provide quantitative detection in samples of various biological fluids.¹⁹ Since the CTGF ELISA antibodies do not cross react with CCN gene family members Cyr61 and Nov, these related proteins are not detected in these assays. Controls are included to ensure that plate to plate variation is <20% and triplicate assays were performed for each sample. For the reported studies, standard curves were generated using rhCTGF and assay minimum detectable concentration was determined to be 2 ng/ml.

We also statistically analysed our data using values of: (1) whole CTGF only, (2) N-terminal fragment only, and (3) ratio of N-fragment/whole CTGF. The overall results using the above three categories are similar to the statistical analysis using the whole CTGF+N fragment values. This implies that increased expression (not fragmentation) of CTGF plays an important part in our findings. The CTGF levels reported here are the values of the whole CTGF+N-terminal CTGF fragments.

Statistics

Results of the immunoassays were expressed as mean (SEM). Kolmogorov-Smirnov test was used to determine normal distribution of the data. The group means were compared and analysed using Fisher's protected least significance

Group	Mean (SEM) (ng/ml)		
	TIMP-1	CTGF	
A:(A1+A2+A3) All PXF	9.76 (1.10)* (n = 56)	4.38 (0.65)† (n = 36)	
A1: (PXF grade I)	5.33 (0.57) (n = 9)	3.44(1.03)(n=5)	
A2: (PXF grade II)	10.34(1.47)(n = 37)	3.89 (0.65) (n = 24)	
A3: (PXF grade III)	11.61 (2.55) (n = 10)	6.74 (2.28) (n=7)	
B: Uveitis	14.92 (3.38) (n = 11)	7.07(1.98)(n = 12)	
C: Controls	5.73 (0.43) (n = 112)	2.35 (0.46) (n = 29)	
The group means were compar analysis of variance (ANOVA) *p <0.01 compare with contro †p <0.05 compare with contro	ed and analysed using Fisher's PLSD indicated a significant difference (Sta (significant for both Fisher's PLSD a l (significant only for Fisher's PLSD to	test and Bonferroni/Dunn test after one way atView, SAS Institute, Cary, NC, USA) and Bonferroni/Dunn tests). est).	

difference (Fisher's PLSD) test and Bonferroni/Dunn test after one way analysis of variance (ANOVA) identified statistically significant differences. a p value of less than 0.05 was considered to be statistically significant for Fisher's PLSD test, and likewise less than 0.0167 for Bonferroni/Dunn test (software: StatView; SAS Institute, Cary, NC, USA). The above two post hoc tests were chosen to account for the known increase rate of type I and II error in Fisher PLSD and Bonferroni/Dunn test respectively.

RESULTS

The PXF group (see table 1) showed a significantly higher TIMP-1 (9.76 (1.10) ng/ml, n = 56 v control 5.73 (0.43) ng/ ml, n = 112) and CTGF (4.38 (0.65) ng/ml, n = 36 ν control 2.35 (0.46) ng/ml, n = 29) concentrations compared with the controls, achieving p value of 0.0002 and 0.048 respectively. The higher the PXF grade the higher the TIMP-1 and CTGF levels as well as presence of glaucoma. None of the PXF grade I eyes have glaucoma, whereas over 50% of the eyes in PXF grade III group were on treatment for glaucoma. Such clinicomorphological observations of the amount of PXF material in the anterior eye segment are potential reliable indicators of disease progression and risk of glaucoma. Histologically, the amount of PXF material in the juxtacanalicular tissue and the outflow filtration area has been proved to correlate with intraocular pressure and optic nerve damage by the Lütjen-Drecoll group.23

We also compared the results under the categories of PXF without glaucoma, PXF with glaucoma, and the controls (see table 2). The PXF glaucoma group had a higher TIMP-1 level than PXF without glaucoma. However, this is not statistically significant with a p value of 0.11. The fact that seven subjects of the PXF glaucoma group were receiving topical prostaglandin analogue (latanoprost 0.005%) treatment may have a role in the disease modification. It has been documented in various studies that latanoprost can alter the expression of MMP/TIMP in human explant organ cultures,²⁴ Tenon fibroblasts,²⁵ ciliary muscle cells,²⁶ and rabbit conjunctiva.²⁷

However, the effect of latanoprost in our current study is difficult to extrapolate because most patients used more than one type of antiglaucoma medication before surgery at variable duration ranging from weeks to years. On the other hand, the CTGF concentration in the PXF glaucoma group (6.03 (1.09) ng/ml, n = 18) is significantly higher than PXF without glaucoma (2.73 (0.45) ng/ml; n = 18), achieving a p value of 0.002 (see fig 1).

We also postulated that PXF glaucoma is a consequential event from an abnormal wound healing response possibly initiated by the aggregation of PXF material in ocular tissue, leading to a low grade inflammatory response with impaired blood-aqueous barrier. As shown in table 1, the uveitis group with known defective blood-aqueous barrier also has very



Figure 1 Comparison of aqueous humour CTGF (connective tissue growth factor) levels among PXF (pseudoexfoliation syndrome) without glaucoma (PXF), treated PXF glaucoma (PXF-G), and control groups. Data were represented as mean (SEM) as depicted on the bar chart. The group means were compared and analysed using Fisher's protected least significance difference (Fisher's PLSD) test and Bonferroni/Dunn test after one way analysis of variance (ANOVA) identified statistically significant differences (software: StatView; SAS Institute, Cary, NC, USA). A p value of 0.05 (Fisher's PLSD) or 0.0167 (Bonferroni/Dunn) was considered to be statistically significant.

Group	Mean (SEM) (ng/ml)		
	TIMP-1	CTGF	
PXF without glaucoma	8.84 (1.34)* (n = 37)	2.73 (0.45) (n = 18)	
PXF with glaucoma	11.55 (1.88)* (n = 19)	6.03 (1.09)* (n = 18)	
Control	5.73 (0.43) (n = 112)	2.35(0.46)(n = 29)	

raised levels of TIMP-1 and CTGF. The above findings in the uveitis group, however, should be considered as preliminary observations as they only represent a small number of samples compared with the controls.

Apart from two samples in the uveitis group which were regarded as outliers where the MMP-9 levels were measured at 52.0 and 52.8 ng/ml respectively, the MMP-9 levels in all the other aqueous humour samples (PXF, n = 55; uveitis, n = 10; controls, n = 85) were low and below the detection limits of our commercially obtained ELISA kits.

DISCUSSION

Degradation of ECM protein is orchestrated by matrix metalloproteinases (MMPs), a family of zinc dependent endopeptidases, whose activity is in turn regulated by their physiological inhibitors known as tissue inhibitors of metalloproteinases (TIMPs). TIMPs are expressed by a wide spectrum of cells and are present in most tissue and body fluids. They also have important regulatory activities independent of MMP inhibition including the promotion of growth, alteration of cell morphology, and modulation of inflammation.²⁸²⁹ As aqueous humour is closely linked to the structural and functional integrity of the trabecular meshwork, the presence of altered MMP/TIMP levels in the aqueous humour may well affect the homeostasis of cells lining the conventional outflow routes. The expression of MMPs and their inhibitors by human trabecular meshwork cells also serves an important role in the ongoing trabecular meshwork ECM turnover and subsequent regulation of trabecular outflow resistance.^{30–33} Bradley et al demonstrated in their perfused human anterior segment organ culture study that TIMPs suppressed aqueous outflow rates.³¹ TIMP-1 is the inhibitor for MMP-9 and it can bind to both the latent and active form of the enzyme.³⁴

Our current study showed a raised TIMP-1 level in PXF aqueous humour. The total MMP-9 in PXF aqueous humour remained low or undetectable, without significant corresponding changes. This supports the hypothesis of dysregulation in the matrix metabolism in PXF where the excessive TIMP-1 prevents the destruction of newly synthesised matrix, leading to PXF material accumulation. The presence of elevated $\text{TGF}\beta_1{}^7{}^8$ as well as reduced oxygen $\hat{\text{tensions}}^{6\,35}$ in PXF aqueous humour may also contribute to the raised TIMP-1 levels, as TIMP-1 promoter contains a hypoxia response element.²⁹ Our findings are consistent with a recent paper from Schlotzer-Schrehardt *et al* showing a significantly higher concentration of TIMP-1 in the aqueous humour of PXF eyes with or without glaucoma, and very low level of MMP-9 (15-55 pg/ml) requiring an ultrasensitive ELISA for quantification in all groups with no significant differences.36 In their study, TIMP-1 levels in the PXF eyes with (n = 30)and without (n = 30) glaucoma are 2-fold and 1.8-fold higher than their controls (n = 30), respectively. Similar results are indicated in our current study where our PXF eyes with (n = 19) and without (n = 37) glaucoma are 2-fold and 1.5fold higher than the controls (n = 112). On the contrary, a smaller study conducted by Gartaganis et al found TIMP-1 levels in 19 normotensive PXF aqueous humour were slightly decreased, though not significantly so, compared to 15 normal controls.37 Nevertheless, comparisons between clinical studies are often difficult, owing to analytical and methodological (different types of ELISA kits used) variations.

CTGF is a member of an emerging family of regulatory proteins grouped together under the acronym CCN (CTGF, Cysteine-rich protein (CyR61), and Nephroblastoma overexpression gene (*Nov*)). Its overexpression is well described in various fibrotic disorders including scleroderma, renal and pulmonary fibrosis, inflammatory bowel disease, and atherosclerosis.^{16 38} CTGF has the ability to affect the ECM metabolism by directly promoting collagen synthesis and regulating the expression of MMPs and TIMPs.³⁹ Its potential role as a therapeutic target in controlling various ocular fibrotic diseases has been highlighted in recent studies related to corneal scarring,⁴⁰ ocular cicatricial pemphigoid,⁴¹ and glaucoma filtration surgery.⁴² CTGF gene expression had also been demonstrated in human trabecular meshwork cells, ciliary body, retinal vascular endothelial cells, proliferative vitreoretinopathy membranes, choroidal neovascular membranes, cataractous plaque, corneal scars, tear fluid, and pterygia.^{43–52}

Our current study showed a significantly higher CTGF concentration in the aqueous humour of PXF glaucoma patients, which is suggestive of the involvement of CTGF in the fibrotic pathology of PXF syndrome and its association with glaucoma. The presence of CTGF had been found in human aqueous humour by western blot analysis^{43 44} and ELISA.¹⁸ In their study, Van Setten *et al* reported the mean concentration of CTGF in 10 normal subjects without PXF as 1.24 (SD 0.26) ng/ml¹⁸ compared with our current study where mean level in 29 controls eyes was 2.35 (SE 0.46) ng/ml. The difference may be attributed to the different ELISA used. Furthermore, the CTGF levels reported in our study are the values of the whole CTGF plus N-terminal CTGF fragments.

There are several mechanisms that might account for the presence of certain growth factors in the aqueous humour. These include selective ultrafiltration of low molecular weight growth factors from serum, local intraocular synthesis and release of growth factors, diffusional exchange of growth factors between aqueous and vitreous humour, and selective transport systems for certain growth factors.2 The bloodaqueous barrier contributes partial control on the types and levels of growth factors that enter the aqueous humour in health and diseases. The drawbacks of our current aqueous humour study are that the raised elements found could be associated with the origin of disease development or, on the contrary, a consequential manifestation of the underlying pathology or maybe both. One could argue that it is the outflow obstruction that leads to the accumulation of TIMP-1 and CTGF. Similarly, the high levels of TIMP-1 and CTGF in the aqueous humour can be caused by excessive synthesis and give rise to the pathology.

The prevalence of PXF syndrome increases with age and its incidence doubles every decade after age 50.53 There is additional evidence recently demonstrating that PXF syndrome is a systemic condition where individuals with this condition appeared to have an increased incidence of hypertension, angina, myocardial infarction, stroke, aortic abdominal aneurysms, hyperhomocystinaemia, sensorineural hearing loss and dementia.^{54–59} Prevalence data for PXF in glaucoma populations vary considerably worldwide.⁶ A previous report from our department shows that 63% of the patients in our glaucoma clinics have PXF syndrome.60 Further work into understanding the mechanism of the development of PXF syndrome could allow us to have new therapeutic insights for treating PXF glaucoma. Regulation of MMP/TIMP expression and anti-CTGF therapy may offer potential therapeutic avenues for controlling PXF related ocular morbidity.

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