

# American Journal of Ophthalmology

## Personalized model to predict keratoconus progression from demographic, topographic and genetic data --Manuscript Draft--

<b>Manuscript Number:</b>	AJO-22-105R1
<b>Article Type:</b>	Original Article
<b>Keywords:</b>	Keratoconus; corneal cross-linking; keratoconus genetics; keratoconus prediction
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<b>Abstract:</b>	<p><b>Purpose :</b> To generate a prognostic model to predict keratoconus progression to corneal cross-linking (CXL).</p> <p><b>Design:</b> Retrospective cohort study.</p> <p><b>Methods :</b> We recruited 5025 patients (9341 eyes) with early keratoconus between January 2011 and November 2020. Genetic data from 926 patients was available. We investigated both keratometry or CXL as end-points for progression and used the Royston-Parmar method on the proportional hazards scale to generate a prognostic model. We calculated hazard ratios (HR) for each significant covariate, with explained variation and discrimination, and performed internal-external cross validation by geographic regions.</p> <p><b>Results:</b> After exclusions, model-fitting comprised 8701 eyes, of which 3232 underwent CXL. For early keratoconus, CXL provided a more robust prognostic model than keratometric progression. The final model explained 33% of the variation in time-to-event: age HR [95% confidence limits] 0.9 [0.90-0.91], maximum anterior keratometry (Kmax) 1.08 [1.07-1.09], and minimum corneal thickness 0.95 [0.93-0.96] as significant covariates. Single nucleotide polymorphisms (SNPs) associated with keratoconus (n=28) did not significantly contribute to the model. The predicted time-to-event curves closely followed the observed curves during internal-external validation. Differences in discrimination between geographic regions was low, suggesting the model maintained its predictive ability.</p> <p><b>Conclusions:</b> A prognostic model to predict keratoconus progression could aid patient empowerment, triage and service provision. Age at presentation is the most significant predictor of progression risk. Candidate SNPs associated with keratoconus do not contribute to progression risk.</p>

<b>Suggested Reviewers:</b>	<p>Marcus Ang  Marcus.Ang@singhealth.com.sg</p> <p>Alfonso Iovieno  alfonsoiovieno@hotmail.com</p> <p>Mani Bhogal  manibhogal@aol.co</p> <p>Mohammed Ziaei  mahdi207@yahoo.com</p>
<b>Opposed Reviewers:</b>	
<b>Response to Reviewers:</b>	<p>Dear Dr Parrish,</p> <p>Re: AJO-22-105, "Personalized model to predict keratoconus progression from demographic, topographic and genetic data</p> <p>We thank the reviewers for their comments and have responded to each point below. Thank you for considering our revisions.</p> <p>I confirm all the authors have reviewed the updated manuscript and responses to reviewers and agree with the changes and responses.</p> <p>I had full access to all the data in the study and takes responsibility for the integrity of the data and the accuracy of the data analysis as well as the decision to submit for publication. Thank you for your time and for considering our study and we look forward to hearing from you.</p> <p>Yours sincerely,</p> <p>Dr Daniel Gore  Consultant Ophthalmologist in Cornea and External Disease</p> <p>Responses to reviewers are found in blue.</p> <p>Reviewer #1: The study was well thought out and designed, with appropriate statistical modelling and methodology.</p> <p>1. Was multicollinearity considered or assessed in the survival models, as some of the independent variables seem highly collinear?</p> <p>Response:  Multicollinearity was considered. Correlations between keratometry variables are plotted in Supplemental Figure 2. In particular, K1 and K2 were highly correlated (<math>R^2 = 0.91</math>), thus K2 was removed from the model. This is discussed on line 241 with the following text:</p> <p>'We chose a model without K2 on the basis of parsimony, which was supported by the fact that K1 and K2 were highly correlated (<math>R^2=0.91</math>)'</p> <p>2. Are there any differences among prognostic factors across regions?</p> <p>Response:  We did not identify any significant differences in prognostic factors across regions and we have added this to the text (line 280)</p> <p>'We did not identify any significant differences in prognostic factors across regions.'</p> <p>3. Table 2: Are there any difference between the univariate and multivariable cohorts?</p>

Response:

The cohort used for univariable and multivariable analysis were identical. However, for multivariable analysis, due to the presence of missing data, the number of eyes where all covariates were available was lower than for univariable analysis. We have added this to the manuscript from line 365.

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We used 4 risk groups for showing model fit and predictive ability (Figures 1 and 3). Figure 2 uses 3 hypothetical patients (not risk groups) to demonstrate the application of the model in a clinical setting.

Reviewer #2: The authors present a very interesting and topical work, which aims to answer the question whether it is possible to predict the progression of keratoconus in the examined eyes on the basis of the measured parameters, which subsequently leads to the necessity of CXL procedure.

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Response:

We have added from line 311

"Of the significant covariates in our model, younger age made the greatest contribution to our model. Thus, one should have a lower threshold for treatment in younger patients."

We have not provided specific limits on the patient's age and keratometry as it is beyond the scope of this study. We have referenced the treatment criteria that we have applied. (Gore et al 2021) and from line 355 we have added:

'The effect of recommendations based on this model on clinical practice is yet to be evaluated.'

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Establishing more robust criteria for patient selection and prompt CXL treatment is one of the most challenging issues in keratoconus management. This manuscript aims at identifying and classifying the variables influencing keratoconus progression. The authors identified age, Kmax and minimum corneal thickness as the most significant

covariates in establishing keratoconus progression, genetic or geographic covariates did not improve the accuracy of their model. Based upon the results, the conclusions of the work are not new. However, the manuscript may deserve further revision in view of the extensive work done by the authors.

6. The weakest point of the manuscript is common to all other publications in the topic, which is the definition of keratoconus progression. The authors should discuss this issue further by revising the most recent literature in the topic.

Response:

In this study, we have chosen to use crosslinking as a surrogate for progression. This is a meaningful endpoint for patients and does not rely simply on meeting defined keratometric thresholds. We recognise that there is variability in the definition of keratoconus progression in the literature and have included some updated references on this point, which also explores the increased variability in more advanced disease, including Hashemi et al 2022, Ozalp et al 2022, Flockerzi et al 2022 and Kreps et al 2020.

We have clarified these points in the text from line 63.

‘We considered the date of numeric progression, as well as the date when CXL was performed, as alternative end-points to define keratoconus progression. Although the use of keratometry as an end-point may appear the more objective method, there is variability on the definition of progression reported in the literature and conclusions may vary with the definition that is adopted.(Vinciguerra et al. 2021; Shajari et al. 2019; Ozalp and Atalay 2022; Hashemi et al. 2022) Repeatability thresholds are not usually tailored to individual eyes (i.e. an increase in Kmax by 1 D is not significant in advanced keratoconus) although there is growing evidence on the variability of measurements in more advanced disease and the need for tailoring numerical progression definitions to the disease state, and distinguishing real progression from inherent measurement variability.(Flynn et al. 2016; Flockerzi et al. 2022; Kreps et al. 2020) ‘

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Thank you for this suggestion. However, we believe that there is value in the genetic report. Assessing the impact of SNPs associated with keratoconus on risk of progression is novel and builds on the recent findings of genetic loci significantly associated with keratoconus. (Hardcastle et al 2021 Commun Biol.) Whilst genetic data was only available for a subgroup, it is still a sizeable subgroup out of a very large number of eyes. Also, our analysis puts an upper bound on the contribution of SNPs associated with keratoconus on its progression.

The geographic data was split to perform internal external validation and is a recognised method of validating the model with the original dataset. Therefore, they are not intended to relate to ethnicity or other variables.

We have now correctly labelled Figure 2.

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generalisability of the model with an external dataset. Therefore, we have not proposed any practice guidance. We plan to test the model before making recommendations on clinical practice based on the model predictions. We have clarified this recommendation in the Discussion (line 355).

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We agree there is variability in reported outcomes related to treatment protocols and also to the definition of failure. The 98% success rate is our own data from our 2-year outcomes report (Ref 7, Gore et al 2021). We have added a range for other publications (88% - 100%) from line 45.

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## Abstract

**Purpose:** To generate a prognostic model to predict keratoconus progression to corneal cross-linking (CXL).

**Design:** Retrospective cohort study.

**Methods:** We recruited 5025 patients (9341 eyes) with early keratoconus between January 2011 and November 2020. Genetic data from 926 patients was available. We investigated both keratometry or CXL as end-points for progression and used the Royston-Parmar method on the proportional hazards scale to generate a prognostic model. We calculated hazard ratios (HR) for each significant covariate, with explained variation and discrimination, and performed internal-external cross validation by geographic regions.

**Results:** After exclusions, model-fitting comprised 8701 eyes, of which 3232 underwent CXL. For early keratoconus, CXL provided a more robust prognostic model than keratometric progression. The final model explained 33% of the variation in time-to-event: age HR [95% confidence limits] 0.9 [0.90-0.91], maximum anterior keratometry (Kmax) 1.08 [1.07-1.09], and minimum corneal thickness 0.95 [0.93-0.96] as significant covariates. Single nucleotide polymorphisms (SNPs) associated with keratoconus (n=28) did not significantly contribute to the model. The predicted time-to-event curves closely followed the observed curves during internal-external validation. Differences in discrimination between geographic regions was low, suggesting the model maintained its predictive ability.

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Editor in Chief  
*American Journal of Ophthalmology*

March 31st, 2022

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I confirm all the authors have reviewed the updated manuscript and responses to reviewers and agree with the changes and responses.

I had full access to all the data in the study and takes responsibility for the integrity of the data and the accuracy of the data analysis as well as the decision to submit for publication. Thank you for your time and for considering our study and we look forward to hearing from you.

Yours sincerely,

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Consultant Ophthalmologist in Cornea and External Disease

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**Personalized model to predict keratoconus progression from demographic, topographic and genetic data**

**Short title: Keratoconus progression**

Howard P Maile<sup>1\*</sup>; Ji-Peng Olivia Li<sup>2\*</sup>; Mary D Fortune<sup>3</sup>; Patrick Royston<sup>4</sup>;  
Marcello T Leucci<sup>2</sup>; Ismail Moghul<sup>1</sup>; Anita Szabo<sup>1</sup>; Konstantinos Balaskas<sup>2</sup>;  
Bruce D Allan<sup>2</sup>; Alison J Hardcastle<sup>1</sup>; Pirro Hysi<sup>5,6</sup>; Nikolas Pontikos<sup>1\*</sup>; Stephen J  
Tuft<sup>2\*</sup>; Daniel M Gore<sup>2\*§</sup>

<sup>1</sup>UCL Institute of Ophthalmology, 11-43 Bath Street, London EC1V 9EL

<sup>2</sup>Moorfields Eye Hospital NHS Foundation Trust, 162 City Road, London EC1V 2PD

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<sup>5</sup>Section of Ophthalmology, School of Life Course Sciences, King's College London

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\* Authors contributed equally

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Keywords: Keratoconus, corneal cross-linking, keratoconus genetics, keratoconus prediction

Page Break

## Introduction

Keratoconus is a common corneal ectasia that causes irregular astigmatism, scarring and loss of vision. Thinning and steepening can progress through childhood and early adulthood, but the shape of most eyes stabilizes by the third or fourth decade. Without intervention, keratoconus can lead to severe visual loss, with approximately 10% of eyes eventually requiring corneal transplantation.<sup>1</sup> Corneal cross-linking (CXL) by topical application of riboflavin, followed by irradiation with UV-A light, can arrest progression of keratoconus in up to 88% to 100% of eyes even when there is relatively advanced disease.<sup>2-6</sup> The potential benefit of CXL is to prevent visual deterioration with a relatively low risk procedure that is cost effective for healthcare providers.<sup>7-9</sup> However, CXL is usually not offered to all patients at presentation because the disease may have already stabilized. In the recent KERALINK study 43% of children <17 years of age at presentation had not progressed after 18 months.<sup>10</sup> The definition of progression also varies with the severity of keratoconus, but for early disease a common threshold is either an increase in the maximum keratometry (Kmax) of >1 dioptre, a change in the manifest refractive spherical equivalent of >0.50 dioptre, or an increase in manifest refractive cylinder of >1 dioptre.<sup>2,11</sup> Depending on the rate of progression this threshold may be passed in a few months, years, or not at all. At the first assessment it can be a challenge to distinguish eyes that are at risk of rapid progression from those where it is safe to monitor. Unnecessary review visits are a burden to the patient and the care system.

We considered the date of numeric progression,<sup>6</sup> as well as the date when CXL was performed, as alternative end-points to define keratoconus progression. Although the use of keratometry as an end-point may appear the more objective method, there is variability on the definition of progression reported in the literature and conclusions may vary with the definition that is adopted.<sup>11-14</sup> Repeatability thresholds are not usually tailored to individual eyes (i.e. an

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70 evidence on the variability of measurements in more advanced disease and the  
71 need for tailoring numerical progression definitions to the disease state, and  
72 distinguishing real progression from inherent variability of measurement  
73 modalities.<sup>15-17</sup> Finally, patients who receive CXL prior to progression must be  
74 censored from the dataset even though these eyes are likely to have been at  
75 risk of progression. This type of informative censoring creates a bias.<sup>18</sup> In  
76 contrast, the time to CXL depends on several variables that include numeric  
77 disease progression, but also incorporates patient-specific risk factors for future  
78 progression. Its strength is that it is an easily comprehensible and meaningful  
79 end-point for patients. It encompasses individual risk factors that are not  
80 considered when imaging is used in isolation and it has been used by others as  
81 defining the event of interest.<sup>19</sup>

82  
83 For these reasons we have used demographic and serial tomography data from  
84 a large cohort of patients to generate a time-to-event model to predict the  
85 probability of an individual progressing to CXL. Because the Cox proportional  
86 hazards method does not generate smooth time-to-event curves, we used the  
87 Royston-Parmar model to achieve direction estimates of the hazard function.<sup>20</sup>  
88 We also performed a further analysis of a subset of patients who had genetic  
89 data in the form of single-nucleotide polymorphisms (SNP) generated as part of  
90 a study to determine keratoconus risk.<sup>21</sup>

## 91 92 **Methods**

### 93 **Cohort**

94 The study protocol was reviewed and approved by the Clinical Audit  
95 Assessment Committee of Moorfields Eye Hospital NHS Foundation Trust  
96 (reference CA17/CED/03). Institutional Review Board (IRB) approval was  
97 obtained and individual patient consent was not required. The study conformed  
98 to the tenets of the Declaration of Helsinki. We identified from the Moorfields  
99 Eye Hospital electronic health record database (OpenEyes) patients aged 13  
100 years and above diagnosed with clinical or suspected keratoconus who  
101 attended our Early Keratoconus Clinic (EKC) between January 2011 and  
102 November 2020. Clinical data included keratometry (Kmax, Front K1, Front K2,

103 Back K1, Back K2), and pachymetry (minimum corneal thickness) captured by  
104 Scheimpflug tomography (Pentacam HR, Oculus GmbH, Wetzlar). We only  
105 included scans with a quality score of 'good' or 'ok', and where multiple scans  
106 were taken on the same day we used the mean value. The date of all CXL  
107 procedures was recorded. The protocol for offering CXL throughout the study  
108 period was, i) a documented history prior to referral to the EKC of our hospital  
109 of significant recent disease progression,<sup>6</sup> ii) a change in contemporary  
110 measurements of 95% above the repeatability limits of the baseline  
111 measurements as shown in Supplemental Table 1 (available at  
112 <http://www.ajo.com>),<sup>6</sup> or iii) a patient considered by a clinician to be at high risk  
113 of progression despite their not fulfilling the above two criteria. Exclusion criteria  
114 included pregnancy or breastfeeding, uncontrolled ocular surface disease or a  
115 minimum corneal thickness less than 375 µm.

116  
117 All the data used for model fitting started from the first appointment in the EKC.  
118 Patient demographics included age, gender, smoking status (current or ex/non-  
119 smoker) ethnicity and postcode. Ethnicity was coded as 1 for 'Black' or 'South  
120 Asian or South Asian British' and 0 for any other category (excluding missing  
121 values). Before model fitting, the pachymetry in microns was divided by 10 to  
122 generate a meaningful scale. For the primary analysis, eyes with any missing  
123 data were excluded. We also explored multiple imputation, which avoids data  
124 exclusion by generating multiple versions of the dataset with missing values  
125 replaced with values sampled from an appropriate distribution. To see whether  
126 genetic data can help predict keratoconus progression, we used 28 candidate  
127 SNPs from a recent keratoconus genome-wide association study that contained  
128 926 patients from Moorfields Eye Hospital.<sup>21</sup> The SNP data was encoded as  
129 either 0 (homozygous reference genotype), 1 (heterozygous genotype), or 2  
130 (homozygous variant genotype). We chose to use an additive encoding, thus  
131 the risk of disease increases additively with the degree of genetic variation.<sup>22</sup>  
132 Anonymized data were then exported to Excel software for analysis (version  
133 15.24 2016, Microsoft Corp.).

## 134 135 **Model Fitting and Covariate Selection**



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136 A Royston-Parmar flexible parametric survival model was fitted to the data to  
137 predict the probability of an eye progressing to CXL.<sup>23</sup> Initial analysis of the  
138 covariates was performed by univariate analysis using the same model  
139 characteristics as the multivariable model. When selecting covariates for the  
140 final multivariable model, we used backwards stepwise selection with a  
141 significance level of 0.05. We used linear covariates for ease of interpretation of  
142 our final model. To create a more parsimonious model we examined the effect  
143 on explained variation and discrimination of removing single variables from the  
144 model.

### 145 146 **Keratometric Progression Sensitivity Analysis**

147 We included a sensitivity analysis in which we investigated keratometric  
148 progression as an alternative end-point. Keratometric progression was defined  
149 using thresholds from Gore et al 2021.<sup>6</sup> When using numerical thresholds to  
150 define progression, the appointments for eyes beyond the date of CXL cannot  
151 be used. However, censoring these eyes at the date of CXL represents  
152 informative censoring. Based on the recommendations of Clarke et al<sup>18</sup> for  
153 investigating the impact of informative censoring, we generated a ‘best case’  
154 dataset where eyes were censored at the CXL date and a ‘worst case’ dataset  
155 where patients were assumed to progress at the CXL date. The corresponding  
156 Kaplan Meier curves were plotted to provide a visual comparison of the two  
157 datasets. A Royston-Parmar model was then fitted on both datasets. We used  
158 the same techniques (backwards stepwise selection, significance level of 0.05)  
159 as described in the previous section to fit the model and compare the explained  
160 variation and hazard ratios.

### 161 162 **Multivariable Model Validation**

163 We validated the model using internal-external cross validation in which we split  
164 the dataset by geographical region.<sup>24,25</sup> For the kth region, the model is fitted on  
165 the full dataset excluding region k and then Kaplan-Meier curves and predicted  
166 survival curves were generated for region k. Seven geographical regions were  
167 created based on the patient’s postcode as shown in Supplemental Figure 1  
168 (available at <http://www.ajo.com>). To quantitatively assess the validation,  
169 Royston and Sauerbrei’s D statistic was calculated for both the model fitted



170 from data excluding region  $k$  ( $D_{(k)}$ ) and also the model applied to region  $k$  ( $D_k$ ).<sup>26</sup>  
171 The difference between these two discrimination metrics ( $D_k - D_{(k)}$ ) was calculated  
172 with its corresponding standard error to assess the predictive ability of the  
173 model. To demonstrate how the model could be used in practice, we include  
174 three hypothetical patients' eyes with different progression risk profiles (high,  
175 medium, low risk) and plot the predicted time-to-event curve for each shown in  
176 Figure 2.

## 178 **Statistical Analysis**

179 The event of interest was defined as the date that the eye underwent CXL. We  
180 calculated the time-to-event as the difference between the first appointment in  
181 our service and the date of CXL (or the last patient appointment in the case of  
182 censoring). Since we had paired observations (eyes), we used variance-  
183 corrected models to account for correlation between eyes and to ensure that  
184 robust standard errors were produced. The choice of scale and selection of  
185 degrees of freedom for the Royston-Parmar model was informed by inspecting  
186 the Akaike information criterion (AIC) and Bayes information criterion (BIC)<sup>20</sup>  
187 and the results of this were balanced with ease of interpretation. See  
188 Supplemental Table 2 and Supplemental Text 1 (available at  
189 <http://www.ajo.com>) for further explanation. Royston and Sauerbrei's D statistic  
190 was used as a measure of discrimination and  $R^2_D$  as a measure of explained  
191 variation (both calculated on the natural scale of the model). Although all of the  
192 primary results were generated from a complete case analysis, we performed  
193 an additional analysis using multiple chained imputation (predictive mean  
194 matching approach with 5 nearest neighbors). Model fitting was performed in  
195 Stata 13 (StataCorp LP, Texas, USA) and the Royston-Parmar model was fitted  
196 using the stpm2 package from Stata 13.

## 198 **Results**

### 199 **Cohort**

200 From a potential of 9,341 eyes (4316 pairs of eyes and 709 individual eyes), the  
201 final model used 8,701 eyes of 4,823 patients, with 3,232 eyes that had CXL.  
202 The mean age was 28.3 years with standard deviation of 7.1 years. We  
203 excluded 640 eyes with missing data. Table 1 summarizes the available

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204 covariates along with missing data percentages. See Supplemental Text 2 and  
205 Supplemental Table 3 (available at <http://www.ajo.com>) for a description of the  
206 multiple imputation results.

### 207 208 **Model Fitting and Covariate Selection (Genetic Data)**

209 We analyzed patients with genetic data separately because this data was only  
210 available for ~14% of patients. Of 926 patients (1852 eyes) with genetic data,  
211 531 eyes were excluded with incomplete keratometry or CXL data, which left  
212 1321 eyes, of which 665 had CXL. With univariate analysis of the 28 SNPs only  
213 rs72631889 was found to be significant ( $P=0.01$ ) (Supplemental Table 4  
214 (available at <http://www.ajo.com>)). We then produced a multivariable model via  
215 backwards selection on this subset of eyes using corneal data, patient data and  
216 rs72631889 as an additional covariate as shown in Supplemental Table 5  
217 (available at <http://www.ajo.com>). However rs72631889, although significant  
218 ( $P=0.005$ ), had a negligible contribution (0.3%) to the explained variation in the  
219 final model.

### 220 221 **Model Fitting and Covariate Selection (Excluding Genetic Data)**

222  
223 The results of the univariate time-to-event analysis on the hazards scale using  
224 a Royston-Parmar flexible parametric model is shown in Table 2. Genetic data  
225 was excluded from this analysis. All variables except smoking status were  
226 significant. The explained variation ( $R^2_D$ ) and discrimination ( $D$ ) were highest for  
227 age (17%) and Kmax (15%) with Front K1, Front K2, Back K1, Back K2 and  
228 pachymetry each explaining 6-10% of the variation. Notably, gender and  
229 ethnicity, although significant in the univariate analysis, did not contribute to  
230 explained variation. The hazard ratios for significant covariates indicate that  
231 increasing age at presentation, greater pachymetry and flatter (less negative)  
232 posterior keratometry values decrease risk of having CXL, whilst steeper  
233 anterior keratometry values and male gender increase the risk of having CXL.

234  
235 When we fitted a multivariable model the significant covariates were age,  
236 Kmax, Front K1, Front K2 and pachymetry (Table 2). When we removed single  
237 variables from the model the effect this had on explained variation and

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238 discrimination is shown in Supplemental Table 6 (available at  
239 <http://www.ajo.com>). Age was the most important covariate (16.7%), with Kmax  
240 contributing ~5% of explained variation. K1, K2 and pachymetry had a small  
241 effect (<1%) when removed individually. We chose a model without K2 on the  
242 basis of parsimony, which was supported by the fact that K1 and K2 were  
243 highly correlated ( $R^2=0.91$ ) as shown in Supplemental Figure 2 (available at  
244 <http://www.ajo.com>). The final fitted model hazard ratios can be seen on the  
245 multivariable column of Table 2. It is notable that an increase in K1 now has a  
246 protective effect in the final model. The explained variation and discrimination  
247 for the final model were 32.7% and 1.43 respectively.<sup>27</sup> The opposing effect of  
248 Kmax and Front K1 can be explained by examining their regression coefficients  
249 before converting to hazard ratios; Kmax has a positive coefficient (0.0795) and  
250 Front K1 has a negative coefficient (-0.0749). This is logically similar to  
251 including the combined covariate (Kmax - Front K1) in the model which can be  
252 viewed clinically as a proxy for irregular astigmatism. We also investigated  
253 combining K1 and K2 into a single covariate as K2-K1 (standard definition of  
254 astigmatism), but the corresponding p value was not significant.

255  
256 Figure 1 visually depicts the result of applying the final model to the original  
257 dataset. As expected, the predicted mean survival curves closely follow the  
258 Kaplan-Meier curves. To demonstrate the use of the model in clinical practice,  
259 survival curves for three hypothetical patients followed for five years are shown  
260 in Figure 2. We have also produced a web application from the model which  
261 can be accessed at <http://beta.moorfieldscxl.com>.

### 262 263 **Keratometric Progression Sensitivity Analysis**

264 The results of the keratometric progression sensitivity analysis can be found in  
265 the Supplementary Material. By examining the Kaplan Meier curves in  
266 Supplemental Figure 3, we can see that the best case time-to-event curve  
267 indicates a 40% survival probability at 5 years whilst the worst case curve  
268 indicates a 27% survival probability at 5 years. This 13% difference in survival  
269 probability at 5 years represents the upper bound of the discrepancy in survival  
270 probability within the data. After fitting the Royston-Parmar model, amongst the  
271 hazard ratios which overlap (age, Kmax, k2), there was reasonable similarity

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272 (Supplemental Tables 8 and 9). Most importantly, the model fitted to the best  
273 case had an explained variation of 11% compared to 23% for the worst case  
274 indicating a significant difference in model performance depending on the  
275 assumptions used for handling eyes which received CXL.

276

### 277 **Multivariable Model Validation**

278 When performing validation using internal-external cross validation, Figure 3  
279 shows the ability of our final model to predict keratoconus progression across  
280 different geographic regions. We did not identify any significant differences in  
281 prognostic factors across regions. The model prediction curves generally follow  
282 the Kaplan Maier curves. Notably, region 5 (South West Greater London) and  
283 region 7 (other regions) have a worse predictive performance than the other  
284 regions, indicating that these regions have different characteristics compared  
285 with the remainder of the dataset used for model fitting. This could be due to  
286 differing patient characteristics, such as complex cases that required referral to  
287 our tertiary referral centre rather than being managed locally. Overall, the  
288 prediction becomes less accurate over time, which is expected due to low  
289 numbers with follow-up beyond three years. Supplemental Table 7 displays  
290 quantitative validation results of the model using internal external validation.  
291 The difference column  $D_k - D_{(k)}$  is a measure of predictive ability. Region 7 (other  
292 regions outside of Greater London) has the greatest discrepancy in  
293 discrimination (-0.26) which indicates that the model fitted when excluding  
294 region 7 had greater discriminative ability than when applied to region 7 alone.

295

### 296 **Discussion**

297

298 In this study we have incorporated demographic, keratometric, and genetic data  
299 to generate a prognostic model of keratoconus progression to CXL. We have  
300 shown that parameters recorded at the first examination (age, Kmax, Front K1,  
301 minimum pachymetry) can produce a time-to-event curve to calculate a  
302 personalized risk for keratoconus progression. Although we chose time to CXL  
303 rather than keratometric progression as the end point for the time-to-event  
304 analysis, we performed a sensitivity analysis using keratometric progression,  
305 and found that a CXL model accounts for a much higher proportion of the

1 306 explained variation (33%) compared to the keratometric model (11% or 23% for  
2 307 best and worst case respectively). The opposing effects of Kmax and Front K1  
3 308 were unexpected, but similar to including the combined covariate (Kmax - Front  
4 309 K1) in the model; a possible explanation is that the opposing effect is the result  
5 310 of an increase in irregular astigmatism. Of the significant covariates in our  
6 311 model, younger age made the greatest contribution to our model. Thus, one  
7 312 should have a lower threshold for treatment in younger patients.

8 313 When applying internal-external cross validation, the survival curves closely  
9 314 followed the Kaplan Meier survival curves for each of the geographic regions,  
10 315 which indicates generalisability, and model discrimination between training and  
11 316 cross validation groups was similar, indicating that the predictive ability is well  
12 317 maintained. Finally, our SNP genetic data had limited additional predictive utility  
13 318 for keratoconus progression. However, the genetic dataset was relatively small  
14 319 (926 patients), and recruitment was based on the presence of keratoconus, as  
15 320 opposed to the severity of keratoconus, or any other index of risk of rapid  
16 321 progression.

17 322  
18 323 The Royston-Parmar model has previously been used to predict the likelihood  
19 324 of the worst eye of patients with keratoconus progressing to corneal  
20 325 transplantation.<sup>28</sup> In their final model, Quartilho et al chose 3 significant  
21 326 covariates: Kmax, age and ethnicity. The reported covariate hazard ratios that  
22 327 overlap with our study (Kmax and age) were different in magnitude but in the  
23 328 same direction. When performing internal validation their model exhibited good  
24 329 predictive ability. They produced time-dependent receiver operating curves  
25 330 using the validation set and found one-year sensitivity and specificity to be  
26 331 92.8% and 94.6% respectively. Using logistic regression, Kato et al. found that  
27 332 the two strongest factors associated with the requirement for CXL were age and  
28 333 Kmax, which is consistent with our findings.<sup>19</sup> Moreover the team went on to find  
29 334 that age combined with corneal tomography maps was able to predict  
30 335 progression and need for crosslinking using deep learning.<sup>29</sup>

31 336  
32 337 An ability to generate personalized time-to-event curves that predict  
33 338 progression to CXL (Figure 2) could directly inform clinical decisions that benefit  
34 339 patient care. Firstly, patients may better understand their own risk for

1 340 progression and feel more confident in choosing their management options.  
2 341 Secondly, for both clinicians and patients, the prediction of progression may  
3 342 contribute to scheduling treatments, including prioritizing patients at high risk of  
4 343 early progression. For example, patients at high risk with a 98% probability of  
5 344 progressing to CXL at 5 years could be offered CXL at the point of first  
6 345 diagnosis without waiting to demonstrate keratometric progression. Medium risk  
7 346 patients may benefit from a period of clinician-led topographic monitoring. For  
8 347 the lowest risk patients, optometry-led monitoring in the community may be  
9 348 sufficient. This risk stratification could be tailored to regions and reflect local  
10 349 needs and resources such as provision of monitoring services in regions with  
11 350 lower risk and greater capacity for CXL in areas with more high risk patients.  
12 351 Finally, when a decision is made to postpone CXL for further monitoring, the  
13 352 time-to-event curve can contribute to decisions on the scheduling of future  
14 353 follow up reviews, with perhaps shorter time periods where the curve is  
15 354 steepest. Recommendations based on this model on clinical practice is yet to  
16 355 be evaluated.

17 356  
18 357 Our study is subject to several limitations inherent to our dataset. First, if  
19 358 patients had CXL at another hospital, this may not be reliably recorded in the  
20 359 source database. This could lead to a very small number of patients being  
21 360 included in the analysis who have already had CXL. Second, ethnicity is a well  
22 361 established risk factor for keratoconus and keratoconus progression,<sup>27,30,31</sup> but  
23 362 ethnicity is now an optional field at patient registration at our institution and this  
24 363 information was unavailable for approximately 50% of our dataset. However,  
25 364 even when we restricted the dataset to those with ethnicity records, it was not  
26 365 found to be a significant covariate. Third, though the cohort used for univariable  
27 366 and multivariable analysis were identical the number of eyes where all  
28 367 covariates were available was lower than for univariable analysis due to  
29 368 missing data. Finally, when we used multiple imputation to generate a  
30 369 multivariable model, ethnicity was still not found to be significant. In the model  
31 370 fitting process we chose to use a simple backwards selection as opposed to the  
32 371 multivariate fractional polynomial (MFP) method.<sup>32</sup> In our initial investigations,  
33 372 the results of MFP yielded nonlinear functional forms of the covariates and,  
34 373 whilst this method may have slightly increased the predictive power of the



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374 prognostic model, the resulting hazard ratios would be very hard to interpret. In  
375 addition, we did not examine time dependent effects for the covariates, which  
376 may provide a more accurate model fit, and future studies should examine this  
377 option. Finally, although no external validation dataset was available, internal  
378 external cross validation allowed us to confirm that our model is generalizable  
379 across geographical regions.

380

381 In conclusion, we have fitted a prognostic model for progression of keratoconus  
382 to CXL which generates a time-to-event curve using age, Kmax, Front K1,  
383 minimum pachymetry from time of presentation. Incorporation of a relatively  
384 small genetic dataset does not improve the explained variation of our model.  
385 Personalized modeling of risk may improve patients' understanding of their  
386 condition and the need for CXL. Such a model may help better improve patients  
387 and aid clinician decision making to CXL to achieve better outcomes and  
388 judicious use of healthcare resources.

389

#### 390 **Disclosures**

391 The authors have no financial disclosures.

392

393 Funding/Support: HM is funded by a Moorfields Eye Charity PhD Studentship  
394 (GR001147). NP is funded by a Moorfields Eye Charity Career Development  
395 Award (R190031A). Moorfields Eye Charity is supported in part by the National  
396 Institute for Health Research (NIHR) Biomedical Research Centre based at  
397 Moorfields Eye Hospital NHS Foundation Trust and UCL Institute of  
398 Ophthalmology. ST, BA and DG acknowledge that a proportion of their financial  
399 support is from the Department of Health through the award made by the  
400 National Institute for Health Research to Moorfields Eye Hospital NHS  
401 Foundation Trust and University College London Institute of Ophthalmology for  
402 a Specialist Biomedical Research Centre for Ophthalmology. The sponsor or  
403 funding organization had no role in the design or conduct of this research.

404

405 Other acknowledgements: none.

406

#### 407 **References**

- 1  
2  
3 409 1. Gordon MO, Steger-May K, Szczotka-Flynn L, et al. Baseline factors  
4  
5 410 predictive of incident penetrating keratoplasty in keratoconus. *Am J*  
6  
7 411 *Ophthalmol.* 2006;142(6):923-930.
- 8  
9 412 2. Wittig-Silva C, Chan E, Islam FMA, Wu T, Whiting M, Snibson GR. A  
10  
11 413 randomized, controlled trial of corneal collagen cross-linking in progressive  
12  
13 414 keratoconus: three-year results. *Ophthalmology.* 2014;121(4):812-821.
- 14  
15 415 3. Caporossi A, Mazzotta C, Baiocchi S, Caporossi T. Long-term results of  
16  
17 416 riboflavin ultraviolet A corneal collagen cross-linking for keratoconus in Italy:  
18  
19 417 the Siena eye cross study. *Am J Ophthalmol.* 2010;149(4):585-593.
- 20  
21  
22 418 4. O'Brart DPS, Chan E, Samaras K, Patel P, Shah SP. A randomised,  
23  
24 419 prospective study to investigate the efficacy of riboflavin/ultraviolet A (370  
25  
26 420 nm) corneal collagen cross-linkage to halt the progression of keratoconus.  
27  
28 421 *Br J Ophthalmol.* 2011;95(11):1519-1524.
- 29  
30  
31 422 5. Koller T, Mrochen M, Seiler T. Complication and failure rates after corneal  
32  
33 423 crosslinking. *J Cataract Refract Surg.* 2009;35(8):1358-1362.
- 34  
35 424 6. Gore DM, Leucci MT, Koay SY, et al. Accelerated Pulsed High-Fluence  
36  
37 425 Corneal Cross-Linking for Progressive Keratoconus. *Am J Ophthalmol.*  
38  
39 426 2021;221:9-16.
- 40  
41  
42 427 7. Salmon HA, Chalk D, Stein K, Frost NA. Cost effectiveness of collagen  
43  
44 428 crosslinking for progressive keratoconus in the UK NHS. *Eye* .  
45  
46 429 2015;29(11):1504-1511.
- 47  
48  
49 430 8. Lindstrom RL, Berdahl JP, Donnenfeld ED, et al. Corneal cross-linking  
50  
51 431 versus conventional management for keratoconus: a lifetime economic  
52  
53 432 model. *J Med Econ.* Published online November 19, 2020:1.
- 54  
55 433 9. Godefrooij DA, Mangen MJJ, Chan E, et al. Cost-Effectiveness Analysis of  
56  
57 434 Corneal Collagen Crosslinking for Progressive Keratoconus.  
58  
59 435 *Ophthalmology.* 2017;124(10):1485-1495.

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- 436 10. Larkin DFP, Chowdhury K, Burr JM, et al. Effect of Corneal Cross-linking  
437 versus Standard Care on Keratoconus Progression in Young Patients: The  
438 KERALINK Randomized Controlled Trial. *Ophthalmology*. Published online  
439 April 20, 2021. doi:10.1016/j.ophtha.2021.04.019
- 440 11. Vinciguerra R, Belin MW, Borgia A, et al. Evaluating keratoconus  
441 progression prior to crosslinking: maximum keratometry vs the ABCD  
442 grading system. *J Cataract Refract Surg*. 2021;47(1):33-39.
- 443 12. Shajari M, Steinwender G, Herrmann K, et al. Evaluation of keratoconus  
444 progression. *Br J Ophthalmol*. 2019;103(4):551-557.
- 445 13. Ozalp O, Atalay E. Belin ABCD Progression Display Identifies Keratoconus  
446 Progression Earlier Than Conventional Metrics. *Am J Ophthalmol*.  
447 2022;236:45-52.
- 448 14. Hashemi H, Panahi P, Asgari S, Emamian MH, Mehravaran S, Fotouhi A.  
449 Best Indicators for Detecting Keratoconus Progression in Children: A Report  
450 From the Shahroud Schoolchildren Eye Cohort Study. *Cornea*.  
451 2022;41(4):450-455.
- 452 15. Flynn TH, Sharma DP, Bunce C, Wilkins MR. Differential precision of  
453 corneal Pentacam HR measurements in early and advanced keratoconus.  
454 *Br J Ophthalmol*. 2016;100(9):1183-1187.
- 455 16. Flockerzi E, Häfner L, Xanthopoulou K, et al. Reliability analysis of  
456 successive Corneal Visualization Scheimpflug Technology measurements in  
457 different keratoconus stages. *Acta Ophthalmol*. 2022;100(1):e83-e90.
- 458 17. Kreps EO, Jimenez-Garcia M, Issarti I, Claerhout I, Koppen C, Rozema JJ.  
459 Repeatability of the Pentacam HR in Various Grades of Keratoconus. *Am J*  
460 *Ophthalmol*. 2020;219:154-162.
- 461 18. Clark TG, Bradburn MJ, Love SB, Altman DG. Survival analysis part IV:  
462 further concepts and methods in survival analysis. *Br J Cancer*.  
463 2003;89(5):781-786.

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- 464 19. Kato N, Negishi K, Sakai C, Tsubota K. Baseline factors predicting the need  
465 for corneal crosslinking in patients with keratoconus. *PLoS One*.  
466 2020;15(4):e0231439.
- 467 20. Patrick Royston PL. *Flexible Parametric Survival Analysis Using Stata:  
468 Beyond the Cox Model*. Stata Press; 2011.
- 469 21. Hardcastle AJ, Liskova P, Bykhovskaya Y, et al. A multi-ethnic genome-wide  
470 association study implicates collagen matrix integrity and cell differentiation  
471 pathways in keratoconus. *Commun Biol*. 2021;4(1):266.
- 472 22. Ding X, Guo X. A Survey of SNP Data Analysis. *Big Data Mining and  
473 Analytics*. 2018;1(3):173-190.
- 474 23. Royston P, Parmar MKB. Flexible parametric proportional-hazards and  
475 proportional-odds models for censored survival data, with application to  
476 prognostic modelling and estimation of treatment effects. *Stat Med*.  
477 2002;21(15):2175-2197.
- 478 24. Baade PD, Royston P, Youl PH, Weinstock MA, Geller A, Aitken JF.  
479 Prognostic survival model for people diagnosed with invasive cutaneous  
480 melanoma. *BMC Cancer*. 2015;15:27.
- 481 25. Royston P, Parmar MKB, Sylvester R. Construction and validation of a  
482 prognostic model across several studies, with an application in superficial  
483 bladder cancer: CONSTRUCTION AND VALIDATION OF PROGNOSTIC  
484 MODEL. *Stat Med*. 2004;23(6):907-926.
- 485 26. Royston P, Sauerbrei W. A new measure of prognostic separation in  
486 survival data. *Stat Med*. 2004;23(5):723-748.
- 487 27. Tuft SJ, Moodaley LC, Gregory WM, Davison CR, Buckley RJ. Prognostic  
488 factors for the progression of keratoconus. *Ophthalmology*.  
489 1994;101(3):439-447.

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- 490 28. Quartilho A, Gore DM, Bunce C, Tuft SJ. Royston–Parmar flexible  
491 parametric survival model to predict the probability of keratoconus  
492 progression to corneal transplantation. *Eye* . 2020;34(4):657-662.
- 493 29. Kato N, Masumoto H, Tanabe M, et al. Predicting Keratoconus Progression  
494 and Need for Corneal Crosslinking Using Deep Learning. *J Clin Med Res*.  
495 2021;10(4). doi:10.3390/jcm10040844
- 496 30. Pearson AR, Soneji B, Sarvananthan N. Does ethnic origin. *Eye* .  
497 2000;14:625-628.
- 498 31. Georgiou T, Funnell CL, Cassels-Brown A, O’Conor R. Influence of ethnic  
499 origin on the incidence of keratoconus and associated atopic disease in  
500 Asians and white patients. *Eye* . 2004;18(4):379-383.
- 501 32. Royston P. *Multivariable Model-Building: A Pragmatic Approach to*  
502 *Regression Analysis Based on Fractional Polynomials for Continuous*  
503 *Variables*. John Wiley; 2008.
- 504  
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507 **Legend**

508

509 **Figure 1:** Chart showing how the Royston-Parmar model fits the entire dataset.

510 We split the eyes into 4 risk groups by their prognostic index: <25th centile (low  
511 risk), 25-50th centile (medium-low risk), 50-75th centile (medium-high  
512 risk), >75th centile (high risk). The number of eyes at risk corresponds to the  
513 Kaplan-Meier curves.

514

515 **Figure 2:** Time-to-event curves that predict the risk of progression to CXL for

516 three hypothetical patient profiles. The blue line represents a high risk patient

517 who has a 95% probability of progressing to CXL at 5 years. The red line is a

518 medium risk patient who has a 48% probability of progressing to CXL at 5

519 years. The green line is a low risk patient who has a 14% probability of

520 progressing to CXL at 5 years. The equation used to generate the curves is:

521  $S(t) = e^{-H(t)}$ , where  $H(t)$  is the cumulative hazard function and is commonly

522 expressed as  $\ln(H(t)) = s(\ln(t)) + x\beta$ , where  $s(\ln(t))$  is a restricted cubic

523 spline function of log time,  $\beta$  is the vector of coefficients and  $x$  is the vector of

524 covariates. For further details of the derivation, we refer the reader to <sup>20</sup>.

525 Abbreviations: pachy, pachymetry

526

527 **Figure 3:** Predicted and observed survival curves for seven postal code regions

528 of Greater London as shown in Supplemental Figure 1 (available at

529 <http://www.ajo.com>) using IECV. We split the eyes into 4 risk groups by their

530 prognostic index: <25th centile (low risk), 25-50th centile (medium-low risk), 50-

531 75th centile (medium-high risk), >75th centile (high risk).

532

533 Abbreviations:

534 AIC: Akaike Information Criterion

535 Back K1: Flat posterior keratometry in the central 3 mm zone

1	536	Back K2: Steep posterior keratometry in the central 3 mm zone
2	537	BIC: Bayes Information Criterion
3	538	CXL: Corneal Cross-Linking
4	539	EKC: Early Keratoconus Clinic
5	540	EPR: Electronic Patient Record
6	541	Front K1: Flat anterior keratometry in the central 3 mm zone
7	542	Front K2: Steep anterior keratometry in the central 3 mm zone
8	543	HR: Hazard Ratio
9	544	IECV: Internal-external Cross Validation
10	545	Kmax: Maximum anterior keratometry
11	546	MFP: Multivariate Fractional Polynomial
12	547	SNP: Single-nucleotide Polymorphism
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**Personalized model to predict keratoconus progression from demographic, topographic and genetic data**

**Short title: Keratoconus progression**

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41 0207 253 3411. ▲  
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43 Keywords: Keratoconus, corneal cross-linking, keratoconus genetics,  
44 keratoconus prediction ▲

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**Introduction**

Keratoconus is a common corneal ectasia that causes irregular astigmatism, scarring and loss of vision. Thinning and steepening can progress through childhood and early adulthood, but the shape of most eyes stabilizes by the third or fourth decade. Without intervention, keratoconus can lead to severe visual loss, with approximately 10% of eyes eventually requiring corneal transplantation.<sup>1</sup> ~~Corneal collagen~~ Corneal cross-linking (CXL) by topical application of riboflavin, followed by irradiation with UV-A light, can arrest progression of keratoconus in up to ~~9888% to 100%~~ 100% of eyes even when there is relatively advanced disease.<sup>2-8,28</sup> The potential benefit of CXL is to prevent visual deterioration with a relatively low risk procedure that is cost effective for healthcare providers.<sup>9-11,29</sup> However, CXL is usually not offered to all patients at presentation because the disease may have already stabilized. In the recent KERALINK study 43% of children <17 years of age at presentation had not progressed after 18 months.<sup>8,10</sup> The definition of progression also varies with the severity of keratoconus, but for early disease a common threshold is either an increase in the maximum keratometry (Kmax) of >1 dioptre, a change in the manifest refractive spherical equivalent of >0.50 dioptre, or an increase in manifest refractive cylinder of >1 dioptre.<sup>2,12,11</sup> Depending on the rate of progression this threshold may be passed in a few months, years, or not at all. At the first assessment it can be a challenge to distinguish eyes that are at risk of rapid progression from those where it is safe to monitor. Unnecessary review visits are a burden to the patient and the care system.

We considered the date of numeric progression,<sup>7,8</sup> as well as the date when CXL was performed, as alternative end-points to define keratoconus progression. Although the use of keratometry as an end-point may appear the more objective method, there is ~~no consensus variability~~ on the definition of progression reported in the literature and conclusions may vary with the definition that is adopted.<sup>12,13,11-14</sup> Repeatability thresholds are not usually tailored to individual eyes (i.e. an increase in Kmax by 1 D is not significant in all eyes) ~~and other relevant patient specific factors are excluded although there is~~

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80 growing evidence on the variability of measurements in more advanced disease  
81 and the need for tailoring numerical progression definitions to the disease state,  
82 and distinguishing real progression from inherent variability of measurement  
83 modalities.<sup>44,16-17</sup> Finally, patients who receive CXL prior to progression must be  
84 censored from the dataset even though these eyes are likely to have been at  
85 risk of progression. This type of informative censoring creates a bias.<sup>46,18</sup> In  
86 contrast, the time to CXL depends on several variables that include numeric  
87 disease progression, but also incorporates patient-specific risk factors for future  
88 progression. Its strength is that it is an easily comprehensible and meaningful  
89 end-point for patients. It encompasses individual risk factors that are not  
90 considered when imaging is used in isolation and it has been used by others as  
91 defining the event of interest.<sup>46,19</sup>

92  
93 For these reasons we have used demographic and serial tomography data from  
94 a large cohort of patients to generate a time-to-event model to predict the  
95 probability of an individual progressing to CXL. Because the Cox proportional  
96 hazards method does not generate smooth time-to-event curves, we used the  
97 Royston-Parmar model to achieve direction estimates of the hazard function.<sup>47,20</sup>  
98 We also performed a further analysis of a subset of patients who had genetic  
99 data in the form of single-nucleotide polymorphisms (SNP) generated as part of  
100 a study to determine keratoconus risk.<sup>48,21</sup>

101  
102 **Methods**  
103 **Cohort**

104 The study protocol was reviewed and approved by the Clinical Audit  
105 Assessment Committee of Moorfields Eye Hospital NHS Foundation Trust  
106 (reference CA17/CED/03). Institutional Review Board (IRB) approval was  
107 obtained and individual patient consent was not required. The study conformed  
108 to the tenets of the Declaration of Helsinki. We identified from the Moorfields  
109 Eye Hospital electronic health record database (OpenEyes) patients aged 13  
110 years and above diagnosed with clinical or suspected keratoconus who  
111 attended our Early Keratoconus Clinic (EKC) between January 2011 and  
112 November 2020. Clinical data included keratometry (Kmax, Front K1, Front K2,  
113 Back K1, Back K2), and pachymetry (minimum corneal thickness) captured by

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114 Scheimpflug tomography (Pentacam HR, Oculus GmbH, Wetzlar). We only  
115 included scans with a quality score of 'good' or 'ok', and where multiple scans  
116 were taken on the same day we used the mean value. The date of all CXL  
117 procedures was recorded. The protocol for offering CXL throughout the study  
118 period was, i) a documented history prior to referral to the EKC of our hospital  
119 of significant recent disease progression,<sup>7</sup> ii) a change in contemporary  
120 measurements of 95% above the repeatability limits of the baseline  
121 measurements as shown in Supplemental Table 1 (available at  
122 <http://www.ajo.com>),<sup>7</sup> or iii) a patient considered by a clinician to be at high risk  
123 of progression despite their not fulfilling the above two criteria. Exclusion criteria  
124 included pregnancy or breastfeeding, uncontrolled ocular surface disease or a  
125 minimum corneal thickness less than 375  $\mu\text{m}$ .

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128 All the data used for model fitting started from the first appointment in the EKC.  
129 Patient demographics included age, gender, smoking status (current or ex/non-  
130 smoker) ethnicity and postcode. Ethnicity was coded as 1 for 'Black' or 'South  
131 Asian or South Asian British' and 0 for any other category (excluding missing  
132 values). Before model fitting, the pachymetry in microns was divided by 10 to  
133 generate a meaningful scale. For the primary analysis, eyes with any missing  
134 data were excluded. We also explored multiple imputation, which avoids data  
135 exclusion by generating multiple versions of the dataset with missing values  
136 replaced with values sampled from an appropriate distribution. To see whether  
137 genetic data can help predict keratoconus progression, we used 28 candidate  
138 SNPs from a recent keratoconus genome-wide association study that contained  
139 926 patients from Moorfields Eye Hospital.<sup>48</sup> The SNP data was encoded as  
140 either 0 (homozygous reference genotype), 1 (heterozygous genotype), or 2  
141 (homozygous variant genotype). We chose to use an additive encoding, thus  
142 the risk of disease increases additively with the degree of genetic variation.<sup>49</sup>  
143 Anonymized data were then exported to Excel software for analysis (version  
144 15.24 2016, Microsoft Corp.).

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## 146 **Model Fitting and Covariate Selection**

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147 A Royston-Parmar flexible parametric survival model was fitted to the data to  
148 predict the probability of an eye progressing to CXL.<sup>20,23</sup> Initial analysis of the  
149 covariates was performed by univariate analysis using the same model  
150 characteristics as the multivariable model. When selecting covariates for the  
151 final multivariable model, we used backwards stepwise selection with a  
152 significance level of 0.05. We used linear covariates for ease of interpretation of  
153 our final model. To create a more parsimonious model we examined the effect  
154 on explained variation and discrimination of removing single variables from the  
155 model.

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### 157 **Keratometric Progression Sensitivity Analysis**

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158 We included a sensitivity analysis in which we investigated keratometric  
159 progression as an alternative end-point. Keratometric progression was defined  
160 using thresholds from Gore et al 2021.<sup>7,8</sup> When using numerical thresholds to  
161 define progression, the appointments for eyes beyond the date of CXL cannot  
162 be used. However, censoring these eyes at the date of CXL represents  
163 informative censoring. Based on the recommendations of Clarke et al.<sup>15,16</sup> for  
164 investigating the impact of informative censoring, we generated a 'best case'  
165 dataset where eyes were censored at the CXL date and a 'worst case' dataset  
166 where patients were assumed to progress at the CXL date. The corresponding  
167 Kaplan Meier curves were plotted to provide a visual comparison of the two  
168 datasets. A Royston-Parmar model was then fitted on both datasets. We used  
169 the same techniques (backwards stepwise selection, significance level of 0.05)  
170 as described in the previous section to fit the model and compare the explained  
171 variation and hazard ratios.

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### 173 **Multivariable Model Validation**

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174 We validated the model using internal-external cross validation in which we split  
175 the dataset by geographical region.<sup>21,22,24,25</sup> For the kth region, the model is fitted  
176 on the full dataset excluding region k and then Kaplan-Meier curves and  
177 predicted survival curves were generated for region k. Seven geographical  
178 regions were created based on the patient's postcode as shown in  
179 Supplemental Figure 1 (available at <http://www.ajo.com>). To quantitatively  
180 assess the validation, Royston and Sauerbrei's D statistic was calculated for

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181 both the model fitted from data excluding region k ( $D_{k^c}$ ) and also the model  
182 applied to region k ( $D_k$ ).<sup>23,24</sup> The difference between these two discrimination  
183 metrics ( $D_k - D_{k^c}$ ) was calculated with its corresponding standard error to assess  
184 the predictive ability of the model. To demonstrate how the model could be  
185 used in practice, we include three hypothetical patients' eyes with different  
186 progression risk profiles (high, medium, low risk) and plot the predicted time-to-  
187 event curve for each shown in Figure 2.

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### 189 **Statistical Analysis**

190 The event of interest was defined as the date that the eye underwent CXL. We  
191 calculated the time-to-event as the difference between the first appointment in  
192 our service and the date of CXL (or the last patient appointment in the case of  
193 censoring). Since we had paired observations (eyes), we used variance-  
194 corrected models to account for correlation between eyes and to ensure that  
195 robust standard errors were produced. The choice of scale and selection of  
196 degrees of freedom for the Royston-Parmar model was informed by inspecting  
197 the Akaike information criterion (AIC) and Bayes information criterion (BIC)<sup>47,20</sup>  
198 and the results of this were balanced with ease of interpretation. See  
199 Supplemental Table 2 and Supplemental Text 1 (available ~~at~~ at  
200 <http://www.ajo.com>) for further explanation. Royston and Sauerbrei's D statistic  
201 was used as a measure of discrimination and  $R^2_{D}$  ~~as~~ as a measure of explained  
202 variation (both calculated on the natural scale of the model). Although all of the  
203 primary results were generated from a complete case analysis, we performed  
204 an additional analysis using multiple chained imputation (predictive mean  
205 matching approach with 5 nearest neighbors). Model fitting was performed in  
206 Stata 13 (StataCorp LP, Texas, USA) and the Royston-Parmar model was fitted  
207 using the stpm2 package from Stata 13.

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### 209 **Results**

#### 210 **Cohort**

211 From a potential of 9,341 eyes (4316 pairs of eyes and 709 individual eyes), the  
212 final model used 8,701 eyes of 4,823 patients, with 3,232 eyes that had CXL.  
213 The mean age was 28.3 years with standard deviation of 7.1 years. We  
214 excluded 640 eyes with missing data. Table 1 summarizes the available

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215 covariates along with missing data percentages. See Supplemental Text 2 and  
216 Supplemental Table 3 (available at <http://www.ajo.com>) for a description of the  
217 multiple imputation results.

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219 **Model Fitting and Covariate Selection (Genetic Data)**

220 We analyzed patients with genetic data separately because this data was only  
221 available for ~14% of patients. Of 926 patients (1852 eyes) with genetic data,  
222 531 eyes were excluded with incomplete keratometry or CXL data, which left  
223 1321 eyes, of which 665 had CXL. With univariate analysis of the 28 SNPs only  
224 rs72631889 was found to be significant (P=0.01) (Supplemental Table 4  
225 (available [at http://www.ajo.com](http://www.ajo.com))). We then produced a multivariable model  
226 via backwards selection on this subset of eyes using corneal data, patient data  
227 and rs72631889 as an additional covariate as shown in Supplemental Table 5  
228 (available at <http://www.ajo.com>). However rs72631889, although significant  
229 (P=0.005), had a negligible contribution (0.3%) to the explained variation in the  
230 final model.

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232 **Model Fitting and Covariate Selection (Excluding Genetic Data)**

233  
234 The results of the univariate time-to-event analysis on the hazards scale using  
235 a Royston-Parmar flexible parametric model is shown in Table 2. Genetic data  
236 was excluded from this analysis. All variables except smoking status were  
237 significant. The explained variation ( $R^2_D$ ) and discrimination (D) were highest for  
238 age (17%) and Kmax (15%) with Front K1, Front K2, Back K1, Back K2 and  
239 pachymetry each explaining 6-10% of the variation. Notably, gender and  
240 ethnicity, although significant in the univariate analysis, did not contribute to  
241 explained variation. The hazard ratios for significant covariates indicate that  
242 increasing age at presentation, greater pachymetry and flatter (less negative)  
243 posterior keratometry values decrease risk of having CXL, whilst steeper  
244 anterior keratometry values and male gender increase the risk of having CXL.

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246 When we fitted a multivariable model the significant covariates were age,  
247 Kmax, Front K1, Front K2 and pachymetry (Table 2). When we removed single  
248 variables from the model the effect this had on explained variation and

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discrimination is shown in Supplemental Table 6 (available ~~at~~at <http://www.ajo.com>). Age was the most important covariate (16.7%), with Kmax contributing ~5% of explained variation. K1, K2 and pachymetry had a small effect (<1%) when removed individually. We chose a model without K2 on the basis of parsimony, which was supported by the fact that K1 and K2 were highly correlated ( $R^2=0.91$ ) as shown in Supplemental Figure 2 (available ~~at~~at <http://www.ajo.com>). The final fitted model hazard ratios can be seen on the multivariable column of Table 2. It is notable that an increase in K1 now has a protective effect in the final model. The explained variation and discrimination for the final model were 32.7% and 1.43 respectively.<sup>24,22</sup> The opposing effect of Kmax and Front K1 can be explained by examining their regression coefficients before converting to hazard ratios; Kmax has a positive coefficient (0.0795) and Front K1 has a negative coefficient (-0.0749). This is logically similar to including the combined covariate (Kmax - Front K1) in the model which can be viewed clinically as a proxy for irregular astigmatism. We also investigated combining K1 and K2 into a single covariate as K2-K1 (standard definition of astigmatism), but the corresponding p value was not significant.

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Figure 1 visually depicts the result of applying the final model to the original dataset. As expected, the predicted mean survival curves closely follow the Kaplan-Meier curves. To demonstrate the use of the model in clinical practice, survival curves for three hypothetical patients followed for five years are shown in Figure 2. We have also produced a ~~free~~web application from the model which can be accessed at

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<https://pontikoslab.com/keprog>~~.http://beta.moorfieldscxl.com.~~

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### **Keratometric Progression Sensitivity Analysis**

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The results of the keratometric progression sensitivity analysis can be found in the Supplementary Material. By examining the Kaplan Meier curves in Supplemental Figure 3, we can see that the best case time-to-event curve indicates a 40% survival probability at 5 years whilst the worst case curve indicates a 27% survival probability at 5 years. This 13% ~~differeed~~difference in survival probability at 5 years represents the upper bound of the discrepancy in survival probability within the data. After fitting the Royston-Parmar model,

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283 amongst the hazard ratios which overlap (age, ~~kmax~~Kmax, k2), there was  
284 reasonable similarity (Supplemental Tables 8 and 9). Most importantly, the  
285 model fitted to the best case ~~data~~ had an explained variation of 11% compared  
286 to 23% for the worst case indicating a significant difference in model  
287 performance depending on the assumptions used for handling eyes which  
288 received CXL.

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### 290 **Multivariable Model Validation**

291 When performing validation using internal-external cross validation, Figure 3  
292 shows the ability of our final model to predict keratoconus progression across  
293 different geographic regions. We did not identify any significant differences in  
294 prognostic factors across regions. The model prediction curves generally follow  
295 the Kaplan Maier curves. Notably, region 5 (South West Greater London) and  
296 region 7 (other regions) have a worse predictive performance than the other  
297 regions, indicating that these regions have different characteristics compared  
298 with the remainder of the dataset used for model fitting. This could be due to  
299 differing patient characteristics, such as complex cases that required referral to  
300 our tertiary referral ~~center~~centre rather than being managed locally. Overall, the  
301 prediction becomes less accurate over time, which is expected due to low  
302 numbers with follow-up beyond three years. Supplemental Table 7 (~~available~~  
303 ~~at <http://www.ajo.com>~~) displays quantitative validation results of the model  
304 using internal external validation. The difference column  $D_k - D_{(k)}$  is a measure of  
305 predictive ability. Region 7 (other regions outside of Greater London) has the  
306 greatest discrepancy in discrimination (-0.26) which indicates that the model  
307 fitted when excluding region 7 had greater discriminative ability than when  
308 applied to region 7 alone.

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### 310 **Discussion**

312 In this study we have incorporated demographic, keratometric, and genetic data  
313 to generate a prognostic model of keratoconus progression to CXL. We have  
314 shown that parameters recorded at the first examination (age, Kmax, Front K1,  
315 minimum pachymetry) can produce a time-to-event curve to calculate a  
316 personalized risk for keratoconus progression. Although we chose time to CXL

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317 rather than keratometric progression as the end point for the time-to-event  
318 analysis, we performed a sensitivity analysis using keratometric progression,  
319 and found that a CXL model accounts for a much higher proportion of the  
320 explained variation (33%) compared to the keratometric model (11% or 23% for  
321 best and worst case respectively). The opposing effects of Kmax and Front K1  
322 were surprising/unexpected, but similar to including the combined covariate  
323 (Kmax - Front K1) in the model; a possible explanation is that the opposing  
324 effect is the result of an increase in irregular astigmatism. Of the significant  
325 covariates in our model, younger age made the greatest contribution to our  
326 model. Thus, one should have a lower threshold for treatment in younger  
327 patients.

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328 When applying internal-external cross validation, the survival curves closely  
329 followed the Kaplan Meier survival curves for each of the geographic regions,  
330 which indicates generalisability, and model discrimination between training and  
331 cross validation groups was similar, indicating that the predictive ability is well  
332 maintained. Finally, our SNP genetic data had limited additional predictive utility  
333 for keratoconus progression. However, the genetic dataset was relatively small  
334 (926 patients), and recruitment was based on the presence of keratoconus, as  
335 opposed to the severity of keratoconus, or any other index of risk of rapid  
336 progression.

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337  
338 The Royston-Parmar model has previously been used to predict the likelihood  
339 of the worst eye of patients with keratoconus progressing to corneal  
340 transplantation.<sup>25,28</sup> In their final model, Quartilho et al chose 3 significant  
341 covariates: Kmax, age and ethnicity. The reported covariate hazard ratios that  
342 overlap with our study (Kmax and age) were different in magnitude but in the  
343 same direction. When performing internal validation their model exhibited good  
344 predictive ability. They produced time-dependent receiver operating curves  
345 using the validation set and found one-year sensitivity and specificity to be  
346 92.8% and 94.6% respectively. Using logistic regression, Kato et al. found that  
347 the two strongest factors associated with the requirement for CXL were age and  
348 Kmax, which is consistent with our findings.<sup>46-49</sup> Moreover the team went on to  
349 find that age combined with corneal tomography maps was able to predict  
350 progression and need for crosslinking using deep learning.<sup>29</sup>

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351  
352 An ability to generate personalized time-to-event curves that predict  
353 progression to CXL (Figure 2) could directly inform clinical decisions that benefit  
354 patient care. Firstly, patients may better understand their own risk for  
355 progression and feel more confident in choosing their management options.  
356 Secondly, for both clinicians and patients, the prediction of progression may  
357 contribute to scheduling treatments, including prioritizing patients at high risk of  
358 early progression. For example, patients at high risk with a 98% probability of  
359 progressing to CXL at 5 years could be offered CXL at the point of first  
360 diagnosis without waiting to demonstrate keratometric progression. Medium risk  
361 patients may benefit from a period of clinician-led topographic monitoring. For  
362 the lowest risk patients, optometry-led monitoring in the community may be  
363 sufficient. This risk stratification could be tailored to regions and reflect local  
364 needs and resources such as provision of monitoring services in regions with  
365 lower risk and greater capacity for CXL in areas with more high risk patients.  
366 Finally, when a decision is made to postpone CXL for further monitoring, the  
367 time-to-event curve can contribute to decisions on the scheduling of future  
368 follow up reviews, with perhaps shorter time periods where the curve is  
369 steepest. Recommendations based on this model on clinical practice is yet to  
370 be evaluated.

371  
372 Our study is subject to several limitations inherent to our dataset. First, if  
373 patients had CXL at another hospital, this may not be reliably recorded in the  
374 source database. This could lead to a very small number of patients being  
375 included in the analysis who have already had CXL. Second, ethnicity is a well  
376 ~~known~~established risk factor for keratoconus and keratoconus  
377 progression,<sup>24,26,27,29,30,31</sup> but ethnicity is now an optional field at patient registration  
378 at our institution and this information was unavailable for approximately 50% of  
379 our dataset. However, even when we restricted the dataset to those with  
380 ethnicity records, it was not found to be a significant covariate. Third, though  
381 the cohort used for univariable and multivariable analysis were identical the  
382 number of eyes where all covariates were available was lower than for  
383 univariable analysis due to missing data. Finally, when we used multiple  
384 imputation to generate a multivariable model, ethnicity was still not found to be

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385 significant. In the model fitting process we chose to use a simple backwards  
386 selection as opposed to the multivariate fractional polynomial (MFP) method.<sup>28,32</sup>  
387 In our initial investigations, the results of MFP yielded nonlinear functional forms  
388 of the covariates and, whilst this method may have slightly increased the  
389 predictive power of the prognostic model, the resulting hazard ratios would be  
390 very hard to interpret. In addition, we did not examine time dependent effects  
391 for the covariates, which may provide a more accurate model fit, and future  
392 studies should examine this option. Finally, although no external validation  
393 dataset was available, internal external cross validation allowed us to confirm  
394 that our model is generalizable across geographical regions.

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396 In conclusion, we have fitted a prognostic model for progression of keratoconus  
397 to CXL which generates a time-to-event curve using age, Kmax, Front K1,  
398 minimum pachymetry from time of presentation. Incorporation of a relatively  
399 small genetic dataset does not improve the explained variation of our model.  
400 Personalized modeling of risk may improve patients' understanding of their  
401 condition and the need for CXL. Such a model may help better improve patients  
402 and aid clinician decision making to CXL to achieve better outcomes and  
403 judicious use of healthcare resources.

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405 **Disclosures**

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406 The authors have no financial disclosures.

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408 Funding/Support: HM is funded by a Moorfields Eye Charity PhD Studentship  
409 (GR001147). NP is funded by a Moorfields Eye Charity Career Development  
410 Award (R190031A). Moorfields Eye Charity is supported in part by the National  
411 Institute for Health Research (NIHR) Biomedical Research Centre based at  
412 Moorfields Eye Hospital NHS Foundation Trust and UCL Institute of  
413 Ophthalmology. ST, BA and DG acknowledge that a proportion of their financial  
414 support is from the Department of Health through the award made by the  
415 National Institute for Health Research to Moorfields Eye Hospital NHS  
416 Foundation Trust and University College London Institute of Ophthalmology for  
417 a Specialist Biomedical Research Centre for Ophthalmology. The sponsor or  
418 funding organization had no role in the design or conduct of this research.

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Other acknowledgements: none.

**References**

1. [Gordon MO, Steger-May K, Szczotka-Flynn L, et al. Baseline factors predictive of incident penetrating keratoplasty in keratoconus. \*Am J Ophthalmol.\* 2006;142\(6\):923-930.](#)
2. [Wittig-Silva C, Chan E, Islam FMA, Wu T, Whiting M, Snibson GR. A randomized, controlled trial of corneal collagen cross-linking in progressive keratoconus: three-year results. \*Ophthalmology.\* 2014;121\(4\):812-821.](#)
3. [Caporossi A, Mazzotta C, Baiocchi S, Caporossi T. Long-term results of riboflavin ultraviolet A corneal collagen cross-linking for keratoconus in Italy: the Siena eye cross study. \*Am J Ophthalmol.\* 2010;149\(4\):585-593.](#)
4. [O'Brart DPS, Chan E, Samaras K, Patel P, Shah SP. A randomised, prospective study to investigate the efficacy of riboflavin/ultraviolet A \(370 nm\) corneal collagen cross-linkage to halt the progression of keratoconus. \*Br J Ophthalmol.\* 2011;95\(11\):1519-1524.](#)
5. [Koller T, Mrochen M, Seiler T. Complication and failure rates after corneal crosslinking. \*J Cataract Refract Surg.\* 2009;35\(8\):1358-1362.](#)
6. [Gore DM, Leucci MT, Koay SY, et al. Accelerated Pulsed High-Fluence Corneal Cross-Linking for Progressive Keratoconus. \*Am J Ophthalmol.\* 2021;221:9-16.](#)
7. [Salmon HA, Chalk D, Stein K, Frost NA. Cost effectiveness of collagen crosslinking for progressive keratoconus in the UK NHS. \*Eye.\* 2015;29\(11\):1504-1511.](#)

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447 8. [Lindstrom RL, Berdahl JP, Donnenfeld ED, et al. Corneal cross-linking](#)  
448 [versus conventional management for keratoconus: a lifetime economic](#)  
449 [model. \*J Med Econ\*. Published online November 19, 2020:1.](#)

450 9. [Godefrooij DA, Mangen MJJ, Chan E, et al. Cost-Effectiveness Analysis of](#)  
451 [Corneal Collagen Crosslinking for Progressive Keratoconus.](#)  
452 [Ophthalmology. 2017;124\(10\):1485-1495.](#)

453 10. [Larkin DFP, Chowdhury K, Burr JM, et al. Effect of Corneal Cross-linking](#)  
454 [versus Standard Care on Keratoconus Progression in Young Patients: The](#)  
455 [KERALINK Randomized Controlled Trial. \*Ophthalmology\*. Published online](#)  
456 [April 20, 2021. doi:10.1016/j.ophtha.2021.04.019](#)  
457 [Gordon MO, Steger-May K, Szezoika-Flynn L, et al.](#)  
458 [Baseline factors predictive of incident penetrating keratoplasty in keratoconus.](#)  
*Am J Ophthalmol*. 2006;142(6):923-930.

459 Wittig-Silva C, Chan E, Islam FMA, Wu T, Whiting M, Snibson GR. A  
460 randomized, controlled trial of corneal collagen cross-linking in progressive  
461 keratoconus: three-year results. *Ophthalmology*. 2014;121(4):812-821.

462 Caporossi A, Mazzotta C, Baiocchi S, Caporossi T. Long-term results of  
463 riboflavin-ultraviolet A corneal collagen cross-linking for keratoconus in Italy: the  
464 Siena eye cross study. *Am J Ophthalmol*. 2010;149(4):585-593.

465 O'Brart DPS, Chan E, Samaras K, Patel P, Shah SP. A randomised,  
466 prospective study to investigate the efficacy of riboflavin/ultraviolet A (370 nm)  
467 corneal collagen cross-linkage to halt the progression of keratoconus. *Br J*  
468 *Ophthalmol*. 2011;95(11):1519-1524.

469 5. [Gore DM, Shortt AJ, Allan BD. New clinical pathways for keratoconus. \*Eye\*.](#)  
470 [2013;27\(3\):329-339.](#)

471 6. [Koller T, Mrochen M, Seiler T. Complication and failure rates after corneal](#)  
472 [crosslinking. \*J Cataract Refract Surg\*. 2009;35\(8\):1358-1362.](#)

473 7. [Gore DM, Leucci MT, Koay S-Y, et al. Accelerated Pulsed High-Fluence](#)  
474 [Corneal Cross-Linking for Progressive Keratoconus. \*Am J Ophthalmol\*.](#)  
475 [2021;221:9-16.](#)

476 8. [Larkin DFP, Chowdhury K, Burr JM, et al. Effect of Corneal Cross-linking](#)  
477 [versus Standard Care on Keratoconus Progression in Young Patients: The](#)  
478 [KERALINK Randomized Controlled Trial. \*Ophthalmology\*. Published online](#)  
479 [April 20, 2021. doi:10.1016/j.ophtha.2021.04.019](#)

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480 11. Vinciguerra R, Belin MW, Borgia A, et al. Evaluating keratoconus  
481 progression prior to crosslinking: maximum keratometry vs the ABCD  
482 grading system. *J Cataract Refract Surg.* 2021;47(1):33-39.

483 12. Shajari M, Steinwender G, Herrmann K, et al. Evaluation of keratoconus  
484 progression. *Br J Ophthalmol.* 2019;103(4):551-557.

485 13. Ozalp O, Atalay E. Belin ABCD Progression Display Identifies Keratoconus  
486 Progression Earlier Than Conventional Metrics. *Am J Ophthalmol.*  
487 2022;236:45-52.

488 14. Hashemi H, Panahi P, Asgari S, Emamian MH, Mehravaran S, Fotouhi A.  
489 Best Indicators for Detecting Keratoconus Progression in Children: A Report  
490 From the Shahroud Schoolchildren Eye Cohort Study. *Cornea.*  
491 2022;41(4):450-455.

492 15. Flynn TH, Sharma DP, Bunce C, Wilkins MR. Differential precision of  
493 corneal Pentacam HR measurements in early and advanced keratoconus.  
494 *Br J Ophthalmol.* 2016;100(9):1183-1187.

495 16. Flockerzi E, Häfner L, Xanthopoulou K, et al. Reliability analysis of  
496 successive Corneal Visualization Scheimpflug Technology measurements in  
497 different keratoconus stages. *Acta Ophthalmol.* 2022;100(1):e83-e90.

498 17. Kreps EO, Jimenez-Garcia M, Issarti I, Claerhout I, Koppen C, Rozema JJ.  
499 Repeatability of the Pentacam HR in Various Grades of Keratoconus. *Am J*  
500 *Ophthalmol.* 2020;219:154-162.

501 18. Clark TG, Bradburn MJ, Love SB, Altman DG. Survival analysis part IV:  
502 further concepts and methods in survival analysis. *Br J Cancer.*  
503 2003;89(5):781-786.

504 19. Kato N, Negishi K, Sakai C, Tsubota K. Baseline factors predicting the need  
505 for corneal crosslinking in patients with keratoconus. *PLoS One.*  
506 2020;15(4):e0231439.

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507 20. Patrick Royston PL. *Flexible Parametric Survival Analysis Using Stata: Beyond the Cox Model*. Stata Press; 2011.

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509 21. Hardcastle AJ, Liskova P, Bykhovskaya Y, et al. A multi-ethnic genome-wide association study implicates collagen matrix integrity and cell differentiation pathways in keratoconus. *Commun Biol*. 2021;4(1):266.

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512 22. Ding X, Guo X. A Survey of SNP Data Analysis. *Big Data Mining and Analytics*. 2018;1(3):173-190.

514 23. Royston P, Parmar MKB. Flexible parametric proportional-hazards and proportional-odds models for censored survival data, with application to prognostic modelling and estimation of treatment effects. *Stat Med*. 2002;21(15):2175-2197.

518 24. Baade PD, Royston P, Youl PH, Weinstock MA, Geller A, Aitken JF. Prognostic survival model for people diagnosed with invasive cutaneous melanoma. *BMC Cancer*. 2015;15:27.

521 25. Royston P, Parmar MKB, Sylvester R. Construction and validation of a prognostic model across several studies, with an application in superficial bladder cancer: CONSTRUCTION AND VALIDATION OF PROGNOSTIC MODEL. *Stat Med*. 2004;23(6):907-926.

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525 26. Royston P, Sauerbrei W. A new measure of prognostic separation in survival data. *Stat Med*. 2004;23(5):723-748.

527 27. Tuft SJ, Moodaley LC, Gregory WM, Davison CR, Buckley RJ. Prognostic factors for the progression of keratoconus. *Ophthalmology*. 1994;101(3):439-447.

530 28. Quartilho A, Gore DM, Bunce C, Tuft SJ. Royston-Parmar flexible parametric survival model to predict the probability of keratoconus progression to corneal transplantation. *Eye*. 2020;34(4):657-662.

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533 29. Kato N, Masumoto H, Tanabe M, et al. Predicting Keratoconus Progression  
534 and Need for Corneal Crosslinking Using Deep Learning. *J Clin Med Res.*  
535 2021;10(4). doi:10.3390/jcm10040844

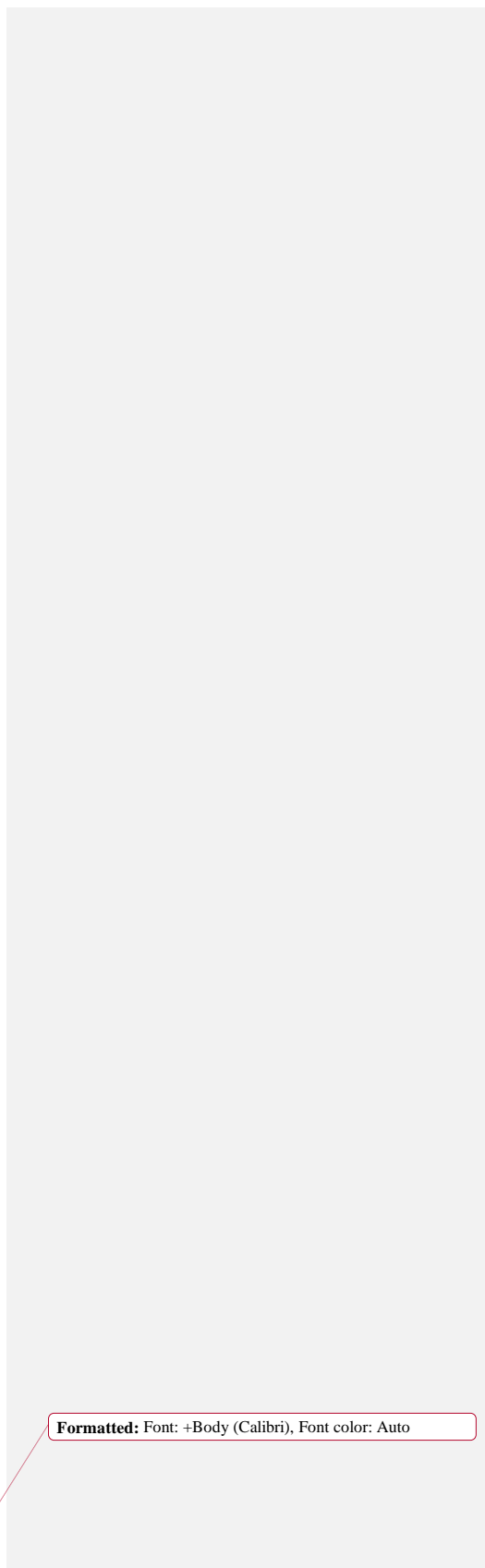
536 30. Pearson AR, Soneji B, Sarvananthan N. Does ethnic origin. *Eye .*  
537 2000;14:625-628.

538 31. Georgiou T, Funnell CL, Cassels-Brown A, O'Conor R. Influence of ethnic  
539 origin on the incidence of keratoconus and associated atopic disease in  
540 Asians and white patients. *Eye .* 2004;18(4):379-383.

541 32. Royston P. *Multivariable Model-Building: A Pragmatic Approach to*  
542 *Regression Analysis Based on Fractional Polynomials for Continuous*  
543 *Variables.* John Wiley; 2008.

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547 ~~9.~~ Salmon HA, Chalk D, Stein K, Frost NA. Cost effectiveness of collagen  
548 crosslinking for progressive keratoconus in the UK NHS. *Eye*.  
549 2015;29(11):1504-1511.

550 ~~10.~~ Lindstrom RL, Berdahl JP, Donnenfeld ED, et al. Corneal cross-linking versus  
551 conventional management for keratoconus: a lifetime economic model. *J Med*  
552 *Econ*. Published online November 19, 2020:1.

553 ~~11.~~ Godofrooij DA, Mangen M-JJ, Chan E, et al. Cost-Effectiveness Analysis of  
554 Corneal Collagen Crosslinking for Progressive Keratoconus.  
555 *Ophthalmology*. 2017;124(10):1485-1495.

556 ~~12.~~ Vinciguerra R, Belin MW, Borgia A, et al. Evaluating keratoconus  
557 progression prior to crosslinking: maximum keratometry vs the ABCD  
558 grading system. *J Cataract Refract Surg*. 2021;47(1):33-39.

559 ~~13.~~ Shajari M, Steinwender G, Herrmann K, et al. Evaluation of keratoconus  
560 progression. *Br J Ophthalmol*. 2019;103(4):551-557.

561 ~~14.~~ Flynn TH, Sharma DP, Bunce C, Wilkins MR. Differential precision of  
562 corneal Pentacam HR measurements in early and advanced keratoconus.  
563 *Br J Ophthalmol*. 2016;100(9):1183-1187.

564 ~~15.~~ Clark TG, Bradburn MJ, Love SB, Altman DG. Survival analysis part IV:  
565 further concepts and methods in survival analysis. *Br J Cancer*.  
566 2003;89(5):781-786.

567 ~~16.~~ Kato N, Negishi K, Sakai C, Tsubota K. Baseline factors predicting the need  
568 for corneal crosslinking in patients with keratoconus. *PLoS One*.  
569 2020;15(4):e0231439.

570 ~~17.~~ Patrick Royston PL. *Flexible Parametric Survival Analysis Using Stata:*  
571 *Beyond the Cox Model*. Stata Press; 2011.

572 ~~18.~~ Hardecastle AJ, Liskova P, Bykhovskaya Y, et al. A multi-ethnic genome-wide  
573 association study implicates collagen matrix integrity and cell differentiation  
574 pathways in keratoconus. *Commun Biol*. 2021;4(1):266.

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575 ~~10.~~ Ding X, Guo X. A Survey of SNP Data Analysis. *Big Data Mining and Analytics*. 2018;1(3):173-190.

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577 ~~20.~~ Royston P, Parmar MKB. Flexible parametric proportional hazards and proportional odds models for censored survival data, with application to prognostic modelling and estimation of treatment effects. *Stat Med*. 2002;21(15):2175-2197.

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581 ~~21.~~ Baade PD, Royston P, Youl PH, Weinstock MA, Geller A, Aitken JF. Prognostic survival model for people diagnosed with invasive cutaneous melanoma. *BMC Cancer*. 2015;15:27.

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584 ~~22.~~ Royston P, Parmar MKB, Sylvester R. Construction and validation of a prognostic model across several studies, with an application in superficial bladder cancer: CONSTRUCTION AND VALIDATION OF PROGNOSTIC MODEL. *Stat Med*. 2004;23(6):907-926.

588 ~~23.~~ Royston P, Sauerbrei W. A new measure of prognostic separation in survival data. *Stat Med*. 2004;23(5):723-748.

590 ~~24.~~ Tuft SJ, Moodaley LC, Gregory WM, Davison CR, Buckley RJ. Prognostic factors for the progression of keratoconus. *Ophthalmology*. 1994;101(3):439-447.

593 ~~25.~~ Quartilho A, Gore DM, Bunce C, Tuft SJ. Royston-Parmar flexible parametric survival model to predict the probability of keratoconus progression to corneal transplantation. *Eye*. 2020;34(4):657-662.

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596 ~~26.~~ Pearson AR, Soneji B, Sarvananthan N. Does ethnic origin. *Eye*. 2000;14:625-628.

598 ~~27.~~ Georgiou T, Funnell CL, Cassels-Brown A, O'Connor R. Influence of ethnic origin on the incidence of keratoconus and associated atopic disease in Asians and white patients. *Eye*. 2004;18(4):379-383.

60 ~~28.~~ Royston P. *Multivariable Model Building: A Pragmatic Approach to Regression Analysis Based on Fractional Polynomials for Continuous Variables*. John Wiley; 2008.

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609 **Figure 1:** Chart showing how the Royston-Parmar model fits the entire dataset.  
610 We split the eyes into 4 risk groups by their prognostic index: <25th centile (low  
611 risk), 25-50th centile (medium-low risk), 50-75th centile (medium-high  
612 risk), >75th centile (high risk). The number of eyes at risk corresponds to the  
613 Kaplan-Meier curves.

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615 **Figure 2:** Time-to-event curves that predict the risk of progression to CXL for  
616 three hypothetical patient profiles. The blue line represents a high risk patient  
617 who has a 95% probability of progressing to CXL at 5 years. The red line is a  
618 medium risk patient who has a 48% probability of progressing to CXL at 5  
619 years. The green line is a low risk patient who has a 14% probability of  
620 progressing to CXL at 5 years. The equation used to generate the curves is:  
621  $S(t) = e^{-H(t)}$ , where  $H(t)$  is the cumulative hazard function and is commonly  
622 expressed as  $\ln(H(t)) = s(\ln(t)) + x\beta$ , where  $s(\ln(t))$  is a restricted cubic  
623 spline function of log time,  $\beta$  is the vector of coefficients and  $x$  is the vector of  
624 covariates. For further details of the derivation, we refer the reader to

~~20. Abbreviations: pachy, pachymetry.~~

Abbreviations: pachy, pachymetry.

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628 **Figure 3:** Predicted and observed survival curves for seven postal code regions  
629 of Greater London as shown in Supplemental Figure 1 (available at  
630 <http://www.ajo.com>) using IECV. We split the eyes into 4 risk groups by their  
631 prognostic index: <25th centile (low risk), 25-50th centile (medium-low risk), 50-  
632 75th centile (medium-high risk), >75th centile (high risk).

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633 Abbreviations:

635 AIC: Akaike Information Criterion

636 Back K1: Flat posterior keratometry in the central 3 mm zone

637 Back K2: Steep posterior keratometry in the central 3 mm zone

638 BIC: Bayes Information Criterion

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639 CXL: Corneal ~~Collagen~~ Cross-~~Linking~~Linking ▲

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640 EKC: Early Keratoconus Clinic ▲

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641 EPR: Electronic Patient Record ▲

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642 Front K1: Flat anterior keratometry in the central 3 mm zone ▲

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643 Front K2: Steep anterior keratometry in the central 3 mm zone ▲

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644 HR: Hazard Ratio ▲

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645 IECV: Internal-external Cross Validation ▲

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646 Kmax: Maximum anterior keratometry ▲

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647 MFP: Multivariate Fractional Polynomial ▲

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648 SNP: Single-nucleotide Polymorphism ▲

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**Table 1:** Summary statistics for the available covariates at the first examination for 9341 eyes recorded at first visit.

Covariate	Type	Mean	SD	N	Missing No. (%)
Front K1 (D)	Numeric	45.31	3.86	8,813	528 (5.7)
Front K2 (D)	Numeric	48.39	4.85	8,839	502 (5.4)
Back K1 (D)	Numeric	-6.53	0.75	7,949	1392 (14.9)
Back K2 (D)	Numeric	-7.23	0.93	8,702	639 (6.8)
Kmax (D)	Numeric	54.14	8.01	8,834	507 (5.4)
Pachymetry (um)	Numeric	462.92	46.15	8,946	395 (4.2)
Age (years)	Numeric	28.28	7.10	9341	0 (0)
Genetic data <sup>a</sup>	Ordinal	N/A	N/A	1141	8020 (85.9)
Self-reported black or asian ethnicity <sup>b</sup>	Categorical (59.9% black or asian)	N/A	N/A	4889	4452 (47.7)
Male gender	Categorical (67% male)	N/A	N/A	9341	0 (0)
Smoker <sup>c</sup>	Categorical (4.5% smoker)	N/A	N/A	9341	0 (0)

Abbreviations: Front K1, flattest anterior keratometry; Front K2, steepest anterior keratometry; Back K1, flattest posterior keratometry; Back K2, steepest posterior keratometry; Kmax: maximum Keratometry; pachymetry: minimum corneal thickness; SD, Standard deviation; N, number of eyes; N/A, not applicable.

<sup>a</sup>Genetic data comprised of 28 SNPs and was encoded in an additive fashion (0,1,2).

<sup>b</sup>1=black or asian, 0=otherwise. <sup>c</sup>0=non-smoker/ex-smoker, 1=current smoker.

**Table 2:** Univariable and final multivariable model for all considered covariables excluding genetic data in the training dataset fitted on the hazards scale with 5 degrees of freedom.

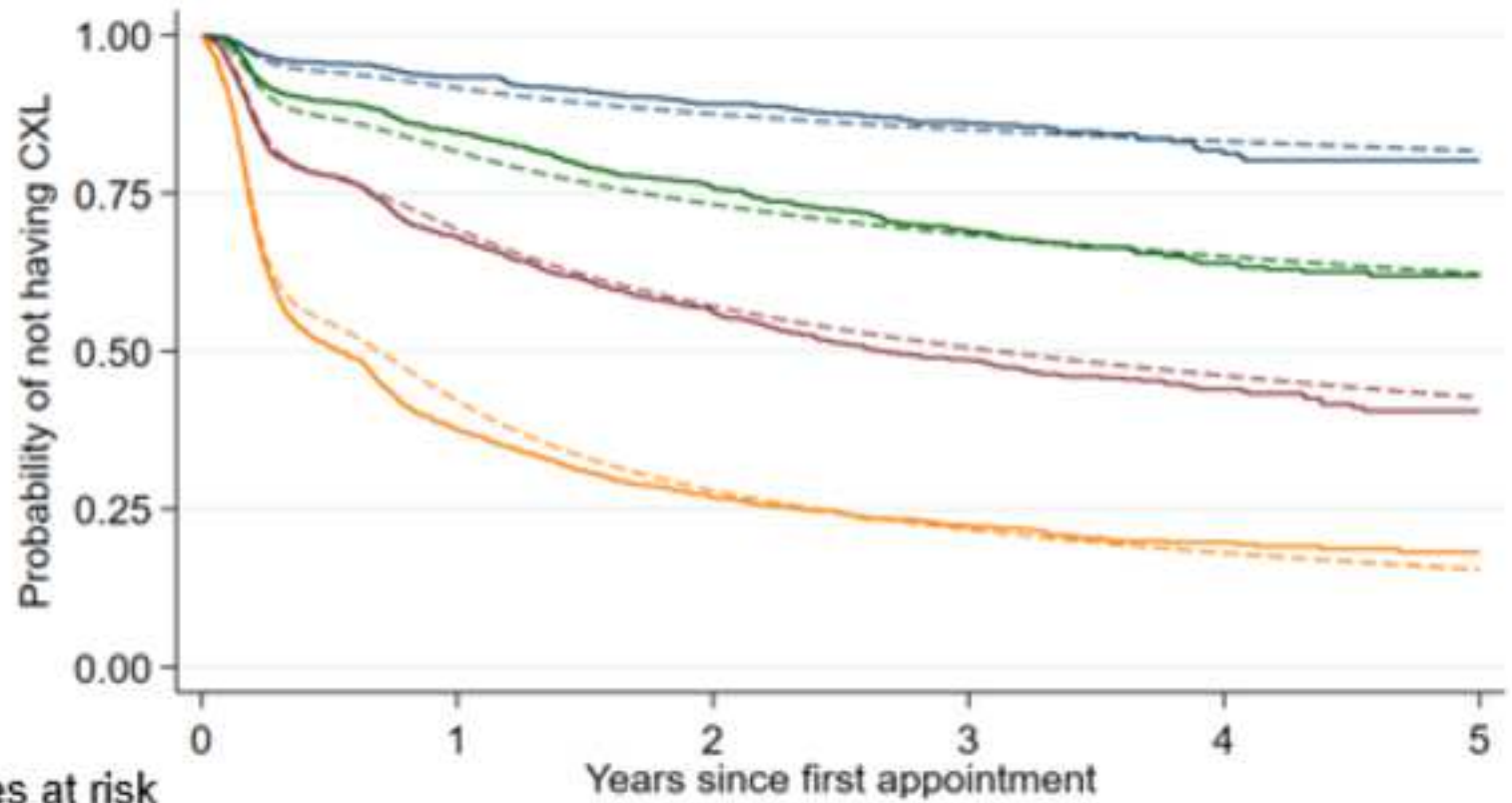
Covariate	Univariable (N=9341)				Multivariable (N=8701)	
	Hazard Ratio [95% CI]	P Value	R <sup>2</sup> <sub>D</sub>	D	Hazard Ratio [95% CI]	P Value
Ethnicity	1.14 [1.02; 1.27]	0.02	0.4%	0.13	N/A	N/A
Smoker <sup>a</sup>	1.07 [0.9; 1.28]	0.46	0.1%	0.05	N/A	N/A
Male Gender	1.11 [1.01; 1.21]	0.02	0.2%	0.10	N/A	N/A
Age at presentation	0.91 [0.9; 0.92]	<0.001	16.7%	0.92	0.9 [0.90; 0.91]	<0.001
Kmax	1.06 [1.05; 1.06]	<0.001	14.9%	0.86	1.08 [1.07; 1.09]	<0.001
Front K1	1.09 [1.08; 1.1]	<0.001	7.0%	0.56	0.93 [0.91; 0.94]	<0.001
Front K2	1.08 [1.07; 1.08]	<0.001	9.8%	0.67	N/A	N/A
Back K1 <sup>c</sup>	0.67 [0.64; 0.71]	<0.001	5.9%	0.51	N/A	N/A
Back K2 <sup>c</sup>	0.7 [0.67; 0.72]	<0.001	8.4%	0.62	N/A	N/A
Pachymetry 10 <sup>b</sup>	0.93 [0.92; 0.94]	<0.001	7.5%	0.58	0.95 [0.93; 0.96]	<0.001

Abbreviations: N, number of eyes; R<sup>2</sup><sub>D</sub>, explained variation; D, Royston and Sauerbrei's D statistic (used as a measure of discrimination); CI, confidence interval; Kmax, maximum keratometry; Front K1, flattest anterior keratometry; Front K2, steepest anterior keratometry; Back K1, flattest posterior keratometry; Back K2, steepest posterior keratometry; pachymetry, minimum corneal thickness; N/A, not applicable due to this variable not being included in the final model.

<sup>a</sup>0=non-smoker/ex-smoker, 1=current smoker.

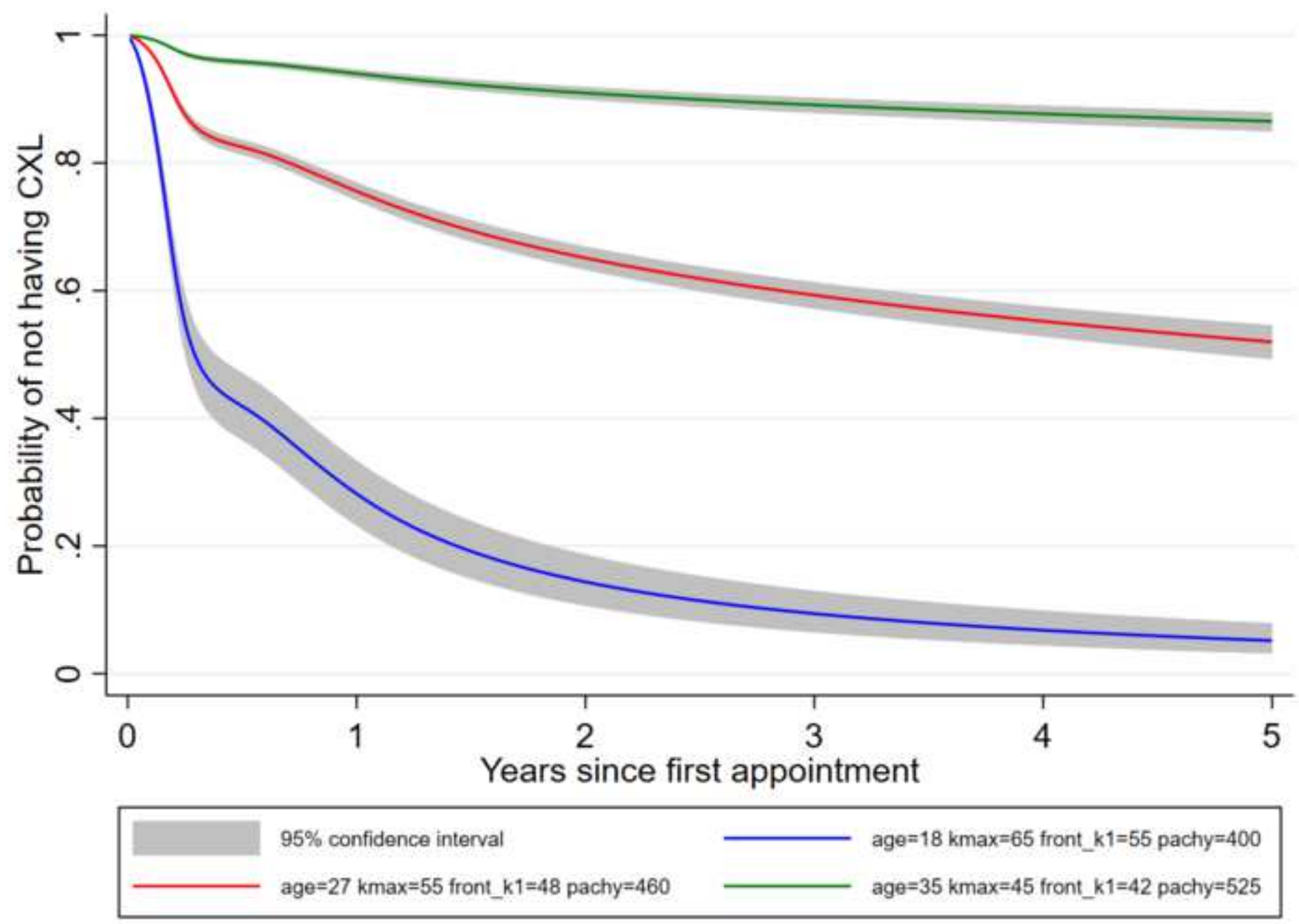
<sup>b</sup>Minimum pachymetry in steps of 10µm.

<sup>c</sup>Back K1 and Back K2 are negative values such that patients with advanced keratoconus are typically associated with large negative values. A hazard ratio below 1 indicates that as measurements become more positive, the risk of progression decreases.

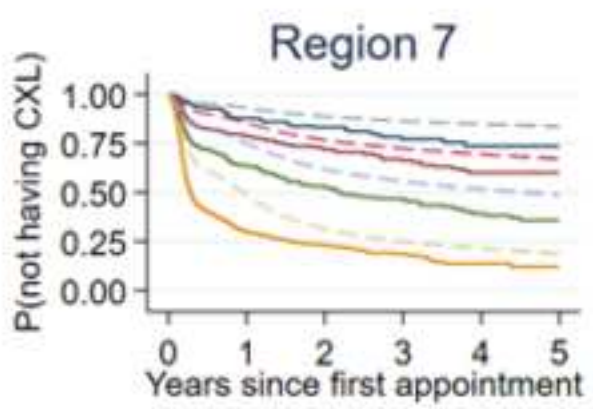
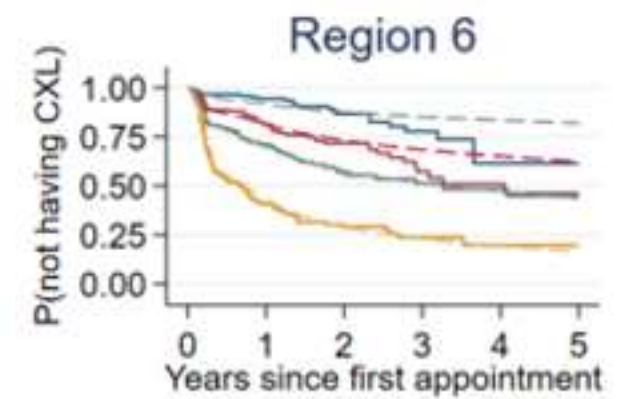
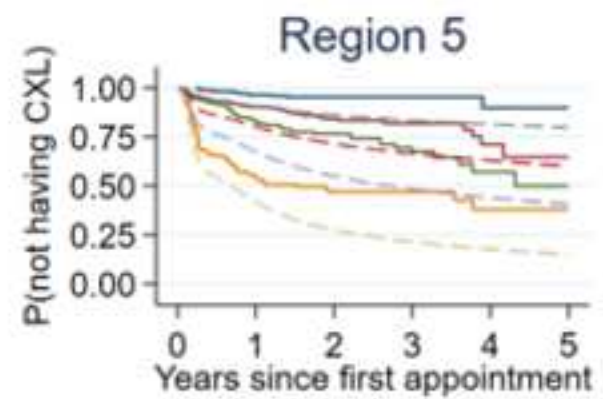
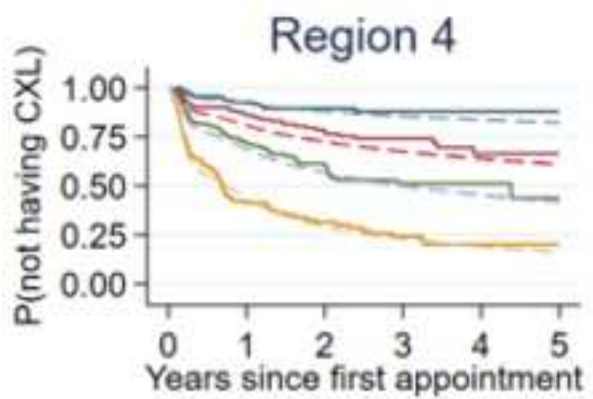
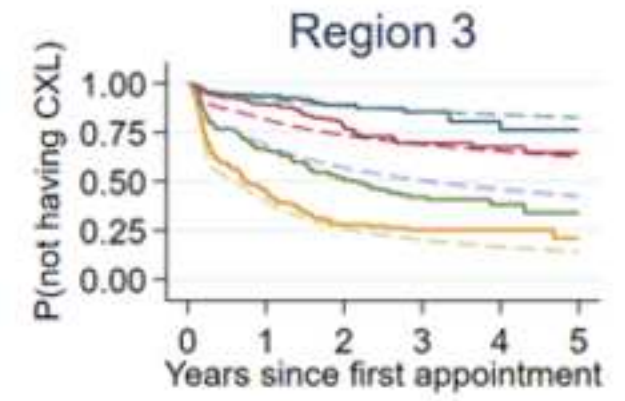
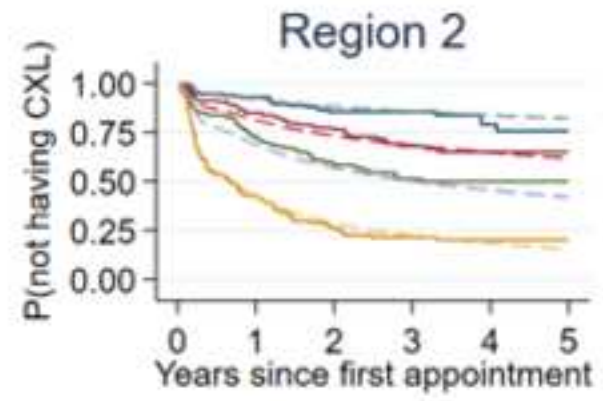
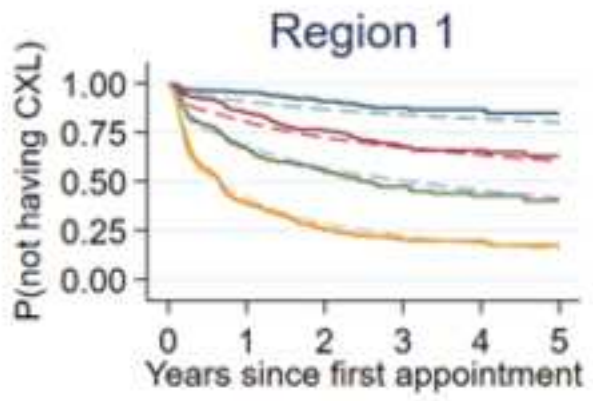


# of eyes at risk	0	1	2	3	4	5
low risk	2175	1429	824	380	163	34
medium-low risk	2175	1399	816	454	218	62
medium-high risk	2176	1134	614	326	143	37
high risk	2175	675	324	168	74	26

	Predicted	Kaplan Meier
low risk	-----	—————
medium-low risk	-----	—————
medium-high risk	-----	—————
high risk	-----	—————

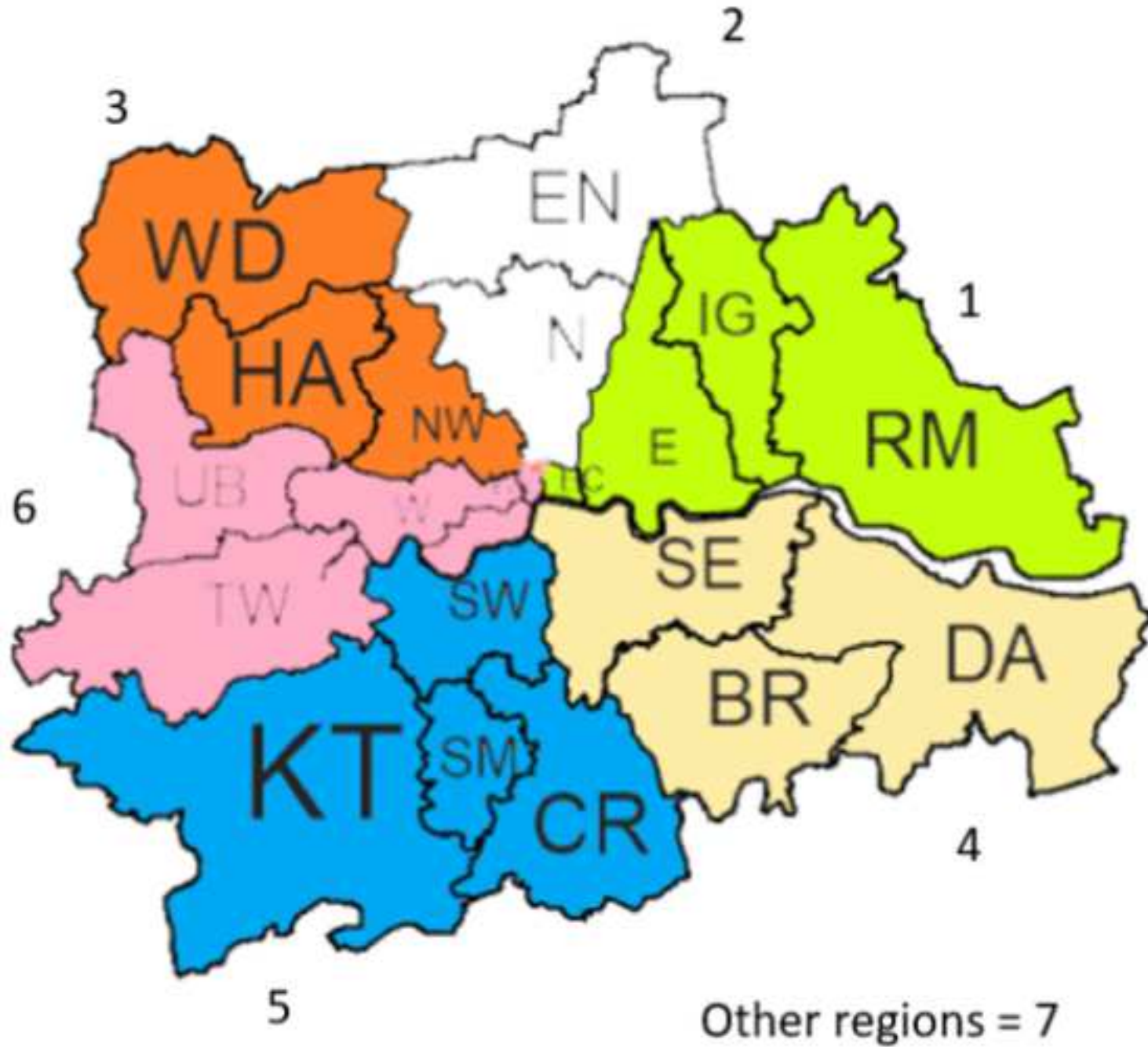




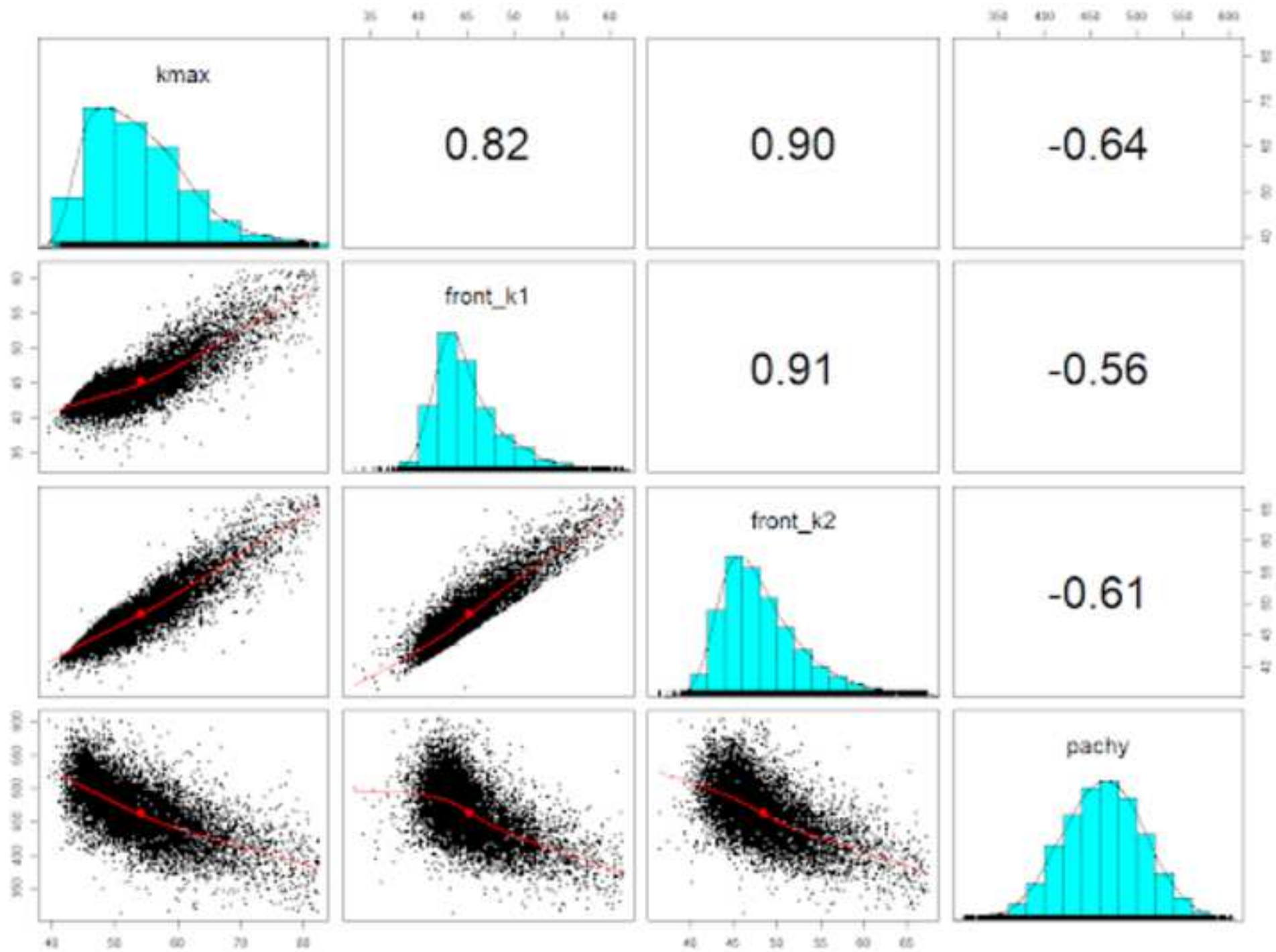




Corneal crosslinking is successful in halting keratoconus progression but providing patients with a personalized visual representation of risk to progression is desirable. This research presents a model to generate projected likelihood of having crosslinking from data collected at presentation. It was trained and validated from a large dataset of 8701 eyes from keratoconus patients. Univariable and multivariable analysis was performed to identify risk factors including single nucleotide polymorphisms associated with keratoconus.

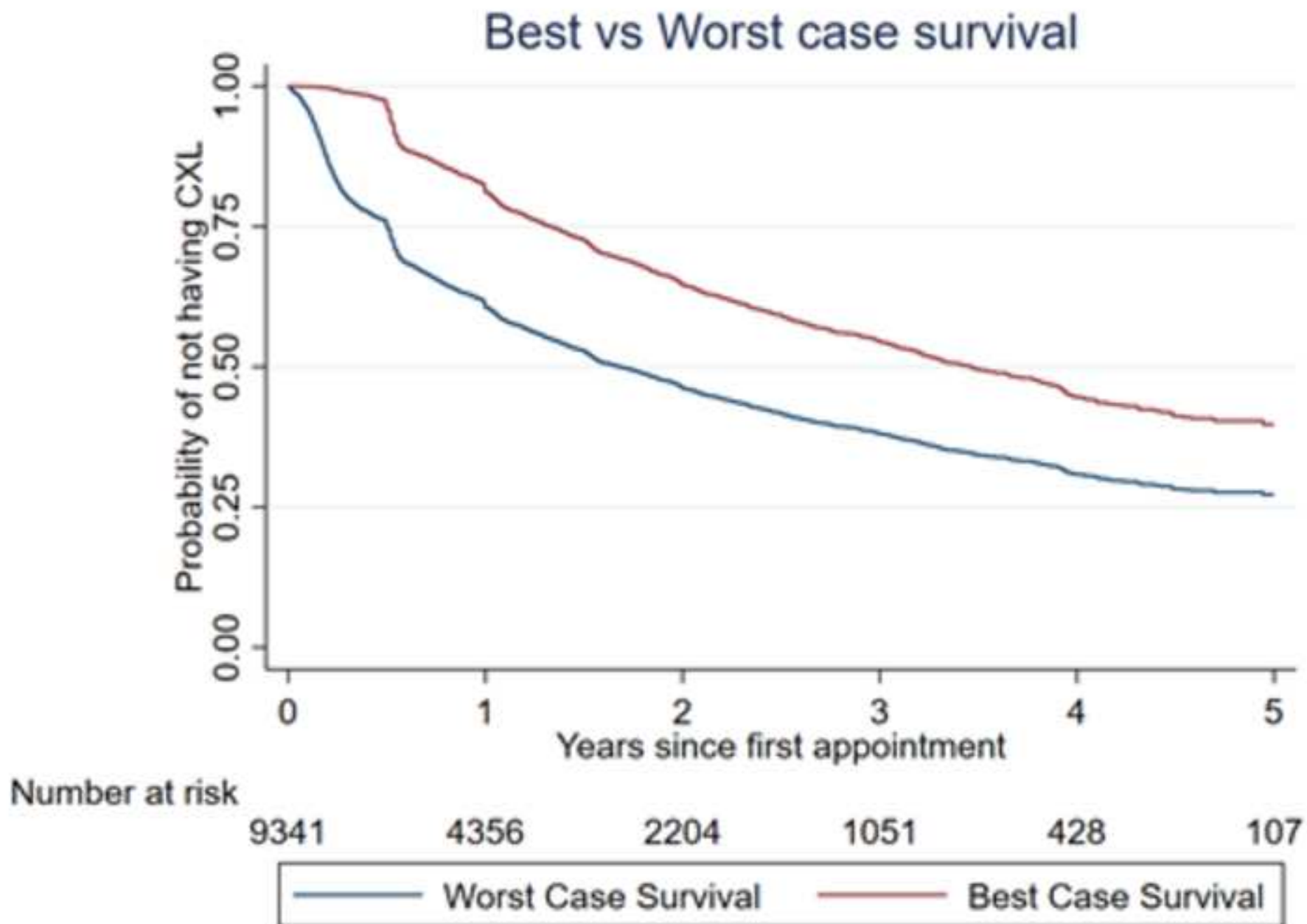


**Supplementary Figure Legend 1.** Geographic region grouping use in IECV. We split greater London into 6 postcode regions around its centre. Any postcodes outside of greater London were classified as the 7th region ('other'). The number of participants per region is given in Table G.



**Supplementary Figure 2:** Correlations amongst anterior keratometry covariates and pachymetry. As explained in the results section, Front K1 and Front K2 are very highly correlated. This correlation is exemplified by the fact that removing one or other of these covariates has negligible effect on the explained variation of the model.

Abbreviations: kmax, maximum Keratometry; front\_k1, flattest anterior keratometry; front\_k2, steepest anterior keratometry; pachy, thinnest point pachymetry



**Supplementary Figure 3:** Kaplan Meier time-to-event curves for keratometric progression in the best (censored at CXL date) and worst (progressed at CXL date) case scenarios.

**Supplementary Table 1.** Defining disease progression in early and moderate/advanced keratoconus prior to corneal cross-linking.

Early (Kmax < 55D) (1 or more)	Moderate/advanced (Kmax ≥ 55D) (1 or more)
<ul style="list-style-type: none"> <li>· ≥ 1 D increase Kmax</li> <li>· ≥ 1 D increase front K1 or K2</li> <li>· ≥ 0.5 D increase back K2</li> <li>· ≥ 16 μm decrease minimum corneal thickness</li> </ul>	<ul style="list-style-type: none"> <li>· ≥ 2.5 D increase Kmax</li> <li>· ≥ 2.5 D increase front K1 or K2</li> <li>· ≥ 22 μm decrease minimum corneal thickness</li> </ul>

Abbreviation: Kmax, maximum keratometry; K1, flat keratometry in the central 3 mm zone; K2, steep keratometry in central 3 mm zone;



**Supplementary Table 2:** Choice of scale and degrees of freedom (DF) using categorized (n=5) continuous covariates. Here it can be seen that 5 DF and the proportional odds model has the optimal (lowest) values for both AIC and BIC. However, we chose to model on the proportional hazards scale because our aim was to build a model that clinicians are familiar with.

DF	Hazard		Odds		Normal	
	AIC	BIC	AIC	BIC	AIC	BIC
1	953.51	1113.8	724.22	884.51	683.78	844.06
2	483.05	649.06	416.68	582.69	511.2	677.21
3	330.09	501.82	307.59	479.32	512.5	684.24
4	305.77	483.23	234.19	411.65	283.45	460.91
5	67.06	250.25	25.8	208.98	88.06	271.24
6	108.82	297.72	62.16	251.07	119.89	308.8

Abbreviations: DF, Degrees of freedom; AIC, Akaike Information Criterion; BIC, Bayesian Information Criterion.

**Supplementary Table 3:** Multiple chained imputation results.

<b>Covariate</b>	<b>Hazard Ratio [CI 95%]</b>	<b>P Value</b>
Age at presentation	0.91 [0.90; 0.92]	<0.001
Kmax	1.10 [1.09; 1.11]	<0.001
Pachymetry 10 <sup>a</sup>	0.97 [0.96; 0.98]	<0.001
Front K1	0.95 [0.93; 0.98]	<0.001
Front K2	0.94 [0.91; 0.96]	<0.001

Abbreviations: CI, confidence interval; Kmax, maximum Keratometry; Front K1, flattest anterior keratometry; Front K2, steepest anterior keratometry

<sup>a</sup>Minimum pachymetry in steps of 10 $\mu$ m.

**Supplementary Table 4:** Univariable results for genetic data. This includes the 28 most significantly associated Single-nucleotide Polymorphisms (SNPs)<sup>1</sup>.

SNP name	Hazard Ratio [95% CI]	P Value	R <sup>2</sup> <sub>b</sub>	D
rs59771807	1.13 [0.72; 1.76]	0.60	0.00	0.08
rs3115850	1.11 [0.96; 1.27]	0.16	0.00	0.12
rs114525117	1.1 [0.85; 1.43]	0.46	0.00	0.08
rs950122	0.97 [0.85; 1.12]	0.71	0.00	0.03
rs115991721	0.72 [0.45; 1.14]	0.16	0.01	0.24
rs55678698	0.93 [0.65; 1.35]	0.72	0.00	0.06
rs6657440	1.03 [0.91; 1.17]	0.63	0.00	0.04
rs72631889	0.4 [0.22; 0.72]	0.002	0.09	0.66
rs12184325	1.09 [0.83; 1.42]	0.53	0.00	0.07
rs74460547	1.2 [0.6; 2.4]	0.60	0.00	0.12
rs72631887	1.29 [0.8; 2.07]	0.29	0.01	0.18
rs4040617	1.11 [0.96; 1.28]	0.17	0.00	0.11
rs116390263	0.96 [0.62; 1.49]	0.85	0.00	0.03
rs116452738	0.66 [0.24; 1.86]	0.43	0.01	0.17
rs3131972	1.06 [0.93; 1.21]	0.35	0.00	0.08
rs3131962	1.08 [0.94; 1.24]	0.29	0.00	0.09
rs13303222	1.02 [0.86; 1.2]	0.85	0.00	0.01

rs4970382	1.04 [0.93; 1.18]	0.48	0.00	0.06
rs116587930	1.26 [0.91; 1.75]	0.17	0.01	0.18
rs11240779	1.11 [0.98; 1.26]	0.10	0.00	0.14
rs4970459	0.98 [0.84; 1.14]	0.79	0.00	0.03
rs12562034	1 [0.84; 1.2]	0.98	0.00	0.00
rs116720794	1.09 [0.84; 1.42]	0.52	0.00	0.07
rs13303101	1.1 [0.94; 1.29]	0.24	0.00	0.11
rs4970383	1.06 [0.93; 1.2]	0.37	0.00	0.08
rs79373928	1.45 [0.82; 2.58]	0.21	0.01	0.24
rs192998324	1.15 [0.72; 1.83]	0.57	0.00	0.10
rs57181708	1.05 [0.88; 1.26]	0.55	0.00	0.04

Abbreviations: N, number of eyes; R<sup>2</sup>, explained variation; D, Royston and Sauerbrei's D statistic (used as a measure of discrimination); CI, confidence interval.

**Supplementary Table 5.** Hazard ratios for genetic sub-analysis model using 1144 eyes containing clinical data and rs72631889 SNP.

Covariate	Hazard Ratio [95% CI]	P Value
Age at presentation	0.92 [0.90; 0.93]	<0.001
Kmax	1.10 [1.07; 1.12]	<0.001
Back K2	1.60 [1.36; 1.87]	<0.001
Pachymetry10 <sup>a</sup>	0.96 [0.94; 0.99]	0.010
rs72631889	0.42 [0.23; 0.77]	0.005

Abbreviations: CI, confidence interval; Back K2, steepest posterior keratometry; Kmax, maximum Keratometry; Back K2, steepest posterior keratometry

<sup>a</sup>Minimum pachymetry in steps of 10 $\mu$ m.

**Supplementary Table 6.** The effect on explained variation and discrimination of removing individual covariates from the full model. Removing age has the most dramatic effect, reducing  $R^2_b$  from 33.3% to 19.7%. Removing Kmax reduced  $R^2_b$  from 33.3% to 28.3%. The other covariates are less important with pachymetry, Front K1 and Front K2 reducing  $R^2_b$  by <1% when removed.

Covariate Removed	$R^2_b$	D
None	0.333	1.45
Age at presentation	0.197	1.01
Kmax	0.284	1.29
Front K1	0.324	1.42
Front K2	0.327	1.43
Pachymetry10 <sup>a</sup>	0.324	1.42

Abbreviations:  $R^2_b$ , explained variation; D, Royston and Sauerbrei's D statistic (used as a measure of discrimination); Kmax, maximum Keratometry; Front K1, flattest anterior keratometry; Front K2, steepest anterior keratometry  
<sup>a</sup>Minimum pachymetry in steps of 10 $\mu$ m.

**Supplementary Table 7.** Internal-external cross-validation across different geographic regions.

Region	N	CXL events	$D_k$	$D_{(k)}$	$D_k - D_{(k)}$	D S.E.
1	2,088	727	1.58	1.38	0.19	0.09
2	946	330	1.31	1.44	-0.12	0.15
3	1,014	366	1.35	1.44	-0.09	0.11
4	711	222	1.59	1.41	0.17	0.16
5	1,050	203	1.60	1.40	0.21	0.14
6	574	216	1.28	1.44	-0.16	0.14
7	2,318	1,168	1.22	1.48	-0.26	0.08

Abbreviations: N, number in region k;  $D_{(k)}$ , discrimination omitting region k;  $D_k$ , discrimination predicted in region k;  $D_k - D_{(k)}$ , difference in discrimination between; S.E., Standard error for  $D_k - D_{(k)}$ .

**Supplementary Table 8.** Best Case Hazard Ratios

Covariate	Hazard Ratio	P Value	95% confidence interval	
age	0.95	<0.001	0.95	0.96
kmax	1.08	<0.001	1.06	1.10
k2	0.91	<0.001	0.88	0.93



**Supplementary Table 9.** Worst Case Hazard Ratios

Covariate	Hazard Ratio	P Value	95% confidence interval	
age	0.93	<0.001	0.92	0.94
kmax	1.09	<0.001	1.08	1.10
k1	0.98	0.02	0.96	1.00
k2	0.95	<0.002	0.93	0.97
pachymetry	0.97	<0.003	0.96	0.98

### **Supplementary Text 1. Choice of Scale and Degrees of Freedom**

Royston-Parmar models can be fit on a number of different scales (Hazard, Odds, Normal) and the degrees of freedom for the baseline spline can also take a range of integer values. Scale selection and degrees of freedom for the Royston-Parmar model was informed by inspecting the AIC, BIC as suggested by Royston et al. <sup>1</sup>. When selecting these two aspects of the model, it is important not to assume linear covariate effects for continuous variables so we categorized each continuous variable into 5 categories first. We subsequently iterated over the 3 different scales (Hazard, Odds, Probit) and 6 degrees of freedom to find the optimal (minimum) AIC and BIC. The results of iterating over both scale and degrees of freedom in order to guide further analysis can be seen in Supplementary Table 2.

1. Patrick Royston PL. Flexible Parametric Survival Analysis Using Stata: Beyond the Cox Model. Stata Press; 2011.

## **Supplementary Text 2. Multiple Imputation results**

When applying multiple chained imputation, all 9,341 eyes were included in the analysis and we imputed the missing values for Front K1, Front K2, Back K1, Back K2, Kmax, pachymetry and ethnicity. We did not attempt to impute missing genetic data because the percentage missing (86%) was deemed to be too high. When we imputed missing data by repeating the model fitting process using multiple chained imputation the hazard ratios were very similar to the complete case analysis (Supplementary Table 3).