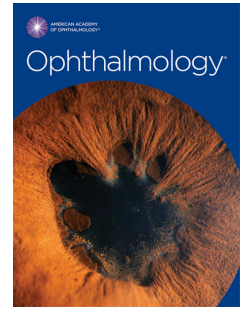


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RP2-associated X-linked Retinopathy: Clinical Findings, Molecular Genetics, and Natural History.

Michalis Georgiou, Anthony G. Robson, Katarina Jovanovic, Thales Antônio Cabral De Guimarães, MD, Naser Ali, Nikolas Pontikos, Sami H. Uwaydat, Omar A. Mahroo, Michael E. Cheetham, Andrew R. Webster, Alison J. Hardcastle, Michel Michaelides

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1 **RP2-associated X-linked Retinopathy: Clinical Findings, Molecular**
2 **Genetics, and Natural History.**

3

4 Michalis Georgiou^{1,2,3}, Anthony G. Robson^{1,2}, Katarina Jovanovic², Thales
5 Antônio Cabral De Guimarães, MD,^{1, 2} Naser Ali,^{1, 2} Nikolas Pontikos,^{1, 2} Sami
6 H. Uwaydat³, Omar A Mahroo^{1,2}, Michael E. Cheetham², Andrew R. Webster^{1,2},
7 Alison J. Hardcastle², Michel Michaelides^{1,2} *

8

9 ¹. Moorfields Eye Hospital, London, UK

10 ². UCL Institute of Ophthalmology, University College London, London, UK

11 ³. Jones Eye Institute, University of Arkansas for Medical Sciences, Little Rock,
12 AR, USA.

13

14 ***Correspondence / Reprints:**

15 Prof. Michel Michaelides,

16 UCL Institute of Ophthalmology, 11-43 Bath Street, London, EC1V 9EL, UK.

17 Email: michel.michaelides@ucl.ac.uk

18 Tel: +44 207 608 6864

19

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35 ABSTRACT

36

37 Purpose:

38 To review and describe in detail the clinical course, functional and anatomical
39 characteristics of *RP2*-associated retinal degeneration.

40

41 **Design:** Retrospective case series.

42

43 Participants

44 Males with disease-causing variants in the *RP2* gene.

45

46 Methods:

47 Review of all case notes and results of molecular genetic testing, retinal
48 imaging (fundus autofluorescence (FAF) imaging, optical coherence
49 tomography (OCT)) and electrophysiology assessment.

50

51 Main Outcome Measures

52 Molecular genetic testing, clinical findings including best-corrected visual acuity
53 (BCVA), qualitative and quantitative retinal imaging analysis, and
54 electrophysiology parameters.

55

56 Results

57 Fifty-four molecularly confirmed patients were identified, from 38 pedigrees.
58 Twenty-eight disease-causing variants were identified; with 20 not previously
59 clinically characterized. Fifty-three patients (98.1%) presented with retinitis
60 pigmentosa. The mean age of onset (range, \pm SD) was 9.6 years of age (1-57
61 years, \pm 9.2 years). Forty-four patients (91.7%) had childhood-onset disease,
62 with mean age of onset of 7.6 years. The commonest first symptom was night
63 blindness (68.8%). Mean BCVA (range, \pm SD) was 0.91 LogMAR (0-2.7, \pm 0.80)
64 and 0.94 LogMAR (0-2.7, \pm 0.78) for right and left eyes respectively. Based on
65 the WHO visual impairment criteria, 18 patients (34%) had low vision. The
66 majority (17/22) showed ERG evidence of a rod-cone dystrophy. Pattern ERG
67 P50 was undetectable in all but 2 patients. A range of FAF findings was
68 observed, from normal to advanced atrophy. There were no statistically

69 significant differences between right and left eyes for ellipsoid zone (EZ) width
70 and outer nuclear layer (ONL) thickness. The mean annual rate of EZ width
71 loss was 219 $\mu\text{m}/\text{year}$ and the mean annual decrease in ONL thickness was
72 4.93 $\mu\text{m}/\text{year}$. No patient with childhood-onset disease had identifiable EZ after
73 the age of 26 years at baseline or follow-up. Four patients had adulthood-onset
74 disease and a less severe phenotype.

75

76 **Conclusions:** This study details the clinical phenotype of *RP2* retinopathy in a
77 large cohort. The majority presented with early-onset severe retinal
78 degeneration, with early macular involvement and complete loss of the foveal
79 photoreceptor layer by the third decade of life. Full-field ERGs revealed rod-
80 cone dystrophy in the vast majority, but with generalised (peripheral) cone
81 system involvement of widely varying severity in the first two decades of life.

82

83 **Abbreviations/Acronyms:** XLRP, X-linked Retinitis Pigmentosa; SD-OCT,
84 Spectral Domain optical coherence tomography; FAF, Fundus
85 autofluorescence; RPE, Retinal pigment epithelium; CFP, Color fundus
86 photography; ISCEV, International Society for Clinical Electrophysiology of
87 Vision; ERG, Electroretinogram; PERG, Pattern ERG; VF, Visual Field; SD,
88 standard deviation; VA, visual acuity.

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90 **INTRODUCTION**

91

92 Retinitis Pigmentosa (RP) is an heterogeneous group of inherited retinal
93 conditions, both in terms of phenotype and genotype, with a prevalence of
94 1/3000 - 1/4000 in the general population.¹ RP can be inherited in an autosomal
95 dominant, autosomal recessive or X-linked pattern.^{1, 2} X-Linked RP (XLRP)
96 cases account for 15% of males with simplex disease.³ XLRP is a severe form
97 of RP, with most affected males presenting with early-onset vision loss (<10
98 years of age), nyctalopia, nystagmus, severely abnormal or undetectable
99 electroretinogram (ERG) and progression to legal blindness by the 3rd to 4th
100 decade.⁴⁻⁶ XLRP patients have symptomatic night blindness from early
101 childhood and are often myopic. *RPGR* and *RP2* disease-causing variants are
102 the commonest causes of XLRP accounting for 80-90% of cases.¹ The on-
103 going gene therapy clinical trials for *RPGR*-associated XLRP⁷ were preceded
104 by multiple studies describing in depth characterization of disease natural
105 history.⁸⁻¹⁴ In contrast, the current literature describing the *RP2* phenotype is
106 limited.

107 *RP2* disease-causing variants are responsible for 5-20% of XLRP.¹⁵⁻²⁰
108 The reports comparing the severity of *RPGR* and *RP2* XLRP have been
109 inconclusive as to which genotype is associated with worse prognosis.^{2, 5, 6, 21,}
110 ²² The genotype-phenotype correlations in *RP2*-associated XLRP are limited.²³
111 Differential diagnosis of *RP2*- or *RPGR*-XLRP is challenging, since no ocular
112 measurement is genotype-specific.^{4, 5} A tapetal-like reflex can be observed both
113 in patients and carriers with *RPGR*- and *RP2*-XLRP.²⁴

114 *RP2* (MIM 312600) is located on Xp11.23 and has a structure similar to
115 cofactor C, which is involved in β -tubulin folding.¹⁹ *RP2* encodes a GTPase-
116 activating protein (GAP) for the small GTPase ARL3, and has a role in
117 trafficking lipidated proteins in the retina to the outer segment of
118 photoreceptors.^{25, 26} Using retinal pigment epithelium (RPE) and 3D retinal
119 organoids differentiated from patient-derived inducible pluripotent stem cells
120 (iPSCs) with an *RP2* premature stop variant, read-through drugs and AAV gene
121 therapy rescued the cellular phenotype, supporting a clinical trial in patients.^{27,}
122 ²⁸ However, there is currently a lack of robust natural history data in genetically
123 proven patients with *RP2*-associated retinopathy. These data are needed to
124 provide better informed advice on prognosis and optimize design of clinical
125 trials including identifying possible robust outcome measures and participant
126 stratification.

127 The current study thereby provides a detailed characterization of the
128 clinical phenotype, molecular basis, and natural history of a large series of
129 patients with *RP2* retinopathy.

130

131 **METHODS**

132

133 *Subject identification and assessment*

134 Males harboring disease-causing variants in *RP2* were identified from
135 Moorfields Eye Hospital (London, UK) and University of Arkansas Medical
136 Science (Little Rock, AR, USA) retinal genetics clinics. All patients included
137 were previously informed and consented. This retrospective study adhered to
138 the tenets of the Declaration of Helsinki and was approved by the local ethics
139 committees.

140

141 *Molecular diagnosis*

142 The majority of patients were screened using a diagnostic targeted next
143 generation sequencing panel for retinal dystrophy. Others were ascertained
144 either via research based whole exome sequencing, or with targeted Sanger
145 sequencing of *RP2*. Variants are annotated according to the Reference
146 Sequence NM_006915. All variants have a gnomAD frequency of 0 (gnomAD
147 v2.1.1). Splice site variants were assessed using SpliceAI
148 (<https://spliceailookup.broadinstitute.org>).

149

150 *Clinical notes*

151 Clinical data extracted included age of onset, visual acuity, slit lamp
152 biomicroscopy and fundoscopy findings. Symptoms at presentation and
153 complications were also recorded. All available data were reviewed, including
154 the findings at the last available follow-up.

155

156 *Best Corrected Visual Acuity (BCVA) and Clinical severity grading*

157 BCVA was assessed monocularly with Snellen chart and converted to
158 logarithmic minimum angle of resolution (LogMAR) for statistical analysis.
159 Jayasundera et al. have described an approach to subdivide *RP2-XLRP*
160 patients into mild, less severe, and severe.²⁰ Patients with relatively late onset
161 severe macular dysfunction were considered less severe. BCVA with different
162 cut-offs for different age ranges, was used as a subjective surrogate for macular
163 function. We adopted and adapted the same clinical severity grading criteria
164 into LogMAR, and applied it for the best seeing eye (**Supplementary Table 1**).

165 In addition, BCVA of the best seeing eye was used to categorize patients
166 into one of four groups based on the World Health Organization (WHO) visual
167 impairment criteria, that defines a person with no or mild visual impairment
168 when presenting VA is ≤ 0.48 LogMAR, moderate impairment when VA is 0.48
169 -1 LogMAR, severe if 1-1.3 LogMAR, and blindness if it is greater than 1.3
170 LogMAR (**Supplementary Table 1**). Low vision corresponds to patients with
171 moderate and severe impairment. Counting fingers vision was given a value of
172 LogMAR 1.98 and hand motion, LogMAR 2.28, light perception and no light
173 perception were specified as LogMAR 2.7 and 3, respectively.²⁹ The BCVA
174 classification criteria are summarized in **Supplementary Table 1**.

175

176 *Electrophysiological testing*

177 Pattern and full-field electroretinogram (PERG; ERG) testing was performed in
178 22 patients, incorporating the standards of the International Society for Clinical
179 Electrophysiology of Vision (ISCEV).^{30, 31} Pattern ERG P50 was used as an

180 objective measure of macular function and the full-field ERG used to assess
181 generalised retinal function of rod and cone systems. The ERG data were
182 compared with a reference range from a group of healthy subjects (age range:
183 10-79 years).^{32, 33} The amplitudes of the main full-field ERG components were
184 plotted as a percentage of the age-matched lower limit of normal or as a
185 difference from the age-matched upper peak time limit, including the dark
186 adapted (DA) 10 ERG a-wave, and the light-adapted (LA) 3 single flash ERG
187 b-wave and the LA 3 30Hz ERG. To address non-Gaussian distribution within
188 the control group, the limits were defined as the lowest amplitude value in the
189 control group minus 5% of the reference range (maximum minus minimum
190 values) for amplitudes or plus 5% of the reference range for peak times.^{34, 35}

191

192 *Fundus Autofluorescence (FAF)*

193 FAF images were obtained using short-wavelength excitation (488 nm) and a
194 scanning laser ophthalmoscope according to previously described methods.³⁶
195 Images were reviewed by one grader (MG) and qualitatively graded.

196

197 *Optical Coherence Tomography (OCT)*

198 The majority of patients seen over the last fifteen years had both OCT and FAF
199 imaging. Horizontal scans acquired using the Heidelberg Spectralis OCT
200 (Heidelberg Engineering, Heidelberg, Germany) were chosen for quantifying
201 the residual ellipsoid zone width (EZW), using the foveal reflex as a reference
202 point. In addition, the device was switched to follow-up mode, so that the same
203 scanning location was imaged at the follow-up visit as the baseline. This
204 enabled comparable measurements to be made between the two visits for a

205 given subject. In others, the analysis described by Tee et al.³⁷ was used to align
206 locations for follow-up measurements of retinal thickness and the EZW
207 (described in detail in **Supplementary Methods**). Vendor supplied Heidelberg
208 Eye Explorer (Heyex) software version 1.6.1.0 was used for image analysis and
209 quantification of EZW, employing the caliper tool.³⁸

210

211 *Statistical Analysis*

212 Statistical analysis was carried out using SPSS Statistics for Windows (Version
213 22.0. Armonk, NY: IBM Corp.). Significance for all statistical tests was set at
214 $P < 0.05$. The Shapiro-Wilk test was used to test for normality for all variables.

215

216 **RESULTS**

217

218 *Molecular Genetics*

219 A total of 54 molecularly confirmed patients were identified, from 38 pedigrees.
220 Twenty-eight variants were identified. The most common variant was c.358C>T
221 p.(Arg120*), identified in 5 pedigrees (13%); 6 variants were identified in 2
222 pedigrees each and all others were restricted to single families. Identified
223 variants included 8 frameshift alterations (28.6%), 7 missense (25.0%), 6
224 nonsense (21.4%) and 3 splice site changes (10.7%). One patient had a whole
225 gene deletion and three patients had smaller deletions. Twenty of the variants,
226 including the deletions, have not been previously clinically characterized.
227 **Figure 1** shows the distribution of the variants across the *RP2* gene/protein.

228 **Supplementary Table 2** details the identified variants including their predicted
229 effect.

230

231 *Phenotype, Age of Onset and Presenting Symptoms*

232 Fifty-three patients (98.1%) presented with RP. Age of onset was documented
233 for 48 patients. The mean age of onset (range, \pm SD) was 9.6 years of age (1-
234 57 years, \pm 9.2 years). Forty-four patients (91.7%) had childhood-onset disease
235 (age range 1-16 years old), with mean age of disease onset (\pm SD) of 7.6 years
236 (\pm 4.1 years). The four patients with adulthood-onset disease had mean age of
237 onset 32.5 years (range: 17-57 years). One patient presented with symptoms
238 and signs consistent with cone-rod dystrophy (CORD, 1.9%) with onset of
239 symptoms at age 10 years.

240 The first symptom/s at disease onset were described in 48 of the patients
241 with RP and included night blindness (n=33, 68.8%), decreased central vision
242 (n=8, 16.7%), both night blindness and decreased central vision (n=4, 8.3%),
243 decreased central vision and peripheral vision loss (n=2, 4.2%). One patient
244 with RP presented with nystagmus (n=1, 2.1%). The patient with CORD
245 presented with decreased central vision and developed night vision difficulties
246 later in life. Clinical data are summarized in **Table 3**.

247

248 *Genotype-phenotype correlations*

249 Null and missense variants were present in childhood-onset and adult-hood
250 onset groups. Of the four patients with adulthood-onset disease, two had
251 frameshift variants with truncation/loss of function, one had a splice site variant
252 with loss of donor splice site and one had a substitution. In the childhood-onset

253 group the phenotype was uniform, with early onset disease and early
254 degeneration. No genotype-phenotype correlations were observed in the
255 current report.

256

257 *Best Corrected Visual Acuity (BCVA)*

258 BCVA was documented in at least one visit for 53 patients and was reduced in
259 all cases. Mean age (range, \pm SD) for baseline BCVA for the whole cohort was
260 23.2 years (3.8-71, \pm 17.4 years), with a mean BCVA (range, \pm SD) of 0.91
261 LogMAR (0-2.7, \pm 0.80) and 0.94 LogMAR (0-2.7, \pm 0.78) for right and left eyes
262 respectively. Forty-three had available longitudinal data, with a mean follow-up
263 (range, \pm SD) of 7.3 years (0.3-30.2, \pm 7.1 years). Mean BCVA change was 0.37
264 and 0.29 Log MAR for right and left eyes, respectively for the follow-up period,
265 and was not statistically significantly different between right and left eyes
266 (paired t-test $P < 0.05$). BCVA data are summarized in **Table 1** and mean
267 baseline BCVA against age is presented in **Figure 2A**.

268

269 *Disease Severity*

270 Based on previously described clinical severity grading criteria
271 (**Supplementary Table 1**): 21 patients had mild disease and 32 patients had
272 severe disease at baseline. Of the 21 patients with mild disease, 18 were seen
273 longitudinally. Eight of those 18 patients met the criteria for severe disease over
274 a follow-up of 9.9 years (SD: \pm 4.8, range: 3-15.1 years). The 10 patients with
275 mild disease at the last follow-up visit, had significantly shorter follow-up time
276 (mean \pm SD: 4.8 ± 4.1 years), with three of them having later onset adulthood
277 disease.

278 Based on the WHO visual impairment criteria: 24 patients (45%) had
279 no or mild visual impairment, 11 patients (21%) had moderate impairment, 7
280 (13%) had severe impairment and 11 (21%) were blind. In total, 18 patients
281 (34%) had low vision. **Figure 2B** depicts the age distribution for each class of
282 visual impairment.

283 *Non-Ocular Manifestations*

284 No non-ocular manifestations were identified. However, ascertainment bias
285 cannot be excluded, as the vast majority of patients were recruited from a
286 stand-alone eye hospital (MEH).

287

288 *Electrophysiology*

289 There was high degree of inter-ocular ERG symmetry based on amplitudes of
290 the DA 0.01, DA 3 and DA 10 ERG a- and b-waves, LA 30Hz ERG and LA 3
291 (single flash) ERG b-waves (slope = 0.94; $r^2= 0.95$) and on the peak times of
292 the DA 10 ERG b-waves and LA 30Hz ERGs (slope = 1.1; $r^2=0.86$).

293 Three of 22 patients had undetectable full-field ERGs under all stimulus
294 conditions (ages 8, 18 and 21 years) and 2 others showed severe and similar
295 reductions of DA and LA ERGs, consistent with a severe rod and cone
296 photoreceptor dystrophy. The majority (n=17), including the 11 with the mildest
297 DA10 ERG a-wave reductions, showed better preservation of LA ERGs than
298 DA 10 ERG a-waves, in keeping with a rod-cone dystrophy (**Figure 3**). All 17
299 patients with a detectable response showed delay in the LA 30Hz ERG,
300 including the majority (n=13) with severe delays of between 10 and 24ms.
301 Pattern ERG P50 was undetectable in all but 2 patients, including patient 12
302 (P50 delayed by 7ms and reduced by >70%; **Figure 4b**) and patient 21 (P50
303 delayed by 10ms and reduced by >25%; **Figure 4c**). **Figure 3** summarizes the
304 electrophysiological findings and patient ages at the time of testing and **Figure**
305 **4** shows representative recordings.

306 There was no significant correlation between age and the amplitudes of
307 the DA 0.01 ERG, DA 10 ERG a- and b-waves, LA 30Hz ERG or LA3 ERG b-

308 waves (maximum $r^2 = 0.083$) or the peak times of the LA 30Hz ($r^2=0.025$),
309 although the narrow age range is highlighted (all but 1 patient were aged
310 between 5 and 21 years). Serial data were available in one child from the age
311 of 7 years and revealed progressive PERG P50 reduction over 5 years and
312 marked worsening of the DA 10 ERG between the ages of 10 and 12 years
313 (**Supplementary Figure 5**).

314

315 *Fundus Autofluorescence (FAF)*

316 FAF imaging was available for 46 patients for at least one visit. At first
317 evaluation, the mean age (\pm SD, range) was 25.1 years (\pm 16.7, 5.8-69.2 years).
318 A range of FAF findings was observed, from normal FAF to advanced atrophy.
319 **Figure 6** shows examples of the different patterns of FAF observed. On
320 qualitative assessment we identified normal FAF in 11 patients (23.9%, mean
321 age \pm SD, range: 15.2 \pm 11.1, 5.8-46.2 years old). Two patients, aged 11 and 24
322 years, had a paracentral macular ring of increased signal; six patients (13%)
323 had a macular ring of increased signal and mid-peripheral patchy changes with
324 a mean age (\pm SD, range) of 18.71 (5.8, 11-25.9) years old; four patients had
325 patchy macular signal and midperipheral changes (8.7%, mean age \pm SD,
326 range: 26.0 \pm 28.1, 6.7-68.2 years old); a further four patients had normal
327 macular signal with patchy midperipheral changes (8.7%, mean age \pm SD,
328 range: 27.4 \pm 23.4, 12.2-62.1 years old), and one patient had patchy macular
329 signal with normal periphery. Eighteen patients (39.1%) had atrophy at a mean
330 age 34.0 years old (\pm 15.1, 15.8-69.2). Three of the patients with advanced
331 atrophy had a choroideremia-like pattern. Atrophy was the commonest pattern
332 on FAF imaging and in total 31 patients (67.4%) had visible changes at the
333 macula at baseline.

334 Follow-up FAF was available for thirty-six patients. The mean (\pm SD,
335 range) follow-up time was 6.0 years (\pm 4.3, 0.6-17.6 years). Nine of nine patients
336 with normal FAF at baseline showed abnormal changes (mean follow-up period
337 of 6.4 years). The FAF showed a high degree of inter-ocular symmetry in all
338 cases, including those that had repeat imaging (examples shown in
339 **Supplementary Figure 7**).

340

341 *Optical Coherence Tomography (OCT)*

342 Forty-six patients had at least one OCT imaging session. Baseline age (\pm SD,
343 range) was 27.2 years old (\pm 17.5, 5.2-69.2 years). EZ width and ONL thickness
344 was not statistically significantly different between right and left eye (paired t-
345 test $P < 0.05$). For further assessment, the mean EZW and ONL thickness for
346 both eyes was calculated for each patient at each visit.

347 Forty-two of the 46 patients had childhood-onset disease. Twenty-three
348 had no identifiable EZ and complete ONL thickness loss at mean age (\pm SD,
349 range) of 36.4 years (\pm 16.0, 17.9-69.2). Nineteen patients (mean age \pm SD,
350 range: 13.5 \pm 5.7, 5.3-25.9 years), had identifiable EZ and residual ONL
351 thickness. Mean EZW was 1493 (\pm 1496, 458-6280 μ m) and mean ONL was 82
352 μ m (31, 31-147 μ m). **Figure 8** presents the distribution of EZW and ONL
353 thickness with age. Sixteen of the patients with identifiable EZ and ONL, had
354 longitudinal assessment, with a mean follow-up of 5.3 years. The mean annual
355 rate of EZW loss was 219 μ m/year and the mean annual decrease in ONL
356 thickness was 4.93 μ m/year. No patient with childhood-onset disease had
357 identifiable EZ after the age of 26 years at baseline or follow-up.
358 **Supplementary Figure 9** shows representative examples of OCT scans from
359 3 adult patients with complete EZ loss.

360 The four patients with adulthood-onset disease (mean age, range: 36.6,
361 20.5-68.2 years old) had evidence of a relatively preserved EZ (mean width
362 3255 μ m, range 615-6500 μ m) and ONL (mean width: 89.63 μ m, range: 64-
363 129 μ m). Three of the four patients had longitudinal assessment after a mean
364 follow-up of 3.9 years. The one patient with age of onset 17 years progressed
365 to complete EZ loss over 7.4 years (age at follow-up: 31 years old). The two
366 patients with later onset disease had stable imaging after 1.4 and 2.9 years
367 (**Figure 10**).

368

369 **DISCUSSION**

370

371 This study details the clinical phenotype in the largest cohort of genetically
372 characterized patients with *RP2*-associated retinopathy to date, including novel
373 genetic findings. Comprehensive electrophysiological testing, natural history
374 and serial retinal imaging data highlight the structural and functional spectrum
375 and variability of the disease, with the aim of informing future patient
376 management and interventional trials. *RP2*-retinopathy is a predominantly
377 childhood-onset, rapidly progressive retinal degeneration, with macular
378 involvement and early complete loss of EZ in most cases.

379 In contrast with some other forms of progressive IRDs,^{39, 40} there was less
380 dissociation of structure and central vision; central vision was severely
381 decreased in all patients with childhood-onset disease and the OCT EZ was
382 undetectable by the age of 26 years in most cases. Electrophysiological testing
383 also revealed PERG evidence of macular dysfunction, severe in all but two
384 cases. Full-field ERGs were mostly consistent with rod-cone dystrophy but the
385 severity of generalized (mainly peripheral) retinal dysfunction varied greatly in
386 children and adolescents of a similar age, ranging from undetectable (severe
387 rod and cone photoreceptor dystrophy) to near-normal cone-mediated ERG
388 components (**Figure 3**).

389 Rare exceptions of adulthood-onset disease with relative preservation of
390 outer retinal structure (**Figure 10**) and the wide range of ERG abnormalities in
391 patients of a similar age (**Figure 3**), highlight the necessity of individual
392 assessments, important to the selection of candidates most suitable for clinical
393 trials and possible future treatment. Patients with complete loss of EZ and
394 geographic atrophy, irrespective of age, are less likely to benefit from attempts

395 to rescue/regain macular function or to arrest progressive maculopathy. There
396 was a rapid rate of progressive EZW reduction and decline in ONL thickness,
397 highlighting a relatively narrow window for intervention. Although clinically
398 significant structural changes are likely to be observed within a short time frame
399 in a clinical trial. However, the severity of degeneration may impose challenges
400 in the accurate measurement of such changes.

401 In the current cohort we identified three patients with a choroideremia-like
402 phenotype similar to some older patients described in a previous study of
403 XLRP,²⁰ with advanced degeneration and of older age. It should be noted that
404 none of those patients had a preserved island of vision and the choriocapillaris
405 atrophy may represent changes secondary to the chronic retinal atrophy. Those
406 cases may highlight the potential value of functional rescue of peripheral retinal
407 function in cases with severe maculopathy, particularly given that some may
408 have near-normal or relatively preserved cone-mediated ERGs (**Figure 3**).

409 Ideally, future prospective studies with standardized imaging acquisition
410 protocols need to establish the inter-session repeatability of measurements
411 before being employed as outcome measurements in trials. Also the use of
412 novel high-resolution imaging techniques such as adaptive optics scanning
413 laser ophthalmoscopy may be more sensitive to change.⁴¹ Prospective natural
414 history studies that monitor patients from a young age will be vital to better
415 establish prognosis, phenotype-genotype correlations and meaningful
416 endpoints for trials. Such studies can inform the design of planned treatment
417 trials, including recruitment criteria, assessments and follow-up time. The pre-
418 clinical work performed both assessing gene therapy, as well as read-through
419 drugs make *RP2*-retinopathy an attractive target for intervention.⁴²

420 The retrospective nature of the current study has inherent limitations.
421 Follow-up intervals were not standardized and the functional assessments did
422 not include visual field testing. Further investigation of female carriers that
423 manifest retinal disease will be of value, in order to determine disease severity
424 and inform counselling; moreover, they may also be candidates for
425 intervention.²⁴

426 This report of a large *RP2*-associated retinal dystrophy cohort helps to
427 define the phenotypic and genetic spectrum. The disorder is characterized by
428 childhood-onset retinal degeneration usually with early macular involvement.
429 Full-field ERGs reveal rod-cone dystrophy in the vast majority, with generalized
430 (peripheral) cone system involvement of widely varying severity in the first two
431 decades of life, and OCT imaging shows early complete EZ loss. Novel
432 therapies for *RP2* are under advanced development and clinical trials are
433 anticipated in the near future. The findings of this study will inform patient
434 management and counselling and are pertinent to the appropriate selection of
435 patients in future clinical trials.

436

437 **FOOTNOTES**

438

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445 SHU, MM, AW and AJH conceived, supervised, and revised the manuscript. All

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451

452 **LEGENDS**

453

454 **Figure 1: Schematic representation of variants in the *RP2* gene and**
455 **protein.**

456 The identified variants are marked along the corresponding location of the *RP2*
457 gene and protein. Black shaded boxes represent the coding exons (exons 1 to
458 5) separated by introns (solid line), with the protein domains (bottom panel)
459 coded by each exon indicated with a dotted line.

460

461 **Figure 2: Visual Impairment**

462 (A) Scattered plot graph presenting mean baseline best corrected visual acuity
463 (BCVA) against age, and (B) stacked scatter plot depicts the age distribution
464 among the different categories of visual impairment based on World Health
465 Organization classification. As expected a greater degree of impairment was
466 present in older patients, with the exception of patients with adulthood-onset
467 disease (open diamonds).

468

469 **Figure 3: Full field ERG**

470 Full field ERG findings summarized in 22 subjects tested according to the
471 ISCEV standard methods; a) The amplitudes of the DA0.01 ERG, DA 10 ERG
472 a-wave, LA 30 Hz ERG and LA 3 ERG b-wave are plotted against the primary
473 axis as a percentage of the age-matched lower limit of the (“normal”)
474 reference range (horizontal broken line), with values arranged in ascending
475 order of DA10 ERG a-wave amplitude for clarity. The LA 30 Hz peak times
476 are plotted as a difference from the age-matched upper limit of normal timing
477 (horizontal dotted line) against the secondary axis. b) The age of the patients
478 at the time of testing, arranged in same order as in a).

479

480 **Figure 4: Representative full-field and pattern ERGs**

481 Patient 4 (a; aged 13 years), 12 (b; 13 years) and 21 (c; 7 years) correspond
482 to the patient numbering used in **Figure 3**. Representative control (“normal”)
483 recordings are shown for comparison (d). Data are shown for the right eyes

484 only, as all showed a high degree of inter-ocular symmetry. Patient traces are
485 superimposed to demonstrate reproducibility. Broken lines replace blink
486 artefacts for clarity. In all 3 patients there is ERG evidence of rod-cone
487 dystrophy. Pattern ERG P50 abnormalities are consistent with macular
488 involvement that is a) severe, b) moderate or c) relatively mild.

489

490 **Figure 6: Fundus Autofluorescence (FAF) Imaging**

491 FAF imaging of six patients with *RP2*-associated retinopathy at different
492 stages of the disease. (A) Normal pattern of autofluorescence. (B)
493 Midperipheral patchy signal, with early patchy foveal pattern. (C)
494 Midperipheral patchy signal, with increased foveal signal. (D) Foveal atrophy,
495 without midperipheral changes. (E) Midperipheral patchy signal, with foveal
496 atrophy. (F) Diffuse atrophic changes.

497

498 **Figure 8: Optical Coherence Tomography (OCT) Graphs**

499 Scattered plots presenting (A) ellipsoid zone width (EZW) and age, and (B)
500 outer nuclear layer (ONL) and age. Greater degree of impairment of structural
501 loss is present in older patients, except for patients with adulthood-onset
502 disease (open diamonds). No patient with childhood-onset disease had
503 identifiable EZ or ONL after the third decade of life.

504

505 **Figure 10: Optical Coherence Tomography (OCT) Imaging**

506 OCT imaging of two patients with *RP2*-associated retinopathy with (A)
507 childhood-onset disease, and (B) adulthood-onset disease. (A) Patient shows
508 progressive loss of the ellipsoid zone over a follow-up of seven years, with no
509 identifiable EZ by the age of 22 years old. (B) Patient had a well-preserved
510 ellipsoid zone at age 46 years old and no ellipsoid zone loss was observed over
511 three years of follow-up.

512

513

514

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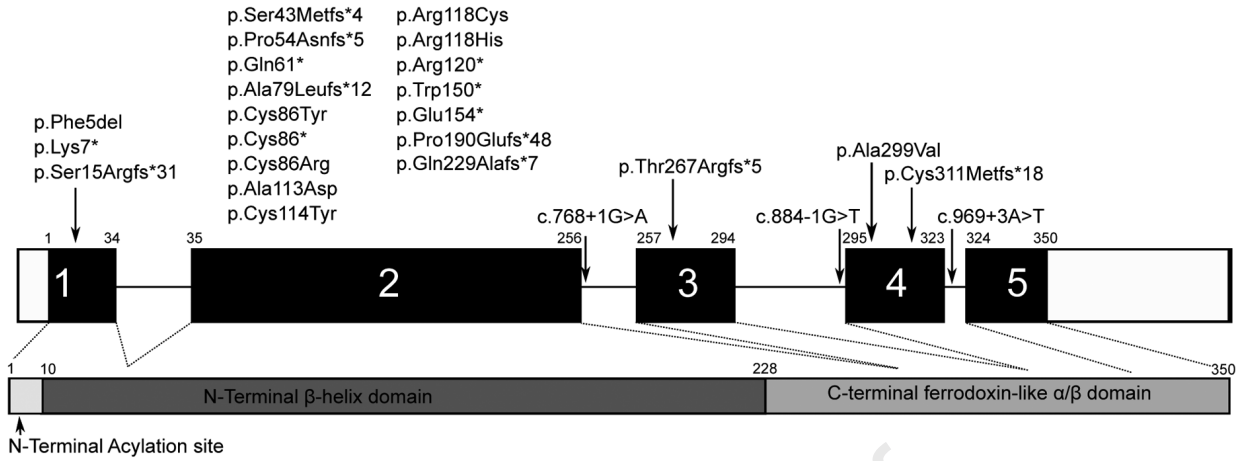
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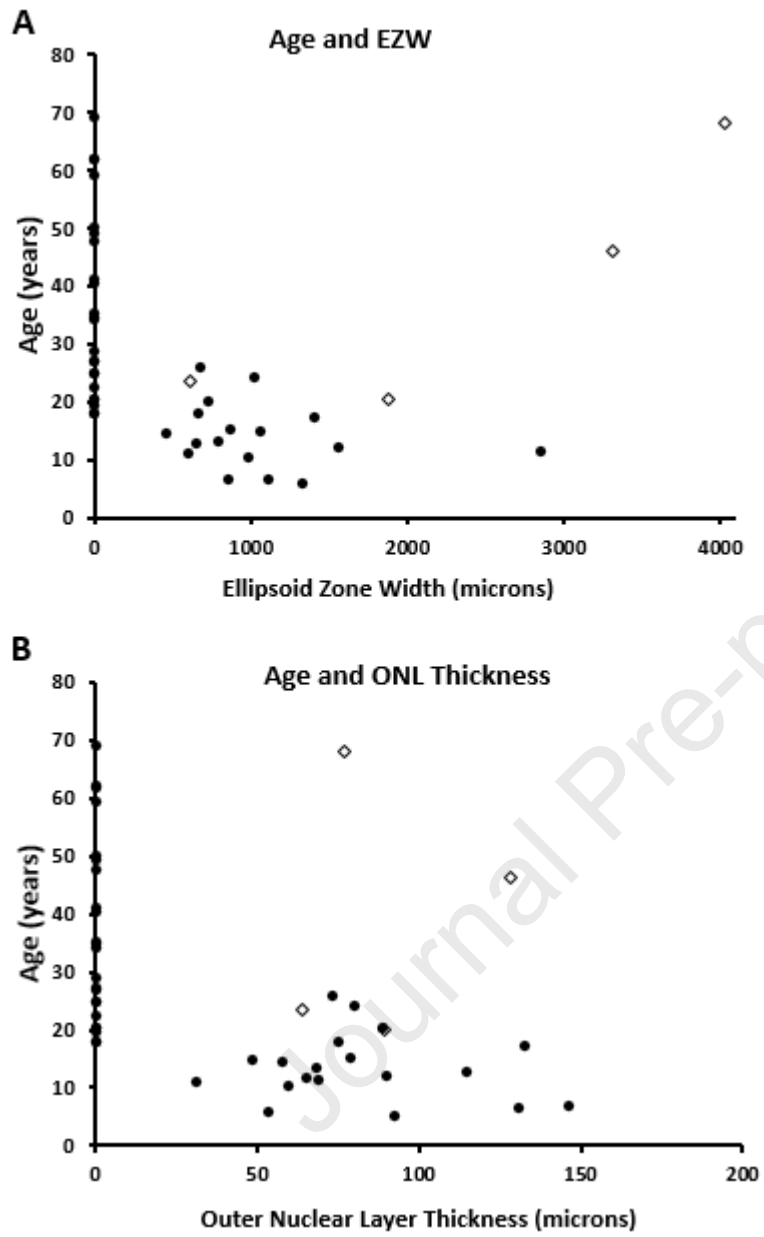
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632

Table 1: Clinical Data and Visual Impairment in *RP2*-Rod-Cone Dystrophy

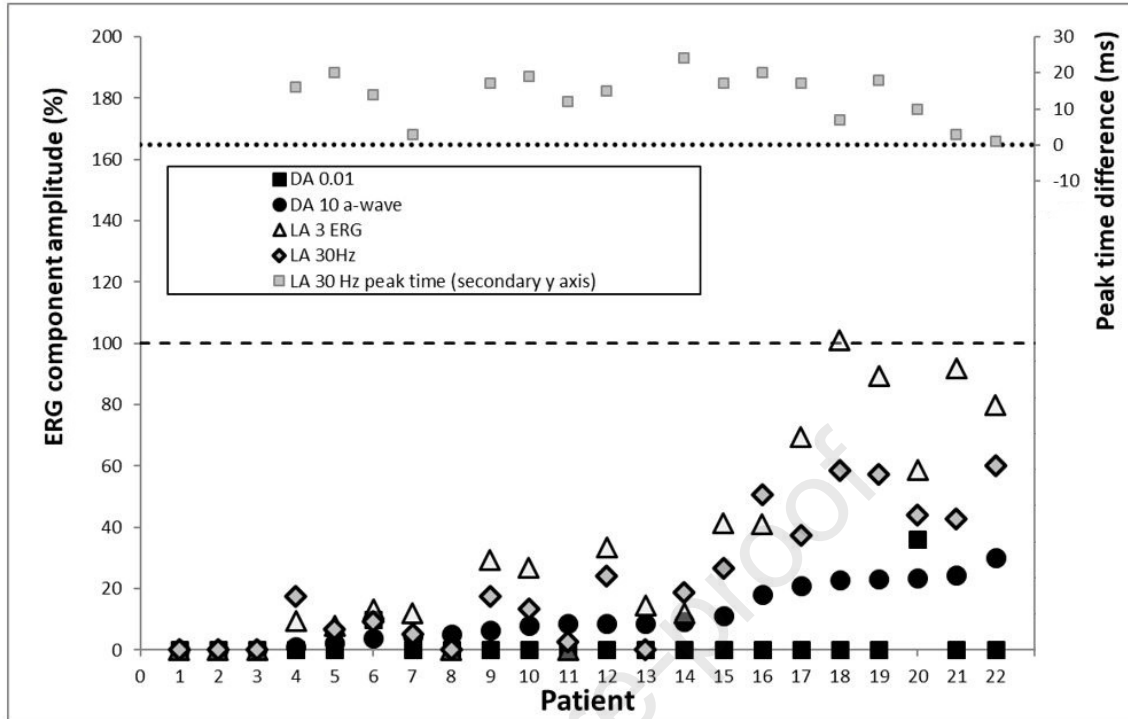
Parameter-Characteristic		
Age of Onset	(n=)	Mean (\pmSD Range,)
Rod-Cone Dystrophy	48	9.63 \pm 9.20, 1-57 years
Childhood-Onset	44 (91.7%)	7.55 \pm 4.10, 1-16 years
Adulthood-Onset	4 (8.3%)	32.5 \pm 16.15, 17-57 years
Presenting Symptoms*		
Night Blindness	33 (68.8%)	
Decreased Central Vision	8 (16.7%)	
Night Blindness and Decreased Central Vision	4 (8.3%)	
Decreased Central and Peripheral Vision	2 (4.2%)	
Nystagmus	1 (2.1%)	
Best Corrected Visual Acuity (BCVA)		
Age at Baseline, n=53		23.2 \pm 17.4, 3.8-71 years
Right Eye Mean BCVA at Baseline		0.91 \pm 0.80, 0-2.7 LogMAR
Left Eye BCVA at Baseline		0.94 \pm 0.78 0-2.7 LogMAR
Mean Follow-up, n=43		7.3 \pm 7.1, 0.3-30.2 years
Right Eye BCVA at Follow-up		1.17 \pm 0.84, 0.16-3.0 LogMAR
Left Eye BCVA at Follow-up		1.16 \pm 0.78, 0.16-3.0 LogMAR
Disease Severity		
Baseline, n=53		
Mild Disease	21 (39.6%)	
Severe Disease	32 (60.4%)	
Follow-up, n=50		
Mild Disease	10 (20%)	
Severe Disease	40 (80%)	
WHO Visual Impairment		
No or mild visual impairment	24 (45%)	
Moderate impairment	11 (21%)	
Severe impairment	7 (13%)	
Blindness	11 (21%)	

*The single patient with cone-rod dystrophy presented with decreased central vision.

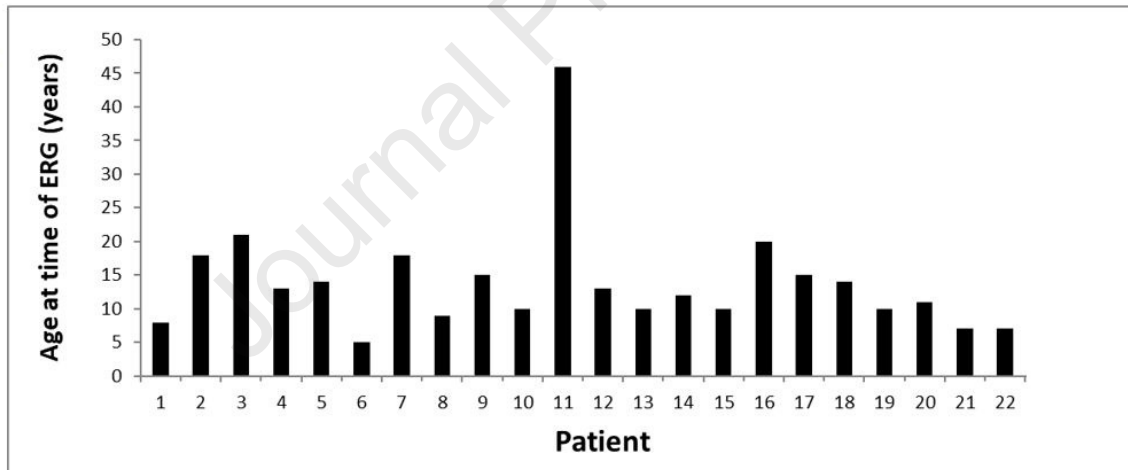


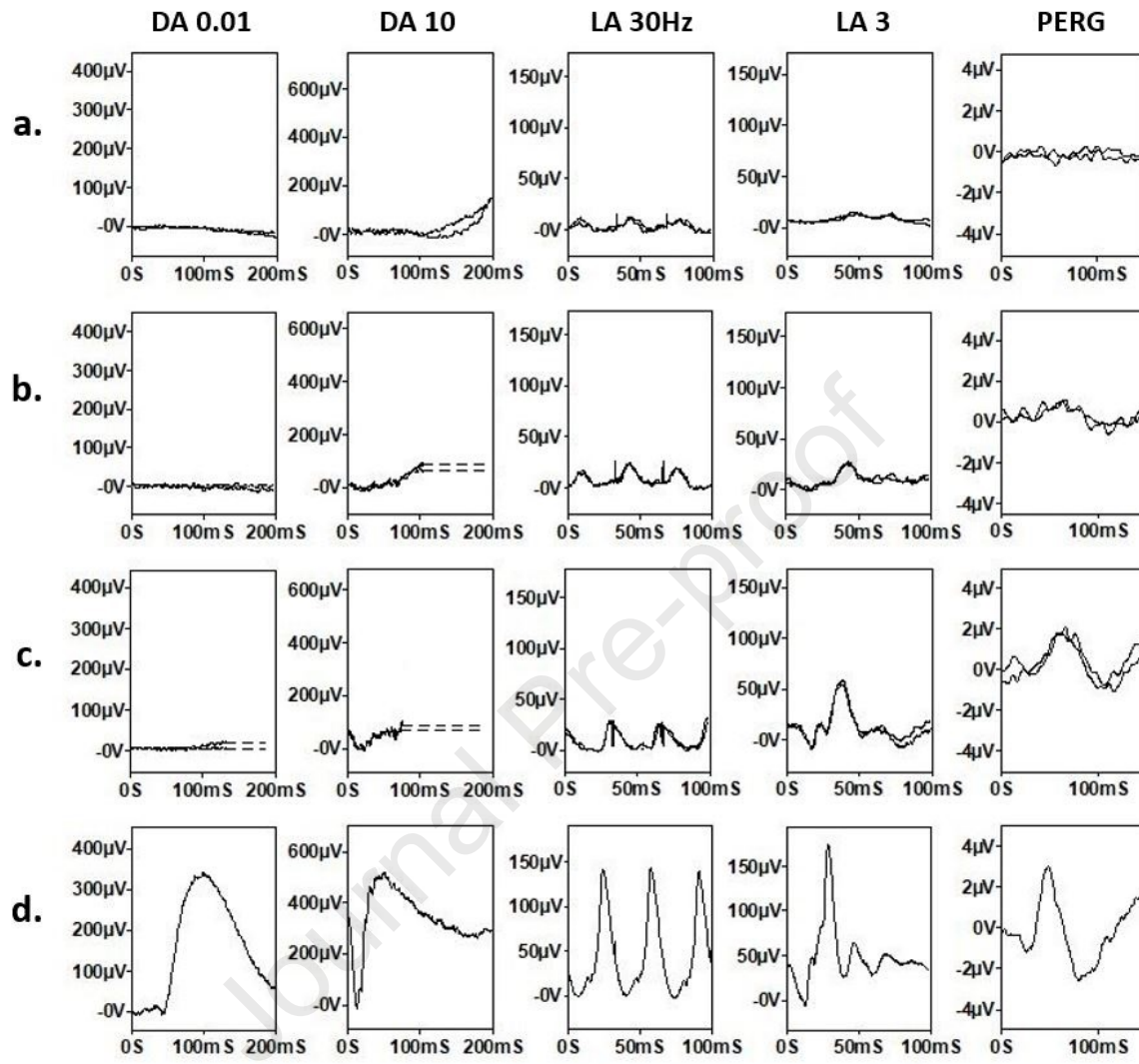


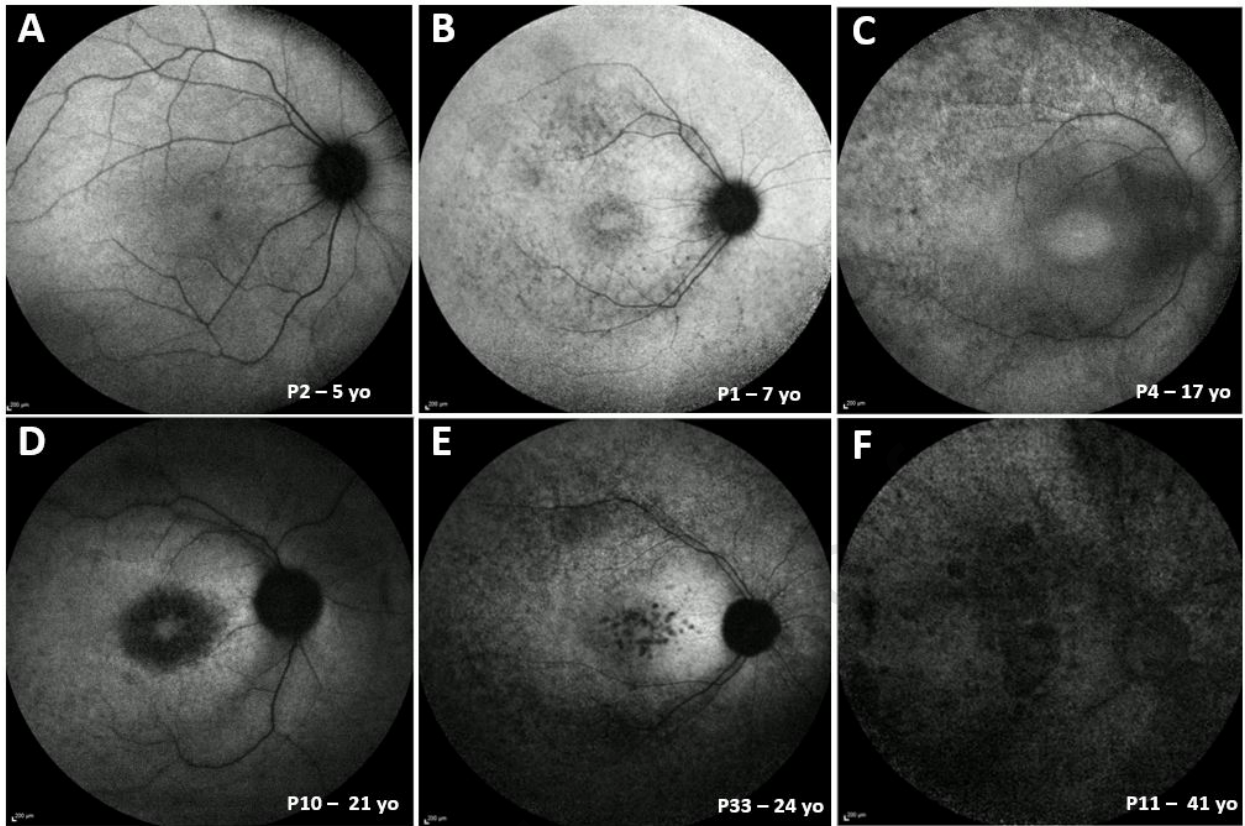
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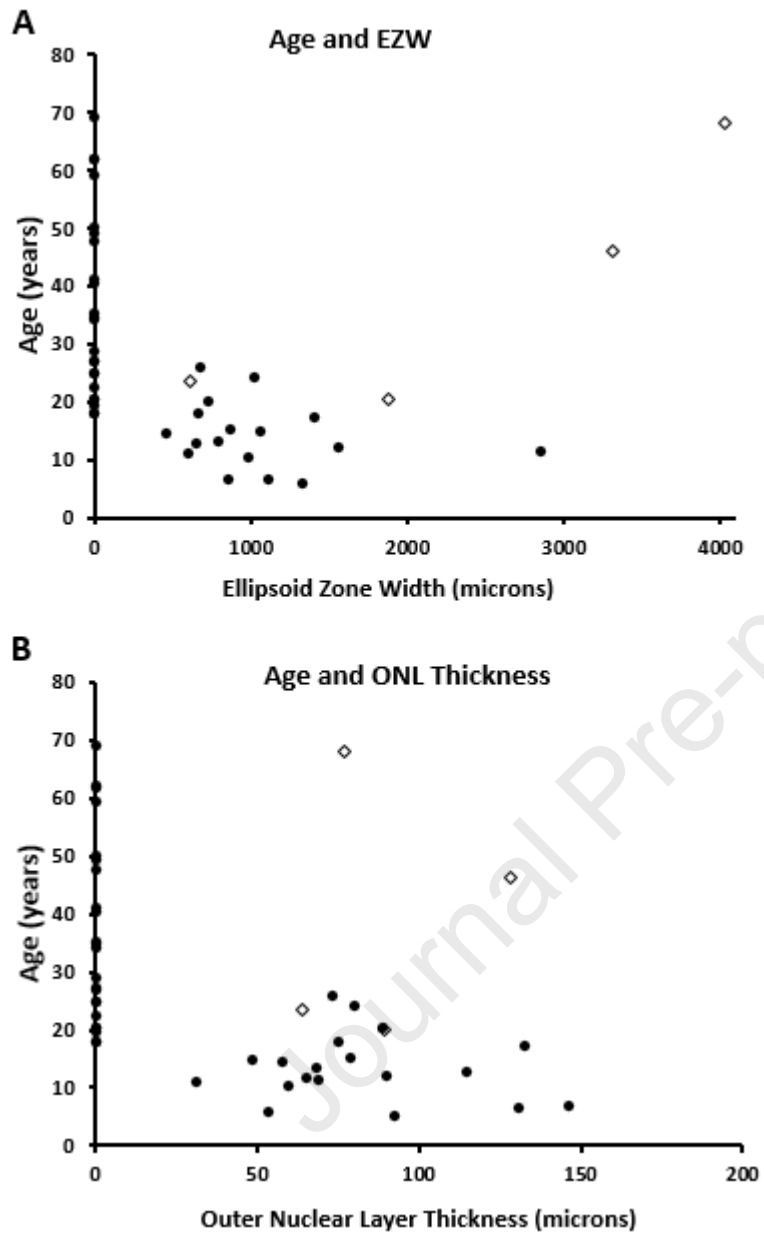
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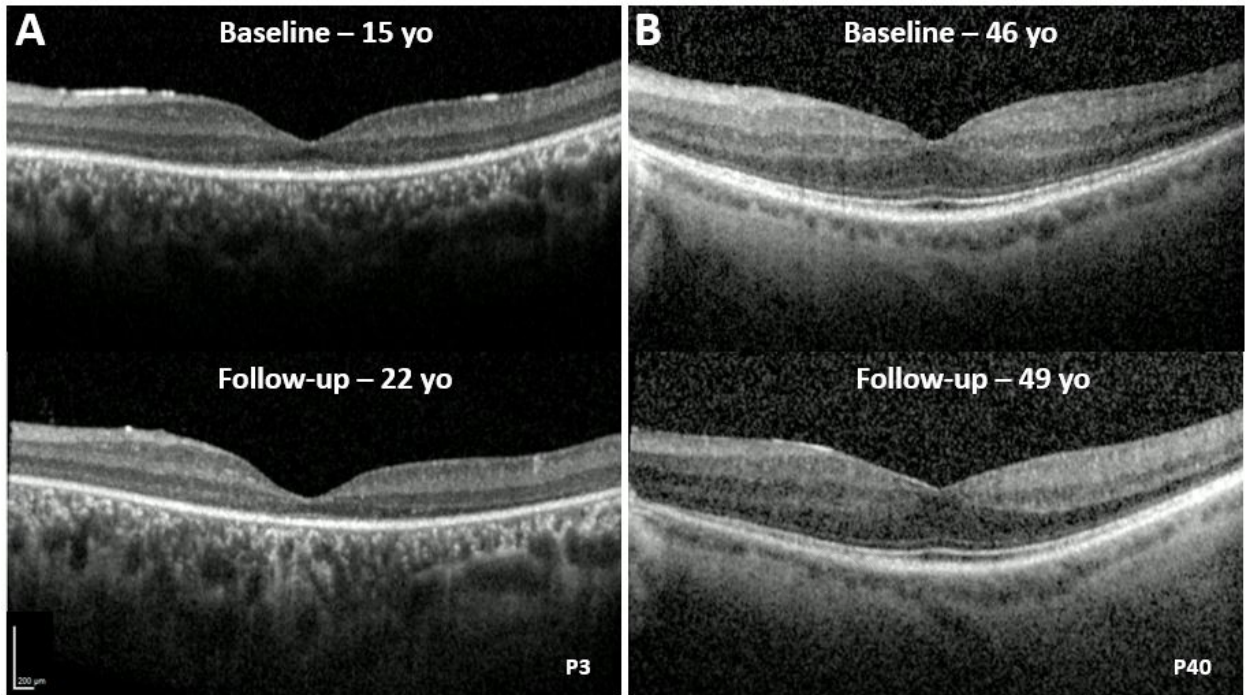






Journal





Precis

Deep phenotyping of the functional and anatomical characteristics of patients with *RP2*-associated retinopathy, in a large cohort, in preparation for planned novel therapeutic interventions.

Journal Pre-proof