Article

Detecting Progression of Retinitis Pigmentosa Using the Binomial Pointwise Linear Regression Method

Shotaro Asano¹,², Akio Oishi³,⁴, Ryo Asaoka¹,⁵⁻⁸, Yuri Fujino¹,⁵,⁹, Hiroshi Murata¹, Keiko Azuma¹, Manabu Miyata³, Ryo Obata¹, and Tatsuya Inoue¹,¹⁰

¹ Department of Ophthalmology, The University of Tokyo, Graduate School of Medicine, Tokyo, Japan
² Department of Ophthalmology, Asahi General Hospital, Asahi, Chiba, Japan
³ Department of Ophthalmology and Visual Sciences, Kyoto University Graduate School of Medicine, Kyoto, Japan
⁴ Department of Ophthalmology and Visual Sciences, Nagasaki University, Nagasaki, Japan
⁵ Department of Ophthalmology, Seirei Hamamatsu General Hospital, Shizuoka, Japan
⁶ Seirei Christopher University, Shizuoka, Japan
⁷ Nanovision Research Division, Research Institute of Electronics, Shizuoka University, Shizuoka, Japan
⁸ The Graduate School for the Creation of New Photonics Industries, Shizuoka, Japan
⁹ Department of Ophthalmology, Shimane University Faculty of Medicine, Izumo, Japan
¹⁰ Department of Ophthalmology and Micro-Technology, Yokohama City University, Kanagawa, Japan

Correspondence: Ryo Asaoka, Department of Ophthalmology, Graduate School of Medicine and Faculty of Medicine, The University of Tokyo, Tokyo 113-8655, Japan. e-mail: rasaoka-tky@umin.ac.jp

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Introduction

Retinitis pigmentosa (RP) is a progressive hereditary retinal disease caused by the degeneration of rod and cone photoreceptors.¹² RP is characterized by night blindness, loss of fine acuity, and a constricted visual field (VF).¹ VF can be measured with a Humphrey field analyzer (HFA) (Carl Zeiss Meditec, Dublin, CA) in eyes with RP,³⁴ which typically have scotomas that enlarge over a period of years owing to the loss of rod and cone function.¹ Visual acuity can remain good even in advanced cases, typically with a small island of remaining central VF.¹ An assessment of central VF, such as the HFA 10-2 test, is of critical importance in evaluating visual disability.¹,⁵ However, a method of evaluating central VF progression in eyes with RP remains to be established.
The VF progression rate is usually evaluated by trend analysis. The mean deviation (MD) trend analysis is frequently used to clinically assess the rate of VF progression, with a HFA. However, it is not an ideal method to detect focal VF progression, because the MD is the average of VF damage over the entire VF. In contrast, an analysis can be undertaken using pointwise linear regression (PLR). Previous studies have reported the advantage of PLR in the early detection of VF progression, particularly in glaucomatous eyes. However, one drawback of PLR is that the progression over the entire VF cannot be analyzed. The application of permutation analyses of PLR (PoPLR) and binomial tests to PLR (binomial PLR) are solutions to this problem. These methods integrate information on the VF progression rate at each test location, and enable an assessment to be made of VF progression as a whole. We previously reported that the binomial PLR method gives more consistent results than MD trend analysis or PoPLR in analyzing HFA 24-2 test results.

There is no treatment for RP, except retinal prosthesis for advanced cases and gene therapy for patients with specific mutations. The inability to evaluate the progression is one of the reasons why clinical trials have not been successful in RP. The establishment of sensitive and reproducible methods of evaluating VF progression is desirable. One example of this effort is a model to predict the central VF sensitivity from the structural damage demonstrated by fundus autofluorescence imaging in RP. In the current study, we investigated whether binomial PLR is useful in evaluating VF progression in the central 10°, in eyes with RP.

### Methods

The study was approved by the Institutional Review Boards of the University of Tokyo and Kyoto University and adheres to the tenets of the Declaration of Helsinki. The study protocols did not require that each patient provide written informed consent. Instead, the protocol was posted at the outpatient clinic to notify participants of the study.

### Subjects

VF data were retrospectively obtained of 196 eyes in 103 patients with RP. Patients were followed in the retinal dystrophy clinics at the University of Tokyo Hospital and Kyoto University Hospital between February 1, 2005, and January 31, 2020. All patients underwent at least 10 measurements with the 10-2 HFA test pattern, excluding the initial VF. Inclusion criteria in this study were (1) typical fundus findings of RP, such as bone spicule pigmentation, arteriolar attenuation, and waxy disc pallor; (2) a decrease in a-wave and b-wave amplitudes or nondetectable full-field electroretinogram; (3) RP was the only disease causing VF damage; (4) no other diseases of the anterior and posterior segments of the eye that could affect VF, including cataracts, except for clinically insignificant senile cataracts; and (5) patient age at least 20 years. Patients who underwent intraocular surgery, including cataract surgery, during the observation period were excluded. VF measurements were performed using HFA with the 10-2 program and the Swedish Interactive Threshold Algorithm standard. Reliability criteria for VF data were applied, including fixation losses of less than 20% and false-positive responses of less than 15%, following the manufacturer’s recommendation. The VF of the left eye was mirror-imaged to that of the right eye for statistical analyses.

### MD Trend Analysis

In the MD trend analysis, the MD value was linearly regressed against time, using 10 VFs (VF1–10; the series of VFs from 1 to 10). We defined a series of VF tests to be significant if the MD slope was less than zero and the calculated P value was less than 0.025. Otherwise, it was not significant. In the current study, the progression assessment for VF1–10 was regarded as a surrogate for the absolute true progression. Similar analyses were carried out with shorter VF sequences (from VF1–5 to VF1–9), and their reliability (consistency) was evaluated through three surrogate measures for the true positive rate (the proportion of both VF1–10 and shorter series progressing [PBP]); the true negative rate (the proportion of both VF1–10 and shorter series not progressing [PBNP]); and the false positive rate (the proportion of inconsistent progression [PIP]). Figure 1 shows the concept of PBP, PBNP, and PIP.

(1) PBP is the probability that both the complete series of VFs (VF1–10) and each of the shorter series of VFs (from VF1–5 to VF1–9) were classified as progressive.

(2) PBNP is the probability that the complete series of VFs (VF1–10) and each of the shorter series of VFs (from VF1–5 to VF1–9) were classified as not progressive.

(3) PIP is the probability that each of the shorter series of VFs (from VF1–5 to VF1–9) was classified as progressive.
as progressive while the complete series of VFs (VF1–10) was classified as not progressive.

We also carried out a Kaplan–Meier survival analysis to compare the time required to detect progression, following our previous studies.13,16,20

(4) The time required to detect progression was assessed using the Kaplan–Meier survival analysis, and compared using a log rank test.

**Binomial PLR**

The calculation of the binomial PLR method is described in detail in our previous reports.13 The assumption in PLR is that VF damage progresses linearly over time. PLR analyses are usually conducted by regressing the total deviation values (as an independent variable) over time (as a dependent variable), similar to the MD and VFI trend analysis in the Guided Progression Analysis in HFA. Here, the null hypothesis

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**Figure 1.** The demonstrative figure of PBP, PBNP, and PIP.

<table>
<thead>
<tr>
<th>VF1-5</th>
<th>VF1-10</th>
</tr>
</thead>
<tbody>
<tr>
<td>Progressive</td>
<td>Progressive</td>
</tr>
<tr>
<td>Non-progressive</td>
<td>Non-progressive</td>
</tr>
<tr>
<td>Non-progressive</td>
<td>Progressive</td>
</tr>
</tbody>
</table>

**Figure 2.** An example of calculating a P value with the binomial PLR method.
Table. Demographics of the Eyes

<table>
<thead>
<tr>
<th>Variables</th>
<th>No. or Mean ± Standard Deviation</th>
</tr>
</thead>
<tbody>
<tr>
<td>No. of eyes</td>
<td>196</td>
</tr>
<tr>
<td>No. of subjects</td>
<td>103</td>
</tr>
<tr>
<td>Eye laterality (right/left)</td>
<td>100/96</td>
</tr>
<tr>
<td>Age, years</td>
<td>47.7 ± 12.1</td>
</tr>
<tr>
<td>MD at the baseline, dB</td>
<td>−16.4 ± 8.0</td>
</tr>
<tr>
<td>Follow-up, years</td>
<td>8.1 ± 1.9</td>
</tr>
<tr>
<td>MD progression rate, dB/y</td>
<td>−0.55 ± 0.53</td>
</tr>
</tbody>
</table>

was that the change of the entire VF was equal to zero. Under this null hypothesis, the slope coefficient $P$ value from linear regression at each test point is uniformly distributed between zero and 1. If we are to accept the null hypothesis, we would expect the numbers of test points to follow a binomial distribution. Hence, the numbers of test points with $P$ values lower than 0.025, 0.050, 0.075, and 0.100 should follow the binomial distribution. If the null hypothesis is rejected, a slope coefficient of zero would be considered unlikely to be the result of random chance. Thus, the significance of the entire VF progression was assessed using these four cut-off $P$ values. The median value was then used to determine progression to merge the four $P$ values.21,22 We defined a series of VF tests to be significant if the slope was less than zero and the $P$ value calculated by binomial PLR was less than 0.025. Otherwise it was not significant. Figure 2 shows an example of calculating a $P$ value with the binomial PLR method. Thereafter we calculated PBP, PBNP, PIP, and the Kaplan–Meier survival analysis with the log rank test, in a similar manner to the MD trend analysis.

PoPLR

PoPLR uses the permutation analyses of PLR to sum the statistical significance for VF deterioration, according to the individual patient’s data (and its variability).14 The PoPLR method was carried out with total deviation values at each test point, using the $R$ package “visualFields” modified to deal with the HFA 10-2 test. Here, we defined a series of VF tests to be significant if the slope was less than zero and the $P$ value calculated by PoPLR was less than 0.025. Otherwise it was not significant. We then calculated the PBP, PBNP, PIP, and the time required to detect a significant progression.

All statistical analyses were carried out using the statistical program $R$ software (version 3.6.3; http://www.r-project.org/). The $P$ values in multiple comparisons were corrected using the Hochberg correction.

Results

Demographics of the eyes are shown in the Table. The average baseline MD was −16.4 ± 8.0 dB, and the initial age of the patients was 47.7 ± 12.1 years. The mean observation period between VF1 and VF10 was 8.1 ± 1.9 years. The average rate of MD progression was −0.55 ± 0.53 dB/y.

Figure 3 shows the PBP with the binomial PLR, PoPLR, and MD trend analysis. These values varied between 0.55 (VF1–5) and 0.95 (VF1–9) with binomial PLR, between 0.33 (VF1–5) and 0.92 (VF1–9) with PoPLR, and between 0.39 (VF1–5) and 0.89 (VF1–9) with MD trend analysis. There was no significant difference among the PBP values obtained with the three methods (all $P$ values > 0.05, in a Wilcoxon signed rank test adjusted for multiple comparisons using the Hochberg correction).

Figure 4 shows the PBNP with binomial PLR, PoPLR, and MD trend analysis. These values varied between 0.67 (VF1–8) and 0.92 (VF1–9) with binomial PLR, between 0.92 (VF1–6, VF1–7, and VF1–8) and 0.94 (VF1–5 and VF1–9) with PoPLR, and between 0.92 (VF1–8) and 0.98 (VF1–5) with MD trend analysis. There was no significant difference among the PBNP

Figure 3. The rates of PBP with binomial PLR, PoPLR, and MD trend analysis. The rates of PBP were compared across binomial PLR, PoPLR, and MD trend analysis. There was no significant difference between the PBP rates of the four methods (all $P$ values > 0.05, in a Wilcoxon signed rank test adjusted for multiple comparisons using the Hochberg correction).
The rates of PBNP with binomial PLR, PoPLR, and MD trend analysis. The rates of PBNP were compared across binomial PLR, PoPLR, and MD trend analysis. There was no significant difference between the PBNP rates of the four methods (all $P$ values > 0.05, in a Wilcoxon signed rank test adjusted for multiple comparisons using the Hochberg correction).

Figure 5. The rates of PIP with binomial PLR, PoPLR, and MD trend analysis. The rates of PIP were compared across binomial PLR, PoPLR, and MD trend analyses. There was no significant difference between the PIP rates of the four methods (all $P$ values > 0.05, in a Wilcoxon signed rank test adjusted for multiple comparisons using the Hochberg correction).

The Kaplan–Meier survival analysis with binomial PLR, PoPLR, and MD trend analysis. The results of the Kaplan–Meier survival analysis with binomial PLR, PoPLR, and MD trend analyses are shown. Log rank tests indicate that the binomial PLR method detected progression significantly earlier than PoPLR and MD trend analysis ($P < 0.01$ and $P < 0.001$, respectively; log rank tests adjusted for multiple comparisons using the Hochberg correction). PoPLR detected progression significantly earlier than MD trend analyses ($p = 0.02$, log rank tests adjusted for multiple comparisons using the Hochberg correction).

Figure 6 shows the PIP for binomial PLR, PoPLR, and MD trend analysis. These values ranged from 0.08 (VF1–9) to 0.33 (VF1–8) with binomial PLR, from 0.06 (VF1–5 and VF1–9) to 0.08 (VF1–6, VF1–7, and VF1–8) with PoPLR, and from 0.02 (VF1–5) to 0.08 (VF1–8) with MD trend analysis. There was no significant difference among the PIP values obtained with the three methods (all $P$ values > 0.05, in a Wilcoxon signed rank test adjusted for multiple comparisons using the Hochberg correction).

Discussion

The reliability (consistency) and sensitivity of the binomial PLR in detecting VF progression was analyzed through HFA 10-2 tests on 196 eyes of 103 patients with RP, and compared with those of MD trend analysis and the PoPLR method. A Kaplan–Meier survival analysis and a log rank test revealed that binomial PLR detected VF progression significantly earlier than the other two methods. However, a possible caveat remains that the earlier detection of progression with the binomial PLR is because of detecting falsely too many progressions. We conducted the PBP, PBNP, and PIP analyses shedding light on this issue, and, as a result, these values were not significantly different among the three approaches. These findings suggest that the binomial PLR detect progression significantly earlier than MD trend analysis and the PoPLR method without sacrificing the accuracy (reliability).

The reliability (consistency) of the diagnosis with each method (i.e., binomial PLR, MD trend analysis, and PoPLR) was investigated by comparing the diagnosis with VF1–10 and those with shorter VF sequences. This is because the VF continuously progresses in RP and the diagnosis should not be changed over time, once diagnosed as progressive. In our previous study, we evaluated the PBP, PBNP,
and PIP through binomial PLR, PoPLR, and MD trend analysis in cases of glaucomatous eyes, using the HFA 10-2 test. The PBP, PBNP, and PIP values of binomial PLR were not significantly different from those of PoPLR and MD trend analysis. The current results on tests of eyes with RP agree with these results, suggesting that binomial PLR is a reliable method of evaluating VF progression in the central 10°, not only in eyes with glaucoma, but also those with RP. Furthermore, similar to the results of our studies of glaucomatous eyes, Kaplan–Meier survival analyses indicated that binomial PLR detected VF progression significantly earlier than PoPLR and MD trend analysis, suggesting that binomial PLR is useful in the early detection of VF progression in RP eyes as well. PoPLR was developed to disclose the progression of the entire VF in glaucomatous eyes at an earlier stage, compared with the MD trend analysis. The current results suggest that this technique is also useful in eyes with RP, but that there remains a greater advantage with the binomial PLR method. In contrast, a possible associated caveat is that the binomial PLR method has not been readily useable in the clinical setting. It would be clinically beneficial to develop software and support tools to detect VF progression, as introduced in this study, similar to PROGRESSOR (Medisoft, Inc., London, UK).

The binomial PLR method analyzes the results of PLR, which uses linear regression analysis. A case could be made for applying a nonlinear regression analysis, such as an exponential model. However, we have shown that linear regression outperforms nonlinear regression in predicting future VF, using the HFA 24-2 test in glaucoma. Moreover, we recently investigated the use of nonlinear regression models using the HFA 10-2 test, and concluded that they showed no advantage over the conventional linear regression method. Besides, a non-negligible problem associated with a nonlinear regression analysis such as the exponential model is that $P$ values cannot be calculated; thus, the binomial PLR method cannot be applied. These results support the view that it would be advantageous to apply binomial tests to PLR, not nonlinear regressions, in glaucoma and also in RP.

The accurate and early detection of VF progression is important in RP management, because the extent of the defects in the VF is related directly to the quality of vision in patients with RP. As suggested in a previous study, visual disability is closely associated with HFA 10-2 test results. The current results suggest that the application of the binomial PLR method to the HFA 10-2 test will aid the estimation of patients’ future condition. In addition, considering the difficulty that remains in devising a treatment strategy to alleviate the progression of RP, the accurate assessment of VF progression is important for the establishment of treatment. Some clinicians prescribe oral vitamin A to patients with RP; however, the treatment outcomes varied across the studies. Docosahexaenoic acid, an omega-3 fatty acid highly present in oily fish, has also been proposed as an additional nutritional treatment, but its effectiveness remains uncertain.

One possible reason for these contradictory results is that the assessments of visual function differed across the studies. For instance, visual acuity was often used in the assessment of visual function. However, visual acuity merely reflects the retinal function around the fovea and tends to be insensitive to the severity of disease. For instance, we previously developed a model to predict VF sensitivity measured with the HFA 10-2 test using visual acuity in RP. The study revealed that the model improved when the assessment of the structural damage demonstrated with fundus autofluorescence imaging was combined, implying the limitation of using visual acuity in establishing the severity of disease in RP. The current results suggest that the binomial PLR method will be useful in assessing VF progression with the HFA 10-2 test when evaluating the effects of any candidate treatment.

There are several limitations in the current study. First, we did not consider the patients’ genetic information. Recently, owing to gene-specific approaches including gene therapy, genetic information is of growing interest in the management of RP. For instance, variants in many genes, such as $ABCA4$, $PRPH2$, and $PROM1$ genes, are associated with various phenotypes. Owing to its retrospective nature, no genetic information was included in this study. Thus, the evaluation of VF progression with certain genetically defined forms of RP remains to be investigated in a future study. Second, the PBP, PBNP, and PIP were calculated with VF$_{1-10}$ as a surrogate for the absolute true progression. The absolute true progression cannot be stated. Thus, the current study may require validation using a simulation dataset. In the current study, we hypothesized the linear regression at each test point. This assumption was based on our previous report that linear regression was most suitable in glaucomatous eyes; however, this point needs further investigation, specifically in RP. We limited our analysis to static perimetry measures because the process to standardize and equate static and kinetic VF measures for this application would require additional extensive modeling. Previously, we reported that microperimetry with fundus tracking (MP-3, Nidek Co. Ltd, Gamagori, Japan) demonstrated a better structure–function relationship in glaucomatous eyes than did HFA. Based on the current study’s results,
the application of a PLR method to the results of microperimetry may also achieve improved assessments in central VF progression in RP. This point remains to be investigated in a future study. Similarly, future studies should concentrate on comparisons of the usefulness of the proposed approach and other nonstandard VAs, such as low luminance VA and contrast sensitivity.48,49

In conclusion, we applied a binomial test to the PLR analysis of the HFA 10-2 test of eyes with RP, and revealed that binomial PLR achieved reliable and earlier detection of VF progression.

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