

1 Predicting visual acuity by using machine learning in patients treated
2 for neovascular age-related macular degeneration

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10 Abbreviations:

11	AMD	Age-related macular degeneration
12	CRT	Central retinal thickness
13	EMR	Electronic medical records
14	ETL	Extract, transform, load
15	DW	Data warehouse
16	GEID	Generic eye identifier
17	ICD-10	International classification codes of diseases version 10
18	MAE	Mean absolute error
19	nAMD	neovascular age-related macular degeneration
20	OCT	Optical coherence tomography
21	PRN	Pro re nata treatment scheme
22	RMSE	Root mean square error
23	VA	Visual acuity
24	XML	Extensible Markup Language

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41 Abstract

42 Objective:

43 To predict, by using machine learning, visual acuity (VA) at 3 and 12 months in patients with
44 neovascular age-related macular degeneration (nAMD) after initial upload of 3 Anti-VEGF
45 (vascular endothelial growth factor) injections.

46 Design:

47 Database study

48 Subjects:

49 For the 3-month VA forecast, 653 patients (379 females) with 738 eyes and an average age
50 of 74.1 years were included. The baseline VA before the first injection was 0.54 logMAR (+/-
51 0.39). 456 of these patients (270 females, 508 eyes, average age: 74.2 years) had sufficient
52 follow-up data to be included for a 12-month VA prediction. The baseline VA before the first
53 injection was 0.56 logMAR (+/- 0.42).

54 Methods:

55 Five different machine learning algorithms (AdaBoost.R2, Gradient Boosting, Random
56 Forests, Extremely Randomized Trees, and Lasso) were used to predict VA in patients with
57 nAMD after treatment with 3 Anti-VEGF injections. Clinical data features came from a data
58 warehouse (DW) containing electronic medical records (41 features, e.g. VA) and
59 measurements features from optical coherence tomography (OCT) (124 features, e.g. central
60 retinal thickness). VA of patient eyes excluded from machine learning was predicted and
61 compared with the ground truth, namely the actual VA of these patients as recorded in the
62 DW.

63 Main Outcome Measure:

64 Difference in logMAR VA after 3 and 12 months after upload phase between prediction and
65 ground truth as defined above.

66 Results:

67 For the 3-month VA forecast, the difference between the prediction and ground truth was
68 between 0.11 logMAR (5.5 letters) mean absolute error (MAE)/0.14 logMAR (7 letters) root
69 mean square error (RMSE) and 0.18 logMAR (9 letters) MAE/0.2 logMAR (10 letters) RMSE.
70 For the 12-month VA forecast, the difference between the prediction and ground truth was
71 between 0.16 logMAR (8 letters) MAE/ 0.2 logMAR (10 letters) RMSE and 0.22 logMAR (11
72 letters) MAE/0.26 logMAR (13 letters) RMSE.
73 The best performing algorithm with regard to forecasts was the Lasso protocol.

74 Conclusions:

75 Machine-learning allowed VA to be predicted for three months after three anti-VEGF
76 injections, with a comparable result to VA measurement reliability. For a forecast after 12
77 months of therapy, VA prediction may help to encourage patients adhering to intravitreal
78 therapy.

79 Introduction

80 In computer science, machine learning is a term that covers diverse approaches to artificial
81 intelligence in computers and is nowadays quite a common phenomenon. The original aim,
82 which was stated as early as 1959, was for computers to have the ability to learn without
83 being explicitly programmed ¹. Many methods were tried over the years, but it took until the
84 1990s when machines started to beat humans. Such stories were then broadcast in the mass
85 media and gained a great deal of publicity. In 1996 and 1997, Garry Kasparov lost a chess
86 match against the IBM super-computer “Deep Blue”; this was the first time that a long-
87 standing world champion (1985-2000) was defeated by a computer. Another well-publicized
88 milestone was reached when IBM Watson beat two expert human players in the television
89 show “Jeopardy” in 2011 ².

90 In the medical sciences involving vision, several machine learning techniques have
91 been applied in various sub-specialties ³. Mostly, glaucoma and retinal imaging-related
92 problems have been addressed. As the number of patients with diabetic retinopathy is
93 continuing to rise ⁴, efforts have recently been made to detect early forms of diabetic
94 retinopathy by means of machine learning ⁵⁻⁷. These programs have been shown to provide
95 a highly sensitive tool for detecting such disease complications ⁸. First approaches have been
96 published for the automatic imaging analysis of optical coherence tomography (OCT) by
97 using machine learning ^{9,10}. Additional studies are underway to automate the detection and
98 classification of pathological features in eye imaging including fundus photography and
99 OCT¹¹.

100 Age-related macular degeneration (AMD) is still one of the leading causes of legally
101 defined blindness in industrial countries ¹². However, the use of anti-VEGF agents such as
102 bevacizumab, ranibizumab, or aflibercept in AMD and other retinal diseases such as diabetic
103 macular edema or venous occlusions have allowed, for the first time, the sustainable
104 improvement, stabilization, or slow-down of disease progression ¹³⁻²⁰. Nevertheless, even
105 after several years of experience, real-life results in patients vary greatly from results
106 obtained in clinical trials ^{21,22}.

107 To improve insights into our own clinical results and to facilitate clinical research, a
108 data warehouse (DW) has been set up in our institution ²³. It comprises data, such as
109 diagnoses, medication, and undergone surgery, from electronic medical records (EMR), as
110 developed in our institution based on the hospital-wide EMR system “SAP i.s.h.med”, from

111 over 320,000 patients. It also contains measurement data from OCT (e.g., central retinal
112 thickness, macular volume) and other measurement devices such as IOLMaster and
113 Pentacam.

114 For this study, we have used five modern machine learning algorithms to predict the
115 outcome of visual acuity (VA) after one year of anti-VEGF treatment in newly diagnosed
116 neovascular AMD patients. The base for the prediction was our DW with its clinical and
117 measurement data. The forecasting of VA might help to reduce the psychological pressure
118 on patients, as many fear sight loss as a consequence of the diagnosis of wet AMD²⁴. In
119 particular, before the first injections, patients are extremely anxious and nervous, as they do
120 not know what the outcome of the therapy will be and whether it is worth pursuing²⁵.

121 Material and Methods

122 Data warehouse as repository

123 To estimate VA after an initial upload of 3 anti-VEGF injections in patients with neovascular
124 AMD after 3 and 12 months of intravitreal therapy, we applied machine learning algorithms
125 on real-life data from our DW. This is updated every night with new and altered information
126 from the EMR (i.s.h.med, Cerner AG, Erlangen, Germany). The nightly transfer from EMR
127 includes diagnoses, clinical data such as VA and intraocular pressure, intravitreal injections,
128 medications, appointments, and surgical operations. Up until December 2016, it
129 incorporated data from 330,801 patients including 44,134 recorded intravitreal injections
130 and 402,001 VA measurements. Our Ethics Committee ruled that approval was not required
131 for this study. This study adheres to the tenets of the Declaration of Helsinki.

132 In addition to clinical data, 75,750 Spectralis OCT (Heidelberg Engineering, Heidelberg,
133 Germany) measurements, but no images, were extracted from the software “Heidelberg Eye
134 Explorer”. It provides an XML (Extensible Markup Language) export, which is a structured file
135 format of measurement values, as in the company’s software. This export contains data like
136 central retinal thickness (CRT), retinal volume and measurements in nine zones grid over the
137 macular region (Figure 1). The XML files have been incorporated into the DW with a custom-
138 made extract, transform, load (ETL) script, which is also programmed in Java. The DW itself
139 was built on a “Microsoft SQL” database server running in a database cluster in the hospital’s
140 datacenter. The PC used for the machine learning algorithms was a quad core Intel i5 (3.30
141 GHz) PC with 32 gigabytes of RAM, running on Ubuntu 16.04 LTS and Python (3.6.0). In

142 Python, the following libraries were used: scipy (Version: 0.18.1), numpy (Version: 1.12.0),
143 pandas (Version: 0.19.2), jupyter (Version: 1.0.0), scikit-learn (Version: 0.18.1), and Theano
144 (Version 0.8.2).

145 [Data Pre-processing](#)

146 For our test, we identified all patient eyes in the DW that, at any stage, received the
147 international classification codes of diseases version 10 (ICD-10) “H35.3”, which stands for
148 age-related macular degeneration, and which received at least 3 injections of anti-VEGF
149 medication (aflibercept, bevacizumab or ranibizumab) as noted in the surgical report. The
150 standard operating procedure for treating 153 nAMD in our institution requires monthly
151 injections for the first three months. After the initial upload of three injections, patients
152 were treated according to the most recent treatment recommendations of the German
153 Ophthalmological Society (Deutsche Ophthalmologische Gesellschaft – DOG)^{26, 27}.
154 Accordingly, a pro re nata scheme (PRN) was recommended until 2015. From 2015 onwards,
155 a treat and extend scheme was recommended alongside the PRN regime. This led to 456
156 patients with 508 eyes for the long-term prediction goal (365 days) and 653 patients with
157 738 eyes for the short-term prediction goal (90 days); these were the patients that remained
158 after the removal of all patients with invalid entries. The patients used for the long-term
159 prediction represent a subgroup of the short-term prediction group.

160 Before algorithms can be trained on these patients, data need to be pre-processed and
161 joined into a single dataset. The data came from the EMR and the OCT measurements. Most
162 of the machine learning algorithms expect a dataset with no missing values; this is difficult to
163 achieve in real-life datasets such as ours. Therefore, we centered all the other values around
164 VA measurements. Consequently, we associated VA values with the most recently available
165 measurements from previous (but not future) records in order to rule out potential data
166 mix-ups shortly before and after an intravitreal medication application.

167 In the data collection process, we treated every eye as a separate patient and associated
168 it with a generic eye identifier (GEID). The newly created dataset included not only the most
169 recent measurement values (e.g., VA and CRT), but also aggregated versions with different
170 timeframes (3 months and 12 months) and different aggregation functions (mean, variance,
171 minimum and maximum). These synthetic features were designed in order to capture trends
172 in the temporal progression of the disease. After all preparations, there were 165 features,
173 which were used for training. 41 features were from the EMR or calculated based on EMR

174 data, like days since first injection. 124 features are based on OCT measurement data. Table
175 1 (available online only) gives an overview of all features used for training.

176 Individual patient eye histories were now built from the dataset created to date.
177 These histories consisted of a specified number of previous temporal consecutive visual
178 events up to a certain time point. It was essentially a condensed version of the past visits of
179 the patient until the third injection was given, by using the resulting dataset, as traditional
180 machine learning algorithms cannot use temporal information directly.

181 [Algorithms used for Visual Acuity Prediction](#)

182 To predict the VA outcome of nAMD patients, we applied five different machine learning
183 algorithms, all of which are freely available online and form part of the previously mentioned
184 libraries. In detail, they were:

- 185 • AdaBoost.R2 ²⁸: This belongs to the so-called boosting methods, which try to reduce
186 the bias of combined estimators by specially built base models (often weak learners).
187 The basic idea of the popular ensemble algorithm AdaBoost is to train an initial weak
188 learner (in our case a simple decision tree) on the original dataset. Afterwards,
189 multiple copies (49 in our case) are trained on the same dataset, but with more focus
190 on examples where the performance was previously poor (i.e., the ones that were
191 difficult to learn). The final prediction output is created by a majority vote of all weak
192 learners.
- 193 • Gradient Boosting ²⁹: This approach, being the second boosting method, is a
194 generalization of boosting to arbitrary differentiable loss functions. At every
195 optimization step, a regression tree is fit to the negative gradient of the loss function.
196 These models are very popular, powerful, and robust against outliers.
- 197 • Random Forests ³⁰: This belongs to the so-called averaging methods, where the basic
198 idea is to build multiple independent models and to average their predictions to
199 retrieve the final prediction. Under the assumption that the errors made by the
200 individual models are also independent, the averaging of their predictions will likely
201 reduce the error. In effect, Random Forests are an ensemble of simple decision trees
202 that are built from a bootstrap sample (i.e., a sample drawn with replacement). In
203 addition, each of the (100) decision trees is trained on a random subset of all
204 available features. Despite this being a very simple model, it has been shown to be
205 extremely powerful in many practical applications.

- 206 • Extremely Randomized Trees ³¹: These are special versions of Random Forests, but
207 instead of choosing the thresholds for the randomly picked features by how
208 discriminative they are, these thresholds are also selected at random. This may
209 enhance the variance of the ensemble at the price of higher bias.
- 210 • Lasso ³²: A linear model trained with L1 prior as a regularizer and coordinate descent
211 for training. It estimates sparse model coefficients (in contrast to L2 prior), i.e., it
212 leads to compact models, where zero is assigned to most of the low gain features.
213 Therefore, it is often used for feature selection.

214 Evaluation of Models

215 To evaluate the performance of our predictive models, we used 10-fold cross validation. We
216 randomly split the available GEIDs (\mathcal{D}) into 10 approximately equally sized, mutually
217 exclusive subsets ($\mathcal{D}_{10}, \mathcal{D}_{11}, \dots, \mathcal{D}_{19}$). In every iteration i of the 10 total iterations, we used
218 all subsets without the i -th ($\mathcal{D} \setminus \mathcal{D}_{1i}$) for training (or validation) and the i -th for testing.
219 Therefore, the tests of the model were performed on patient eyes that the models did not
220 previously see. For evaluation, either the data (e.g. VA) of the last one, two, three or four
221 visits prior to the third injection were available. Resulting accuracy was stratified by the
222 number of previous visits. The computing time for the training and testing of the models was
223 about 8 hours. To make the test results more comparable, we used fixed but initially
224 randomly generated seeds for shuffling the GEIDs before splitting. To provide a quality
225 measure for our predictions per algorithm, which shows how close they are to the ground

226 truth, we indicate the root of the mean squared error (RMSE= $\sqrt{\frac{1}{N} \sum_{i=1}^N (\tilde{y}_i - y_i)^2}$ for

227 N=number of predictions per fold) and the mean absolute error (MAE= $\frac{1}{N} \sum_{i=1}^N |\tilde{y}_i - y_i|$) for 90

228 and 365 days' prediction as compared with the ground truth, which is the actual patient VA
229 as measured in clinics. When describing accuracy of machine learning algorithms, usually
230 both values are indicated. RMSE does penalize outliers more, thereby allowing you to
231 choose the more robust algorithm. This is especially helpful, if the MAE is comparable
232 between tested algorithms. MAEs are usually used in medical publications only.

233 Results

234 The demographics of all patients/eyes included in the study are summarized in Table 2. For
235 the long-term prediction, out of 456 patients, 270 were female (59.2%). For the shorter
236 prediction period, out of 653 patients, 379 were female (58%). The average of patients was
237 slightly above 74 years in both groups. The VA at first injection was 0.54 (+/- 0.39) logMAR
238 for the long-term prediction cohort and 0.56 (+/- 0.42) logMAR for the short-term group.
239 Tables 3 and 4 show the accuracy of the applied machine learning algorithms used to predict
240 VA (in logMAR scale) as compared with the ground truth (i.e., the outcome as recorded
241 during the patient's visit in our clinic) after 90 days and 365 days, respectively. In each table,
242 the different numbers of previous visits, e.g., condensed histories of the patient's eye with
243 varying complexity, were used to predict the VA after 90 or 365 days. Predicted values were
244 compared with the closest VA measurement available. The best learner in all cases was the
245 L1 regularized linear model Lasso. If only data of one previous visit before the third injection
246 was taken into account for validation, the RMSE of VA over the ground truth was 0.14
247 logMAR (equals 7 letters) for the short-term prediction (90 days) and 0.23 logMAR (equals
248 11.5 letters) for the long-term prediction. The MAE of VA over the ground truth was 0.11
249 logMAR (5.5 letters) for the short-term prediction and 0.16 logMAR (8 letters) for the long-
250 term prediction. The RMSE was lowest (0.2 logMAR – 10 letters), if data of 4 previous visits
251 before the third injection were taken into account. For the MAE there was no improvement,
252 if 4 previous visits were included in validation. Figures 2 and 3 show the difference in visual
253 acuity between prediction and the ground truth, which are the VA values captured in our
254 clinic. Figure 4 shows the weights of features for the visual acuity prediction for 90 days
255 (Figure 4a) and for one year (Figure 4b). Figure 5 shows the change of VA between the true
256 three-month baseline and predicted VA after 3 months and one year.

257 Discussion

258 Several approaches have been made to predict progression in patients with AMD. Some of
259 them use genetic information to predict progression to an advanced form of wet AMD³³.
260 Others are based on imaging information from OCT and assess, for example, the presence
261 and size of drusen^{10,34,35}. Based on the familiar Comparison of AMD Treatments Trials
262 (CATT) trial, two studies evaluated predictors for vision outcomes after one respectively two
263 years of treatment^{36,37}. Several rules of thumb could be identified: For example, older age,

264 better baseline VA and greater total fovea thickness, were independently associated with
265 less improvement in VA after at least one year of therapy. These predictors are applicable in
266 many patients, but it can't be used in individual patients to predict his or her vision outcome
267 after a certain time. In this study, we have been able to predict VA in individual patients with
268 neovascular AMD undergoing anti-VEGF therapy after a further 90 or 365 days by using
269 machine learning based on real-life data from an EMR and OCT measurement data, but no
270 OCT images. Figure 4 shows that also in our study, previous VA values have great influence
271 on our prediction model. We also noted (figure 5), that predicted VA values, both short-term
272 as long-term, were different from the ground-truth at the 3-month baseline suggesting that
273 after month 3 changes in regard to VA during active therapy can be observed.

274 As expected, VA was best predicted for the short-term goal of 90 days. The best machine
275 learning algorithm was able to predict this with a RMSE of 7 letters (+/- 1.5 letters) or a MAE
276 of 5.5 letters (+/- 1 letter). The prediction was within the variability of measurements of VA
277 in a large eye clinic in which a real change can be detected if it is greater than 0.15 logMAR
278 ³⁸. The best VA prediction (RMSE: 0.20 logMAR (+/- 0.08), MAE: 0.16 logMAR (+/- 0.04)) for
279 365 days after further therapy was also close to this value of 0.15 logMAR. As mentioned
280 before, the RMSE is more sensitive to outliers, as larger errors are punished more severely.
281 This can be also observed in Figures 2 and 3. One would assume that the prediction is better
282 the more previous encounters are included, but for four previous visits, this was not
283 applicable. One reason might be that too few patients were included in the testing.

284 VA is, in most medical retina clinical trials, the primary endpoint, although it has not only
285 scientific value, but it gives an indication as to how patients are affected in their daily lives.
286 As is well known, VA is closely correlated with patient quality of life ²². A recently published
287 review article analyzing 14 studies that assess the psychological impact of anti-VEGF
288 treatments of wet AMD showed that patient often feared persisting vision loss and inability
289 to lead an independent life²⁴. A reduction in VA also leads to a reduction in quality of life and
290 the ability to perform daily duties. One of the most precious activity of daily life for many is
291 the ability to drive a car, which is often a necessity for leading an independent life in more
292 rural areas. Approximately 90% of patients have been shown to meet the requirements to
293 drive a car after 24 months of intravitreal anti-VEGF therapy ³⁹. The prediction of their VA
294 outcome after a defined time alleviates the anxieties of patients and motivated them to
295 pursue Anti-VEGF therapy, which requires frequent consultations with an unknown number

296 of planned injections. On the other hand, mental support for patients with an unfavorable
297 prognosis needs also to be provided. This helps to reduce the incidence of depression, if
298 initiated at a suitable time ⁴⁰.

299 Further refinement can be achieved by the use of patient genetic data. Genetic factors
300 are well known as being involved in disease progression and are more precise in conjunction
301 with further data. It has led to the creation of the “AMD risk calculator”, which can be
302 accessed online and for which various entries need to be made (e.g., age, sex, smoking
303 status, body dimensions, and different AMD gene mutations) ⁴¹. A combination of these
304 approaches might help to build a robust decision support tool for AMD, which might help to
305 deliver evidence-based personalized treatment for wet AMD patients.

306 Major information about the disease state and reaction to therapy is included in OCT
307 images. In this study, we have used numerical measurement information such as CRT from
308 the vendor’s software. However, this provides limited information concerning anatomical
309 aspects of the retina during the examination. Future approaches could make use of the OCT
310 scans itself and learn directly from the image by using deep learning. The use of this more
311 advanced technology could provide major performance gains. Initial promising studies
312 involving the application of deep learning to OCT scans have been published. For example,
313 use of deep learning technology has made it possible to classify normal versus age-related
314 macular degeneration OCT scans⁹. In another study, drusen progression over time at the
315 level of a single druse could be predicted over a mean follow-up of 37.8 months, as drusen
316 volume is known to be predictor for progression to late AMD¹⁰. Further innovations in this
317 field are expected and will likely contribute to more precise VA forecasts.

318 Interestingly, from a computer science perspective, a simple linear model such as Lasso
319 outperforms complex ensemble-based approaches such as Gradient Boosting Machines
320 (GBM), which usually achieve better results. Moreover, bagging with 10 estimators does not
321 really help to increase the performance indicating that the bagged estimators are not able to
322 learn various aspects of the data and make independent errors. Even training with Lasso’s
323 selected features (all non-zero model variables) did not enhance other learners. A linear
324 model thus seems to grasp the internal relationship in the given data relatively well. In this
325 case, the greater capacity of the other learners seems to lead to slight overfitting.

326 However, more work is necessary to improve real-life data preparation and cleaning, as
327 data input errors occur in daily routine work. An approach with regard to this limitation

328 might be the creation of national data registries, as more data should improve prediction
329 quality ⁴². The presented model relied on data of previous visits before VA prediction after
330 the third injection could be done. This limitation might be overcome in the future, if more
331 clinical data will be available and also OCT image recognition using deep learning is
332 integrated in a model. This would enable instant VA prediction on day of diagnosis or e.g. in
333 second opinion cases. Also, randomized clinical trials must be set up to test the reliability
334 and accuracy of decision support software.

335 In conclusion, the prediction of VA by using real-life clinical data is a valuable first
336 step to developing a clinical decision support software. It demonstrates the value of well-
337 structured EMRs. It can be used to optimize anti-VEGF treatment in neovascular AMD
338 patients and the allocation of resources to those who need it most.

339 Figure legends

340 *Figure 1: This figure shows nine sectors, which are placed over the macula within the Heidelberg Eye Explorer software, CO*
341 *being the fovea. Data measurements in the XML files were given for the whole grid as well as for each individual sector.*
342 *Therefore, features of an individual sector in Table 1 were count nine times. The letters T (temporal), I (inferior), N (nasal)*
343 *and S (superior) indicate the anatomic position of the sector.*

344 *Figure 2: Plots showing differences between visual acuity prediction (red line) and ground truth (green line) for a prediction*
345 *for 90 further days of treatment. On the x-axis, a consecutive number for each patient eye in the test set is shown. Positive*
346 *values on y-axis mean improvement of visual acuity, negative values mean deterioration. A graph is plotted for every*
347 *prediction depending on previously included visits (see headline for the exact number). VA=visual acuity (logMAR).*

348 *Figure 3: Plots showing differences between visual acuity prediction (red line) and ground truth (green line) for a prediction*
349 *for 365 further days of treatment. On the x-axis, a consecutive number for each patient eye in the test set is shown. Positive*
350 *values on y-axis mean improvement of visual acuity, negative values mean deterioration. A graph is plotted for every*
351 *prediction depending on previously included visits (see headline for the exact number). VA=visual acuity (logMAR).*

352 *Figure 4: The plot shows the weight of the different features for the visual acuity prediction task over 3 months (4a) and 1*
353 *year (4b). The light blue bar indicates how important the feature was on average for the model on the different test runs (10*
354 *in total). The dark blue bar gives an intuition of how instable the feature was in the different test runs in terms of an*
355 *importance vs. variance ratio. The more of the light blue bar is overlayed with the dark blue bar, the more instable the*
356 *feature was during the test runs. T2=Outer temporal measurement section according to the ETDRS grid as placed on the*
357 *macula, E11.30=ICD-10 code for diabetic type 2 eye complications, I2=outer inferior section of the ETDRS grid, VPP=volume*
358 *in a grid sector of the ETDRS grid, CO=central location of the ETDRS grid.*

359 *Figure 5: This figures shows the difference between true three-month visual acuity and predicted visual acuity after 90 days*
360 *(figure 5b)/365 days (figure 5b). All values are given in logMAR units. The vertical axis indicates the relative frequency of*
361 *each VA delta value.*

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