

CORRIGENDUM

Genetic and environmental factors strongly influence risk, severity and progression of age-related macular degeneration

Wenqiu Wang, Katarzyna Gawlik, Joe Lopez, Cindy Wen, Jie Zhu, Frances Wu, William Shi, Samuel Scheibler, Huimin Cai, Ram Vairavan, Alexander Shi, Weldon Haw, Henry Ferreyra, Ming Zhang, Sherman Chang and Kang Zhang

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Since the publication of this Research Article, the authors have noticed the Legends of Figures 3–5 were incorrect and they should be corrected as follows:

Figure 3. Performance of the prediction model for 7-year progression to geographic atrophy (GA) based on 10 genetic variables. **(a)** Receiver operating characteristic curve (ROC) for the 10-SNP model was generated for learning (blue line) and testing (red dashed line) sets by using the binary logistic regression analysis with a 10-fold cross-validation method. **(b)** Prediction success parameters were calculated for testing set. **(c)** Variable importance ranking is showing relative scores generated by TreeNet software and positioning the predictors from the most important to the least important. NPV, negative prediction value; PPV, positive prediction value.

Figure 4. Performance of the prediction model for 7-year progression to choroidal neovascularization (CNV) based on 27 variables (genetic and environmental factors). **(a)** Receiver operating characteristic curve (ROC) for the 25-SNP plus smoking and BMI model was generated for learning (blue line) and testing (red dashed line) sets by using the binary logistic regression analysis with a 10-fold cross-validation method. **(b)** Prediction success parameters were calculated for testing set. **(c)** Variable importance ranking is showing relative scores generated by TreeNet software and positioning the predictors from the most important to the least important. NPV, negative prediction value; PPV, positive prediction value.

Figure 5. Distribution of the risk score computed in the prediction models. **(a)** AMD lifetime risk score distribution in the study population. AMD cases are shown in red and controls in blue. **(b)** GA progression risk score distribution in the study population. GA progressors are shown in brown and non-progressors in green. **(c)** CNV progression risk score distribution in the study population. CNV progressors are shown in purple and non-progressors in green.

The online version of the article has not been corrected.