Supplementary Online Content


**eTable.** Five-year risk of developing advanced AMD in predictive models with and without genetics (*CFH Y402H* and *ARMS2 A69S* genotypes) for a 75-year-old individual.

This supplementary material has been provided by the authors to give readers additional information about their work.
eTable 1. Five-Year Risk (%) of Developing Advanced AMD in Predictive Models With and Without Genetics (CFH Y402H and ARMS2 A69S Genotypes) for a 75-Year-Old Individual

<table>
<thead>
<tr>
<th>Positive Baseline Variables</th>
<th>AREDS Simple Scale Score</th>
<th>Model without Genetics* (Phenotype**, Demographic and Environmental Factors***)</th>
<th>Model with Genetics** (Phenotype**, Demographic and Environmental Factors***, Genetics*)</th>
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<tbody>
<tr>
<td></td>
<td></td>
<td>NN/NN</td>
<td>NR/NR</td>
</tr>
<tr>
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<td>1 (Positive family history)</td>
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<tr>
<td>2 (Current smoker and advanced AMD in 1 eye)</td>
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<td>3 (Positive family history, current smoker, and advanced AMD in 1 eye)</td>
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</table>

*Genetics: NN/NN = nonrisk genotype: CFH TT/ARMS2 GG; NR/NR = heterozygous genotype: CFH CT/ARMS2 GT; RR/RR = risk genotype: CFH CC/ARMS2 TT.

**Phenotype: AREDS Simple scale score; presence of very large drusen (250 microns or greater in diameter) in at least 1 eye; presence of advanced AMD in 1 eye (geographic atrophy or neovascular AMD).

***Baseline demographic/environmental factors: age (years); positive family history of AMD (first-degree relative); current smoker.