

Supplementary Table S2. Full risk model for progression to advanced age-related macular degeneration (AMD), geographic atrophy (GA) and neovascular disease (NV)

	Advanced AMD (948/5421 eyes)	P value	GA (487/5421 eyes)	P value	NV (495/5421 eyes)	P value
	ESTIMATES (\pm SE)*		ESTIMATES (\pm SE)*		ESTIMATES (\pm SE)*	
Demographic						
male	0.03 (\pm 0.08)	0.69	0.19 (\pm 0.11)	0.068	-0.15 (\pm 0.11)	0.15
white	0.68 (\pm 0.36)	0.06	1.99 (\pm 1.03)	0.053	0.17 (\pm 0.41)	0.67
> high school	-0.04 (\pm 0.07)	0.61	0.07 (\pm 0.11)	0.53	-0.15 (\pm 0.11)	0.17
Behavioral						
Current smoker	0.96 (\pm 0.13)	<0.001	0.74 (\pm 0.2)	0.0002	0.98 (\pm 0.19)	<0.001
Past smoker	0.14 (\pm 0.08)	0.088	0.06 (\pm 0.11)	0.61	0.28 (\pm 0.11)	0.011
BMI 25-29	0.15 (\pm 0.09)	0.078	0.14 (\pm 0.12)	0.26	0.19 (\pm 0.12)	0.14
BMI \geq 30	0.28 (\pm 0.1)	0.003	0.25 (\pm 0.14)	0.06	0.32 (\pm 0.14)	0.018
Baseline Severity Scale						
1						
2	1.46 (\pm 0.3)	<0.001	2.39 (\pm 0.65)	0.0002	1.08 (\pm 0.34)	0.002
3	1.77 (\pm 0.32)	<0.001	2.66 (\pm 0.62)	<0.001	1.46 (\pm 0.36)	<0.001
4	2.60 (\pm 0.25)	<0.001	3.29 (\pm 0.62)	<0.001	2.32 (\pm 0.29)	<0.001
5	3.26 (\pm 0.26)	<0.001	4.21 (\pm 0.6)	<0.001	2.88 (\pm 0.28)	<0.001
6	3.69 (\pm 0.25)	<0.001	4.70 (\pm 0.6)	<0.001	3.27 (\pm 0.28)	<0.001
7	4.31 (\pm 0.25)	<0.001	5.73 (\pm 0.59)	<0.001	3.27 (\pm 0.28)	<0.001
8	4.63 (\pm 0.25)	<0.001	6.14 (\pm 0.59)	<0.001	3.37 (\pm 0.3)	<0.001
Genetic variants						
Complement pathway						
CFH Y402H: rs1061170					0.27 (\pm 0.07)	<0.001
CFH: rs1410996	0.29 (\pm 0.07)	<0.001	0.29 (\pm 0.1)	0.003		
CFH R1210C: rs121913059	1.08 (\pm 0.26)	<0.001	0.91 (\pm 0.44)	0.037	1.04 (\pm 0.3)	<0.001

<i>C3</i> R102G: rs2230199	0.15 (± 0.06)	0.007		0.17 (± 0.08)	0.033
<i>C3</i> K155Q: rs147859257	0.70 (± 0.16)	<0.001	0.83 (± 0.19)	<0.001	
Angiogenesis pathway					
<i>TGFBR1</i> : rs334353				-0.18 (± 0.09)	0.044
Immune/inflammatory pathway					
<i>ARMS2/HTRA1</i> : rs10490924	0.36 (± 0.05)	<0.001	0.22 (± 0.07)	0.002	0.50 (± 0.07)
<i>TNFRSF10A</i> : rs13278062				-0.13 (± 0.07)	0.066
Extracellular matrix					
<i>COL8A1</i> : rs13095226	0.17 (± 0.08)	0.023	0.24 (± 0.11)	0.028	
DNA repair/protein binding					
<i>RAD51B</i> : rs8017304	-0.20 (± 0.06)	<0.001		-0.27 (± 0.08)	<0.001
<i>HSPH1/B3GALT1</i> : rs9542236	0.12 (± 0.05)	0.019		0.15 (± 0.07)	0.038

* Estimates were calculated for time to progression using age as the time scale with eyes as the unit of analysis, adjusted for sex, race, education, smoking history, BMI, AMD baseline eye-specific severity scale, AREDS treatment, and the genetic variants associated with the respective outcome.
