

Evidence from Case-control and Longitudinal Studies Supports Associations of Genetic Variation in *APOE*, *CETP* and *IL6* with Human Longevity

AGE

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Supplementary Table 2: Allele frequency case-control comparisons of middle-aged and oldest-old Danish individuals (both genders combined).

Gene	SNPs genotyped	p<0.05	p<0.01	p<0.005	p<0.001
<i>CETP</i>	20	6	0	1	1
<i>APOE</i>	3	2	0	1	0
<i>HSPA14</i>	6	4	1	2	0
<i>SIRT6</i>	3	1	0	1	0
<i>TGFB1</i>	7	2	0	0	0
11 additional genes	63	0	0	0	0
Observed	102	15	1	5	1
Expected		5.1	1.02	0.51	0.10