

Table S1. Minor allele frequency enrichment summary of nonsense variants in the Old Order Amish (n=1,725) compared to 1000g_(EUR) and ESP_(EA) populations.

1000g_(EUR)

	Ratio >1	Ratio <1	Ratio Unknown	Total
<i>P</i> < 0.05	67	79	0	146
<i>P</i> > 0.05	30	32	0	62
<i>P</i> N.C.	34	4	20	58
Total	131	115	20	266

ESP_(EA)

	Ratio >1	Ratio <1	Ratio Unknown	Total
<i>P</i> < 0.05	103	84	0	187
<i>P</i> > 0.05	22	23	0	45
<i>P</i> N.C.	6	8	20	34
Total	131	115	20	266

Abbreviations: 1000g_(EUR), Total European Ancestry population from 1000 Genomes; ESP_(EA), European American population from the National Heart, Lung, and Blood Institute (NHLBI) Exome Sequencing Project (ESP); Ratio, minor allele frequency in the Old Order Amish divided by the minor allele frequency in the relevant population; N.C., not calculated.