Low-frequency and rare variants may contribute to elucidate the genetics of major depressive disorder

Supplementary Information

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This supplementary information includes Table S1 and Figure S1.

Table S1. Breakdown of descriptive statistics for MDD cases and controls. Std Err, standard error.

Cohort	Status	Sex	Number	Age Mean	Age Std Err
	MDD	Male	70	40.96	1.45
Mexican-		Female	133	39.81	1.05
American	Control	Male	80	39.34	1.36
		Female	116	39.03	1.13
	MDD	Male	216	38.15	0.94
European-		Female	257	39.30	0.79
Ancestry	Control	Male	211	32.63	0.78
		Female	286	32.41	0.73

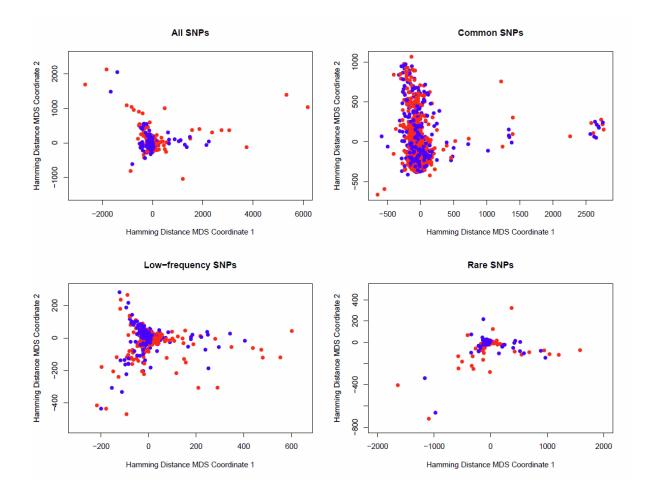


Figure S1. Multi-Dimensional Scaling (MDS) visualization based on Hamming distance for European-ancestry cohort without far outliers.