Table S3. Number of top predicted genes that ovrelap between different models at different cutoffs in the Mixed-Pop cross-validation scheme. KNN, K-Nearest-Neighbor; EN, Elastic-Net; SS, Single-SNP.

Top genes	KNN-EN	KNN-SS	EN-SS	All models
10	7	5	6	4
20	14	11	12	9
50	36	30	37	26
100	74	64	72	58
200	143	131	149	117