

Supplemental Figures

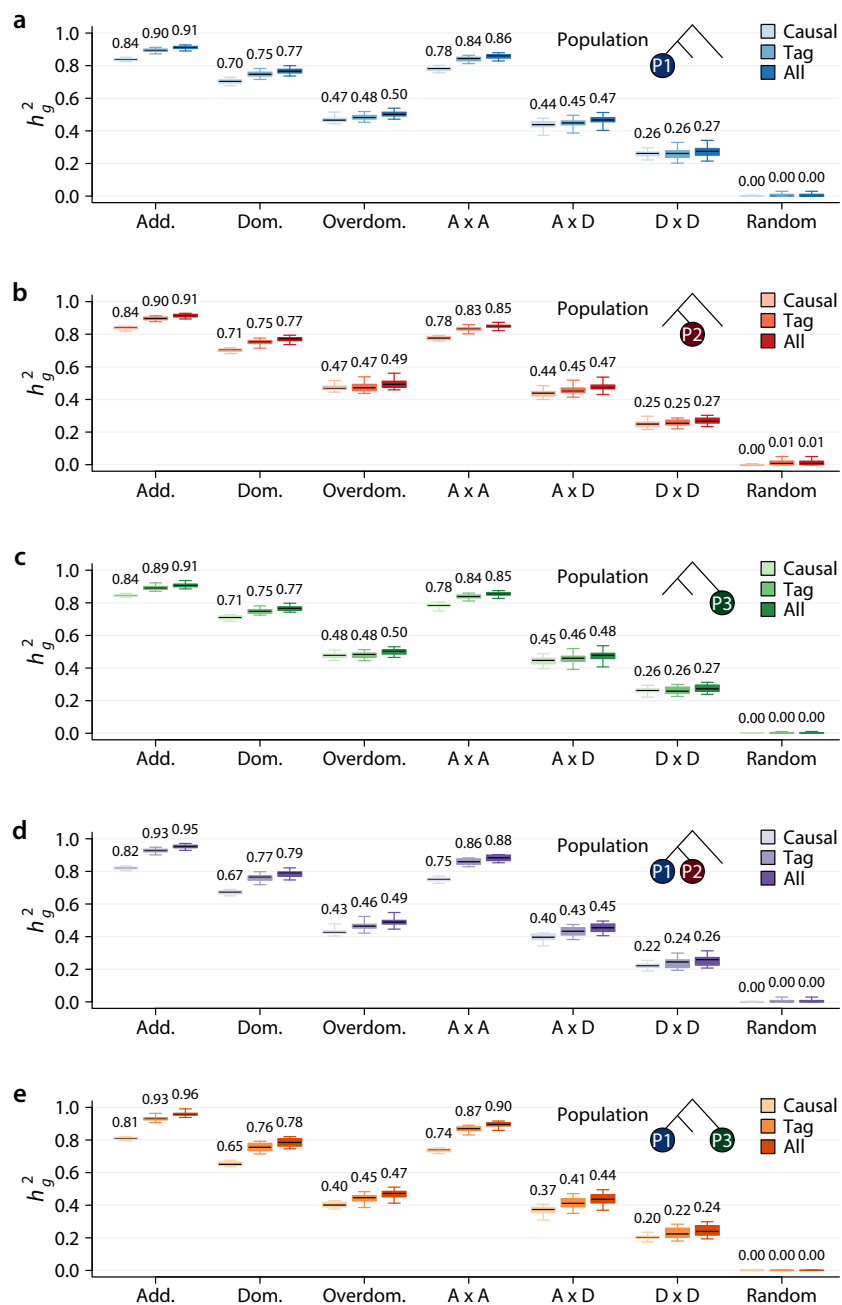


Figure S1. Genomic heritability in different simulated populations. Genomic heritability was plotted for different population samples. (a) population P1; (b) population P2; (c) population P3; (d) population P1 + P2; and (e) population P1 + P3.

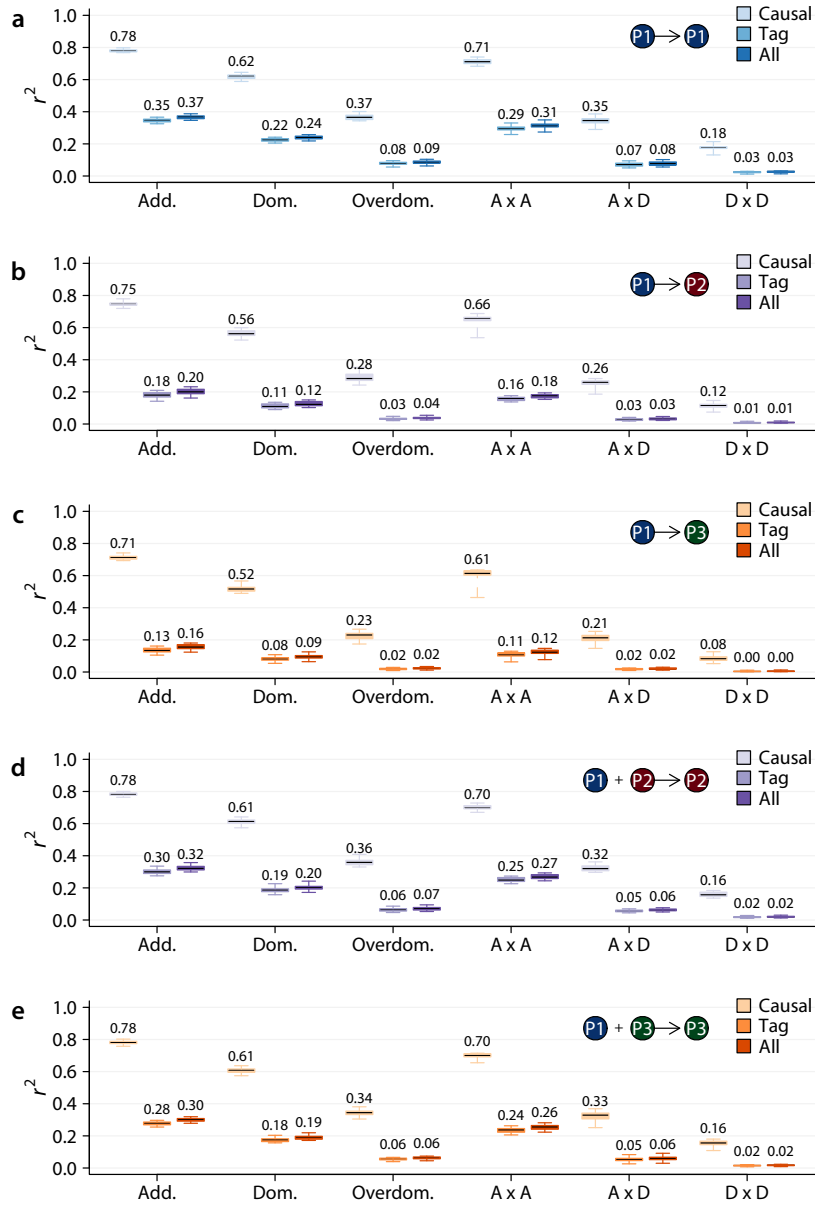


Figure S2. Polygenic prediction under different genetic architecture in different populations. Accuracies of polygenic prediction under different genetic architecture in (a) fit model in P1, predict in A; (b) fit in P1, predict in P2; (c) fit in P1, predict in P3; (d) fit in P1 + P2, predict in P2; (e) fit in P1 + P3, predict in P3.