

Supplementary Datafile 3: Analyses Related to Discussion and Table S3.1 show the Relationship of Lead SNPs from Meta-analysis to Rare CNVs and Mutations Previously Associated with Neurodevelopmental Genomic Disorders.

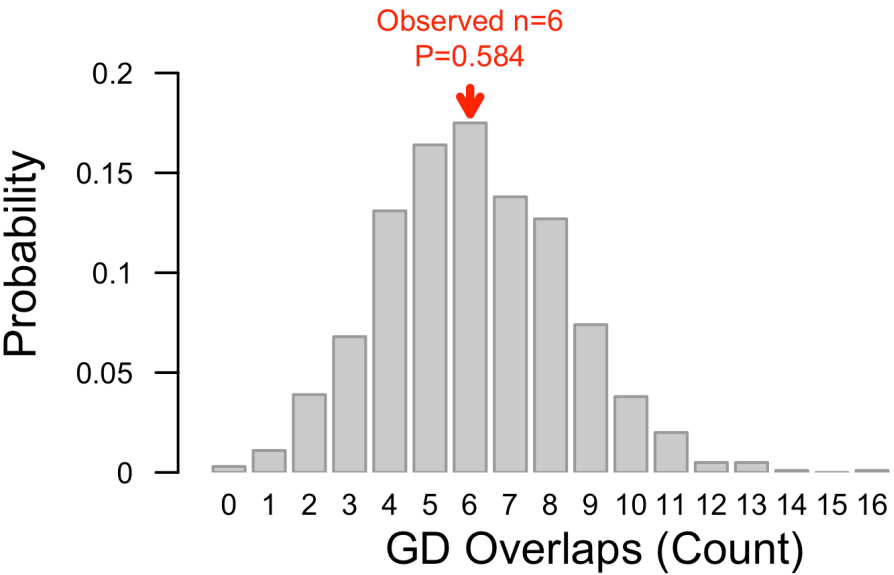
Data S3.1: Sentinel SNPs are not overrepresented among genomic disorder loci.

Data S3.2: Genes linked to sentinel SNPs are not overrepresented among dominant-acting ASD risk genes.

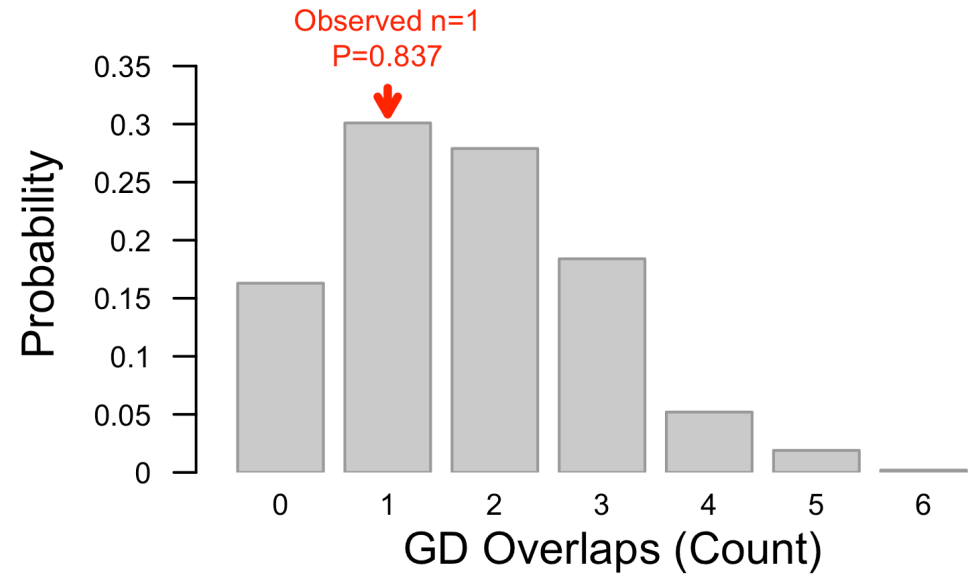
Data S3.3: Genes linked to sentinel SNPs are not enriched for damaging de novo mutations in ASD cases.

Fig SD3.1

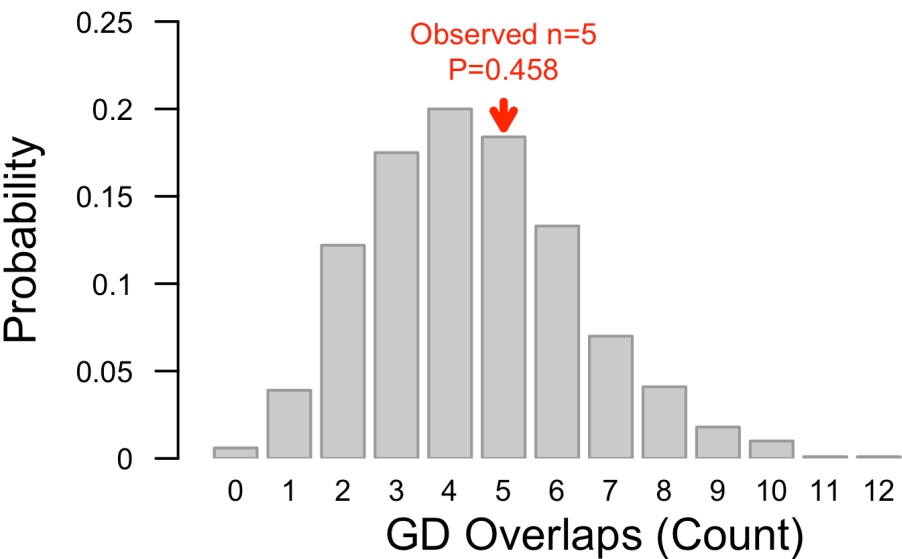
All SNPs



Disorder-Specific SNPs



Pleiotropic SNPs



Highly Pleiotropic SNPs

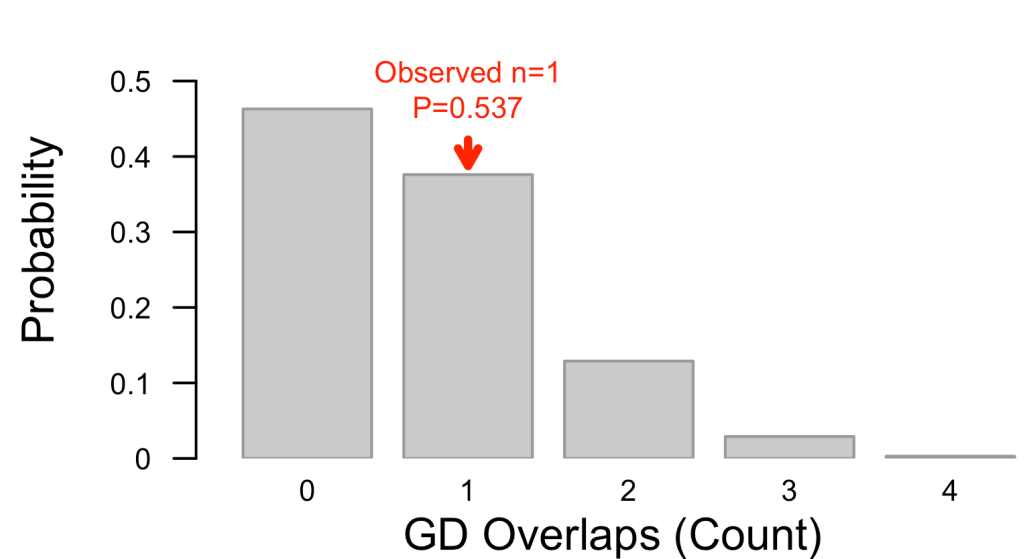


Fig SD3.2

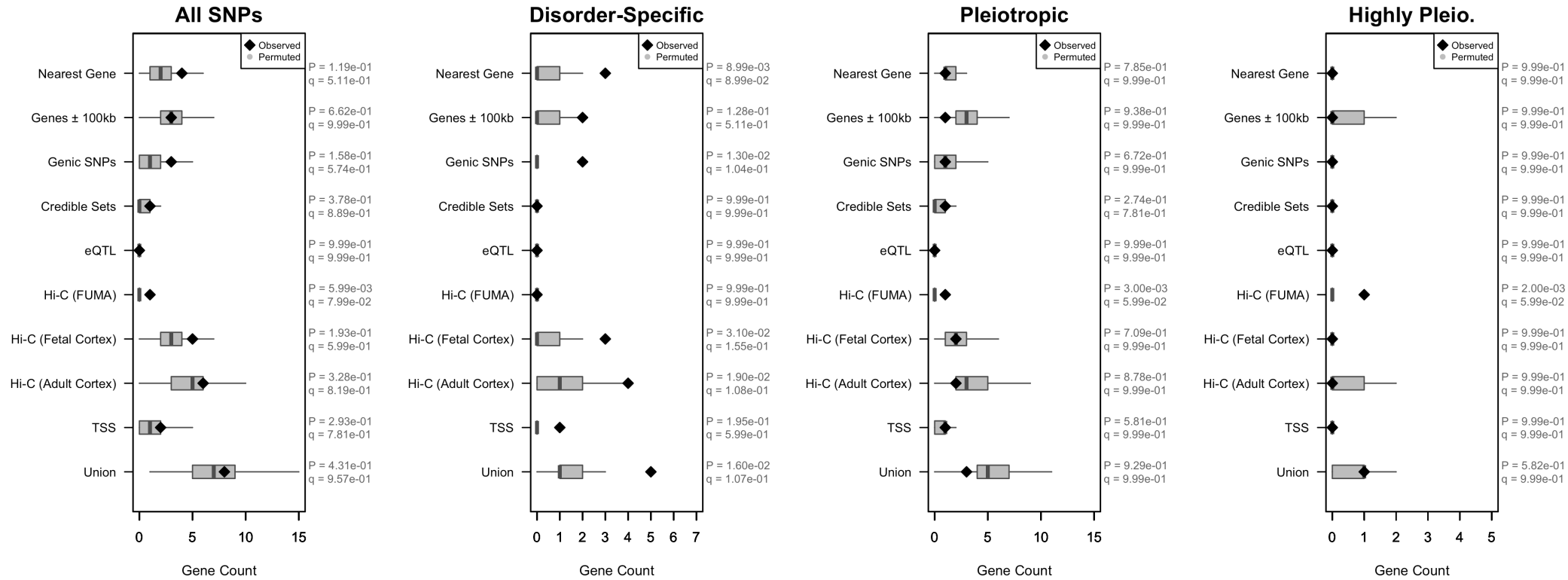


Fig SD3.3

