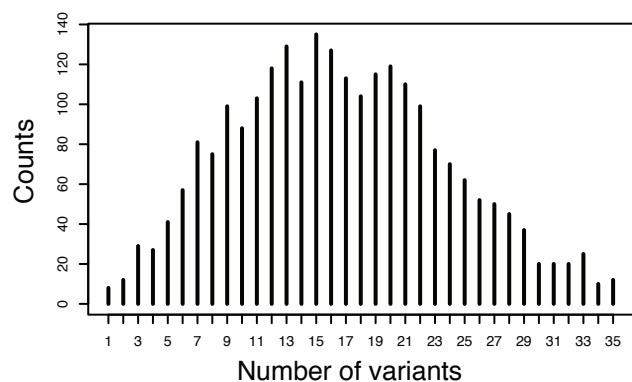
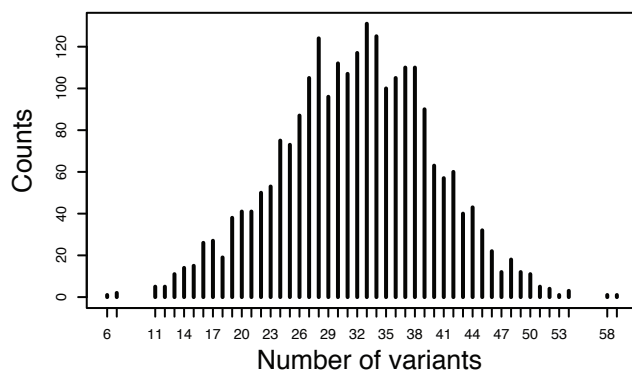


**S4 Figure: Distribution of number of causal variants and total number of simulated variants tested per locus under different architectures.**

**A** Number of variants with assigned effects



**B** Total number of variants tested



**C** Number of total segregating variants vs. number of causal variants under each architecture (VE=1%)

