S13 Figure: Power of gene-based methods in 3K vs. 10K samples, as a function of ratio of causal to total number of variants per locus.

*All loci simulated in both 3K and 10K samples are shown as grey points in both panels **A**) and **B**).*

Red points are loci simulated in 3K samples; blue points are loci simulated in 10K samples. In both sample sizes, SKAT-O p-values are more significant as the ratio of causal to total number of variants increases.