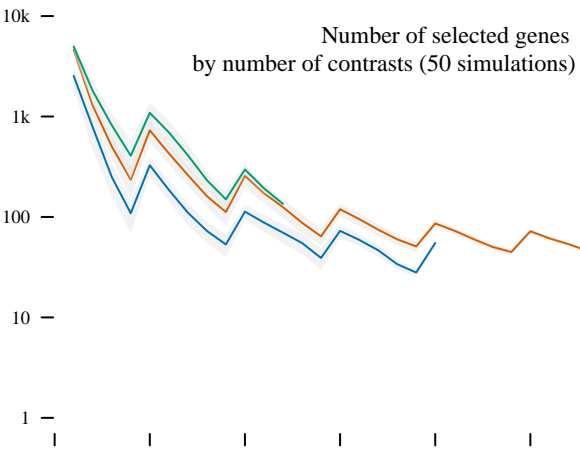


Number of selected genes
by number of contrasts (50 simulations)



Number of selected genes
by p-value cutoff

