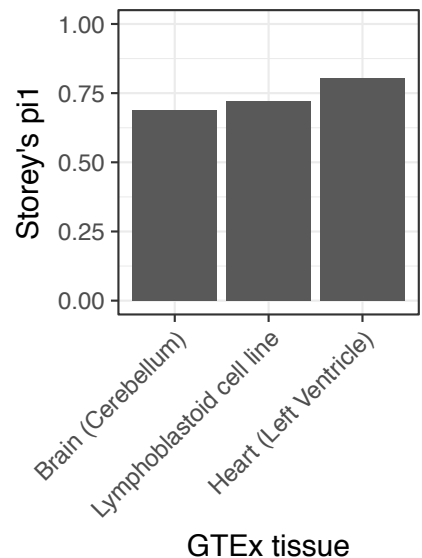
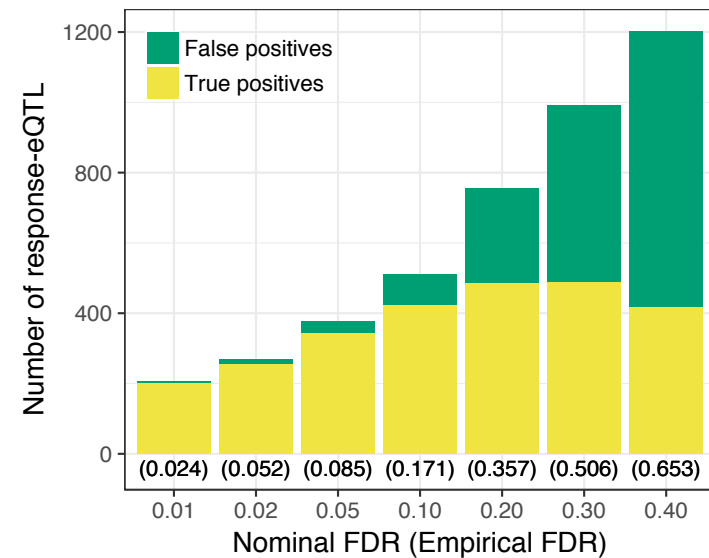


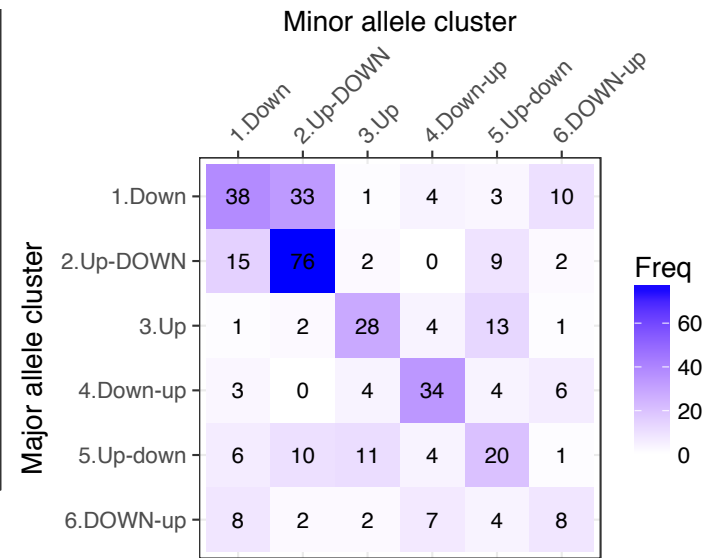
a Marginal eQTL replicate in independent heart data



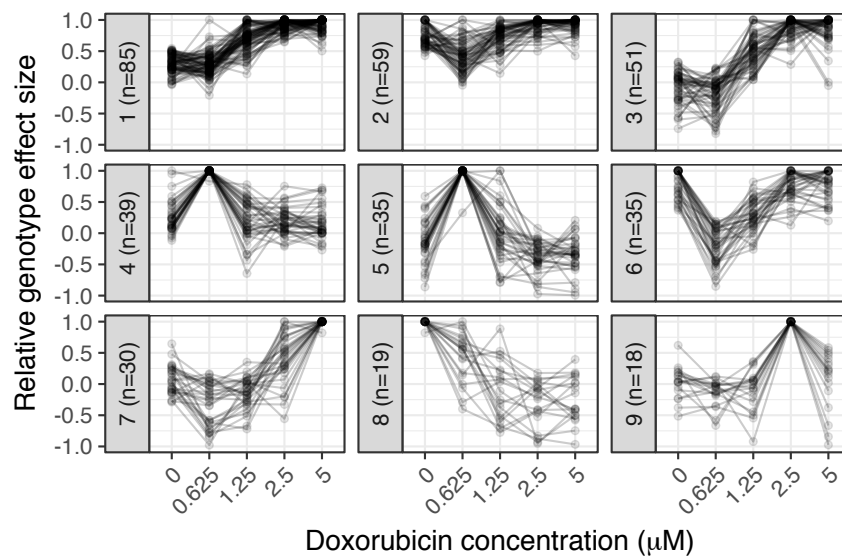
b Detection of 100s of variants modulating transcriptomic response in *cis*



c 46% of response-eQTL switch response pattern
54% modulate the same pattern



d Relative genotype dosage effect sizes form distinct patterns



e An example response-eQTL where minor-allele homozygotes show a markedly different response

