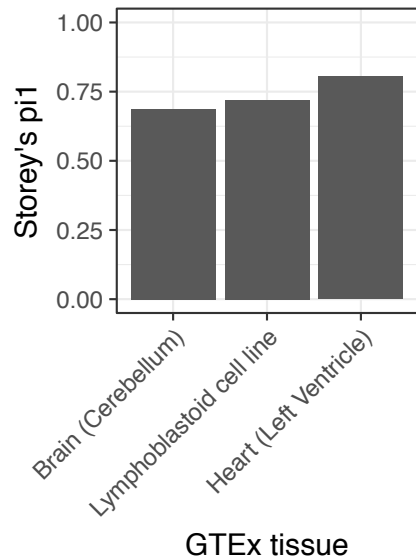
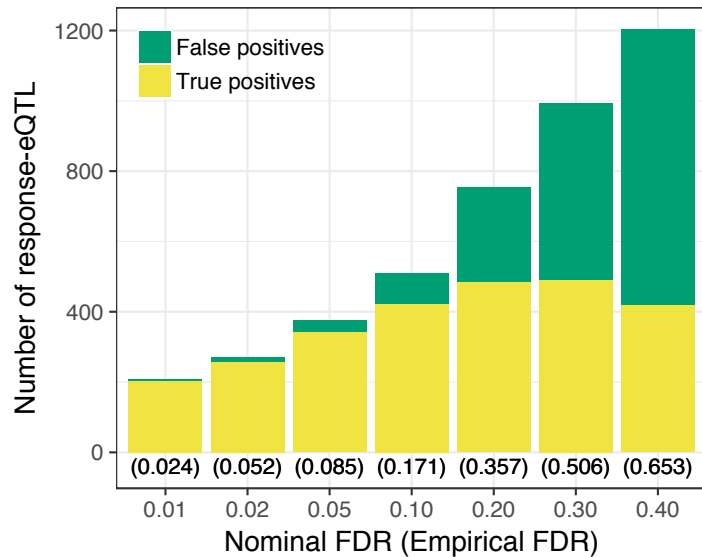


a Marginal eQTL replicate in independent heart data



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



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

