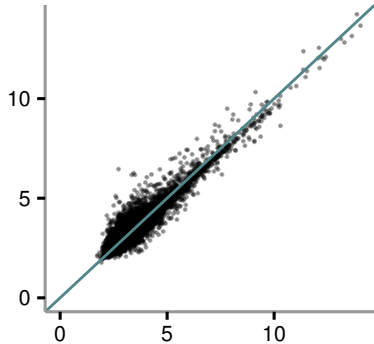
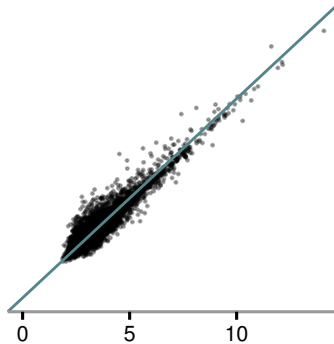


eQTL $-\log_{10}(\text{p-value})$
y = SNP + Batch + Transcriptome diversity

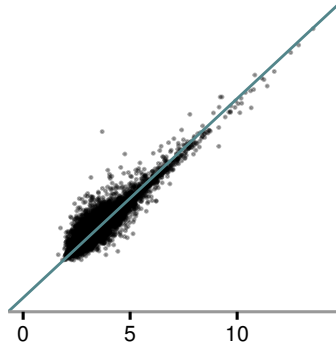
kidney



liver



lung



eQTL $-\log_{10}(\text{p-value})$
y = SNP + Batch