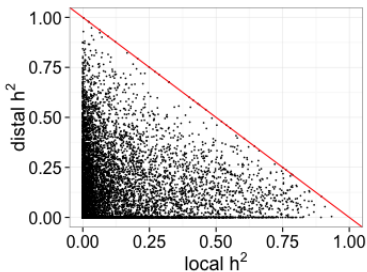


local = SNPs within 1Mb of gene
distal = SNPs on non-gene chrs



local = SNPs within 1Mb of gene
distal = known eQTLs on non-gene chrs

