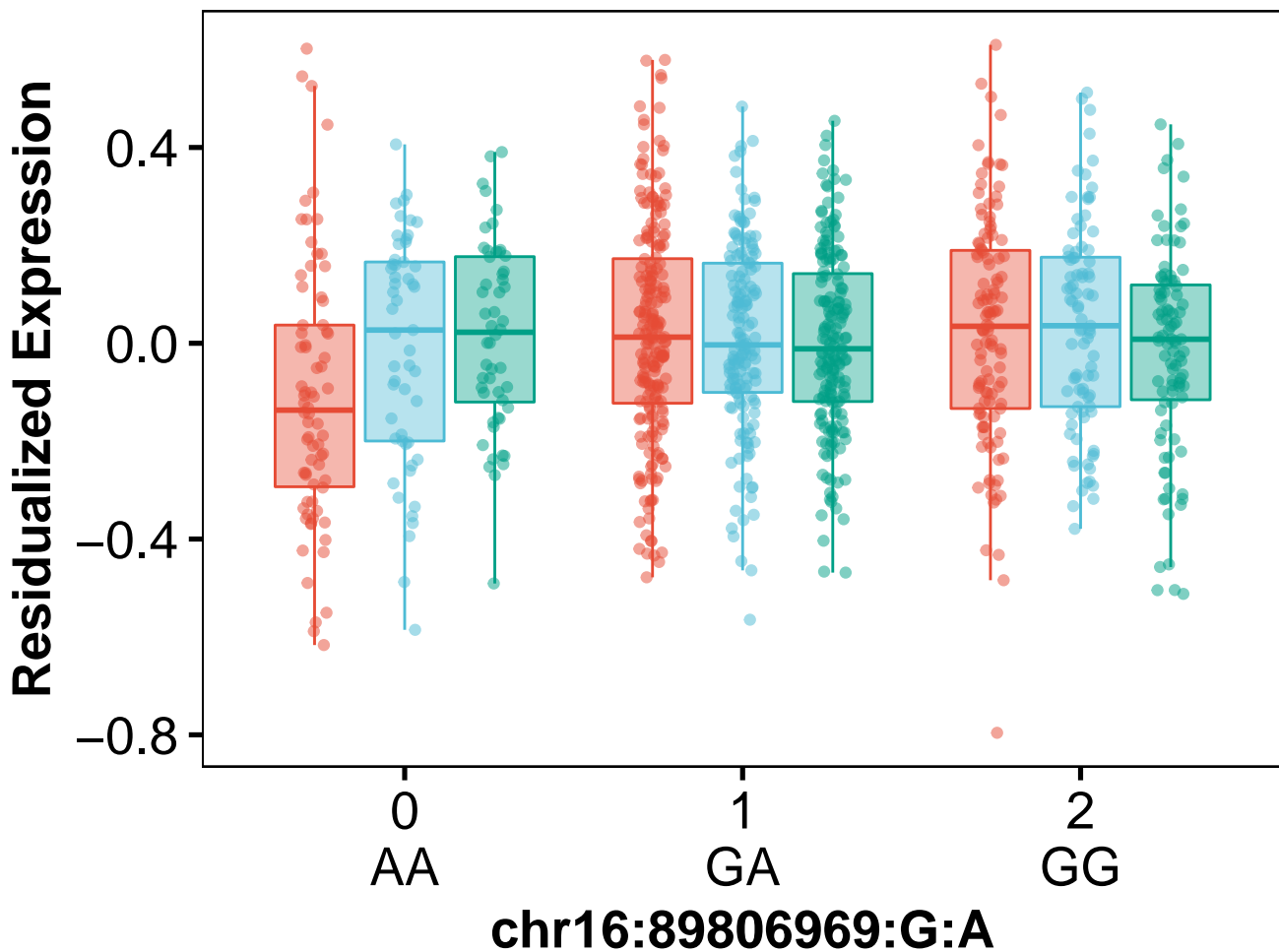


ENSG00000261574.1
eQTL lfsr < 0.025
SZ GWAS pvalue: 3.4e-09
SZ risk allele: G



Region  Caudate  DLPFC  HIPPO