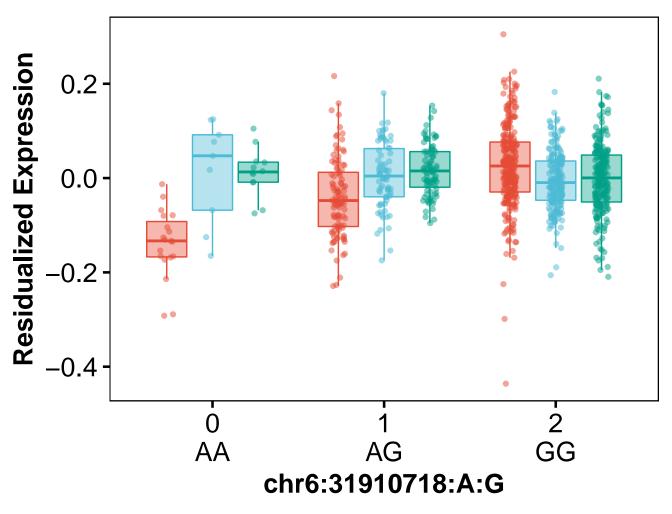
EHMT2
ENSG00000204371.11
eQTL q-value: 2.6e-20

SZ GWAS pvalue: 5.3e-15 SZ risk allele: G



Region = Caudate = DLPFC = HIPPO