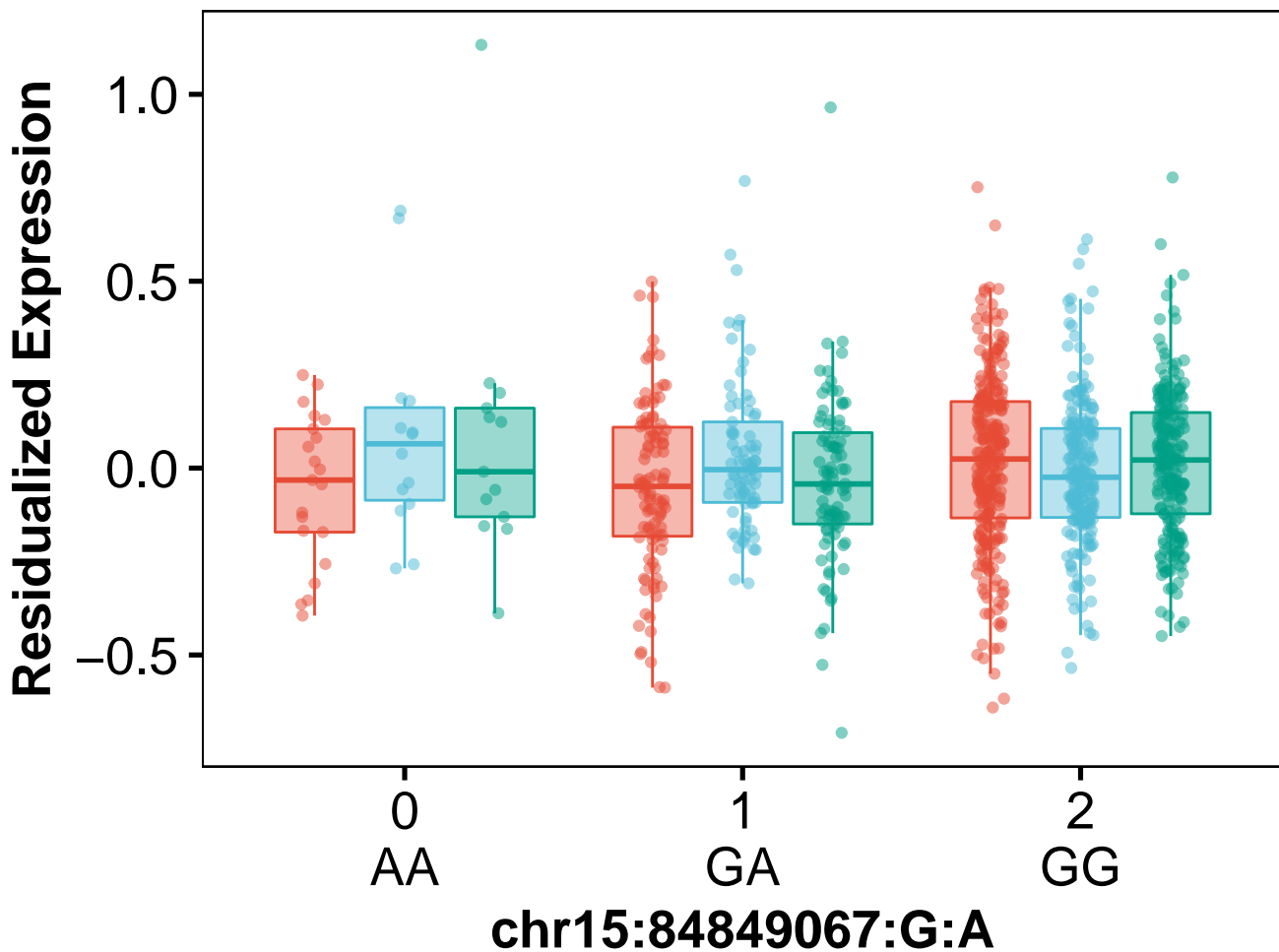


NMB
ENSG00000197696.9
eQTL lfsr < 0.014
SZ GWAS pvalue: 6.6e-10
SZ risk allele: G



Region  **Caudate**  **DLPFC**  **HIPPO**