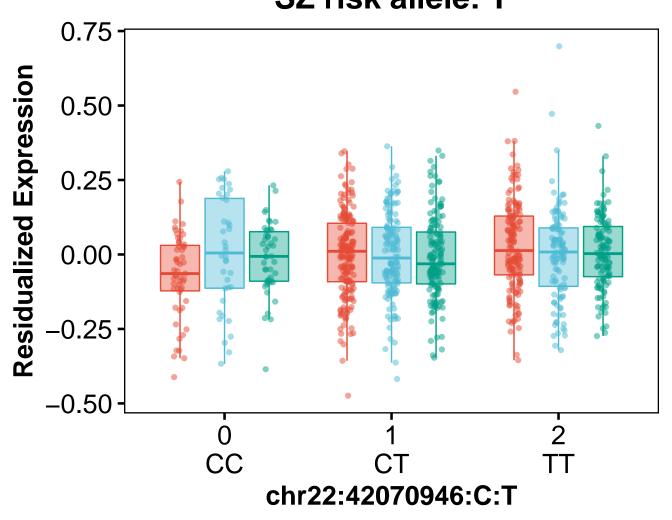
OLA1P1 ENSG00000213790.2 eQTL lfsr < 0.024 SZ GWAS pvalue: 2.1e-10 SZ risk allele: T



Region = Caudate = DLPFC = HIPPC