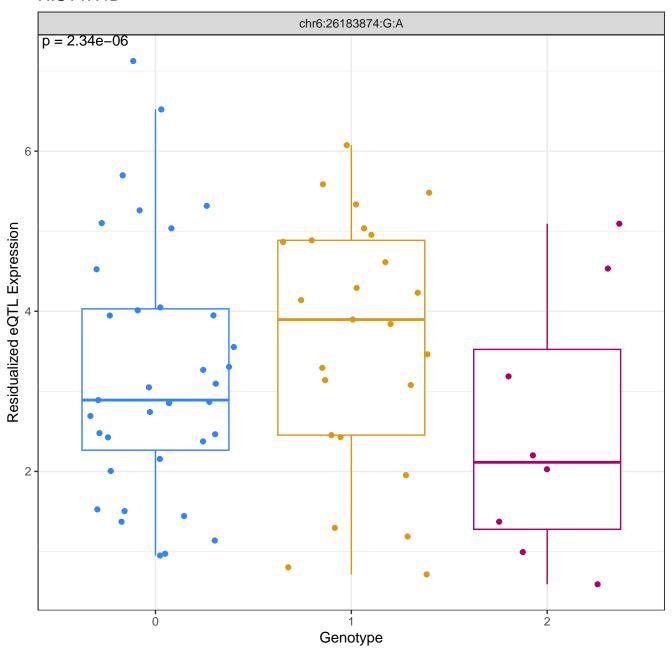
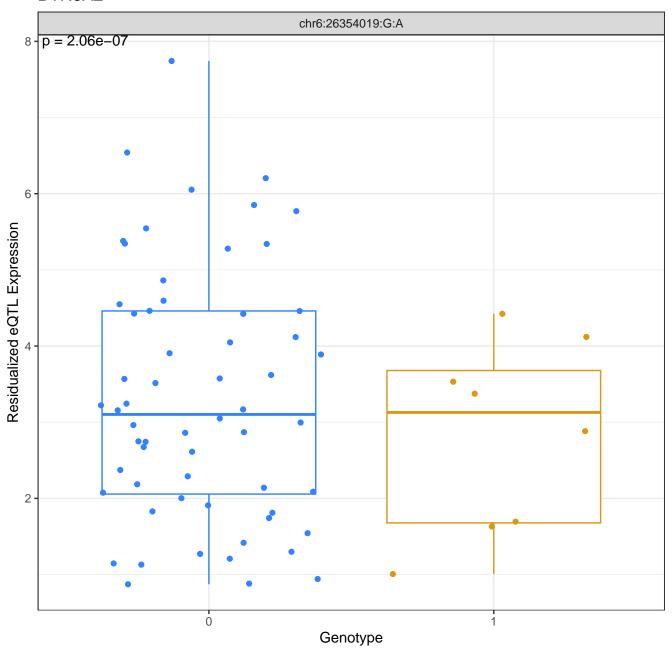
NEK4 chr3:52550699:A:G p = 1.21e - 0710.0 -Residualized eQTL Expression 7.5 -5.0 -2.5 -2 Ö i Genotype

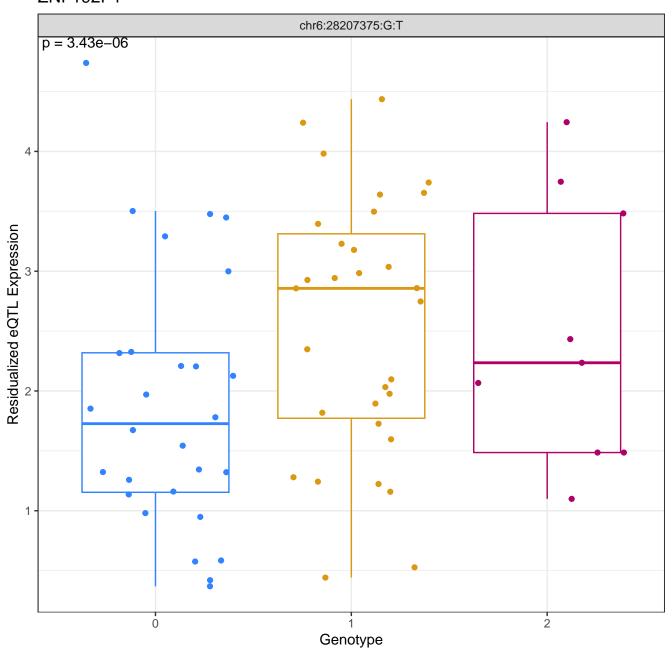
HIST1H4D



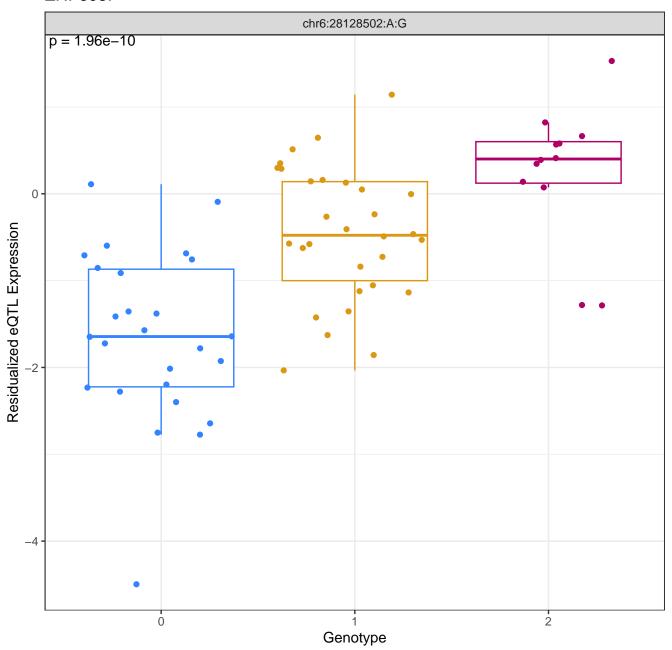
BTN3A2



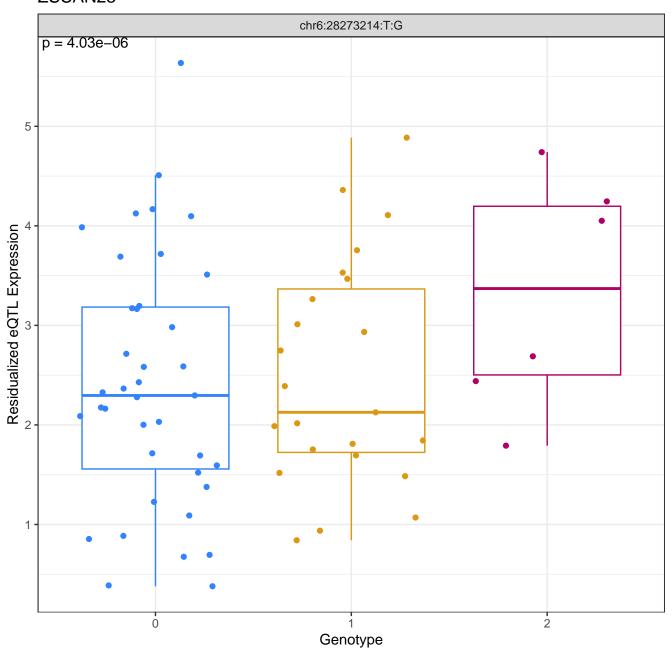
ZNF192P1

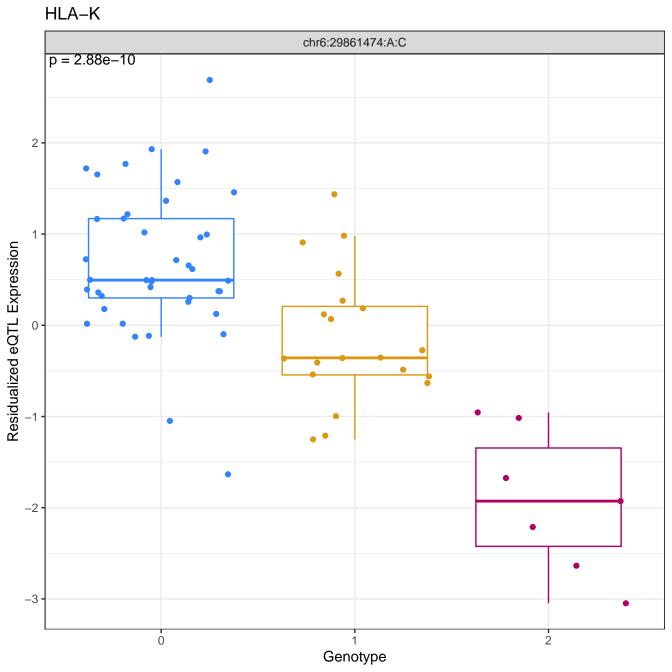


ZNF603P

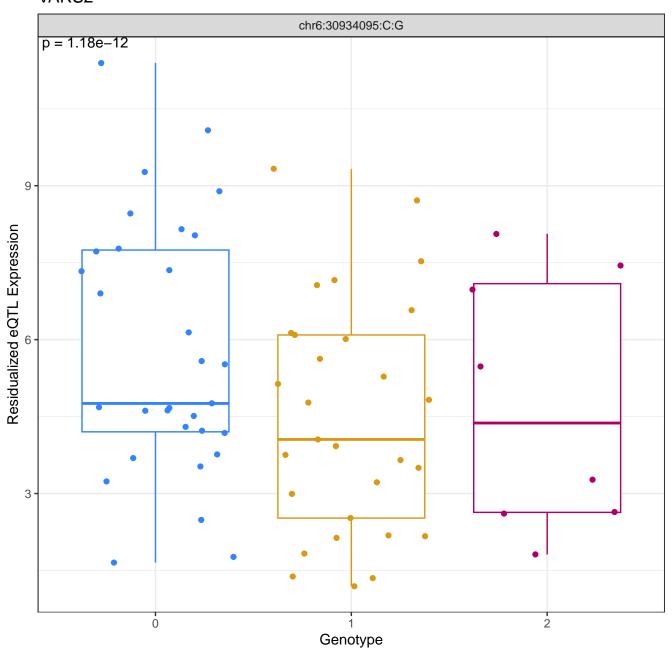


ZSCAN23





VARS2



PSORS1C1 chr6:31025536:C:T p = 5.37e - 080 -Residualized eQTL Expression **-4** -2 i Genotype Ö

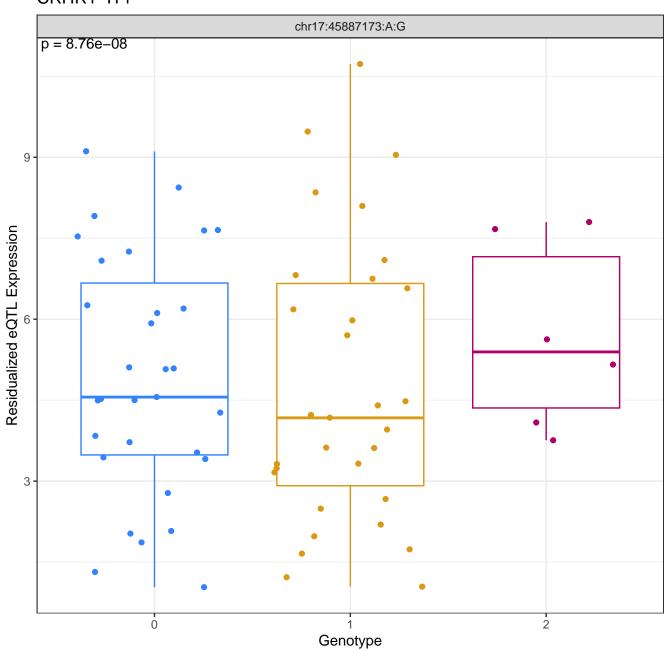
HLA-DMA chr6:32895181:C:G p = 6.71e-08 6 -Residualized eQTL Expression Ö Genotype

LRRC37A4P chr17:45701314:T:A p = 5.41e - 1610.0 -7.5 -Residualized eQTL Expression 5.0 2.5 -0.0 -2 Ö Genotype

DND1P1 chr17:45910839:A:G p = 3.41e-07 0 -Residualized eQTL Expression -3 **-**2 Ö i Genotype

RP11-707O23.5 chr17:45817181:C:T p = 7.44e-17 5. Residualized eQTL Expression 0 2 Ö i Genotype

CRHR1-IT1



KANSL1-AS1 chr17:45817181:C:T p = 6.32e-12 Residualized eQTL Expression -2 **-**-3 **-**2 i Genotype Ö

LRRC37A

