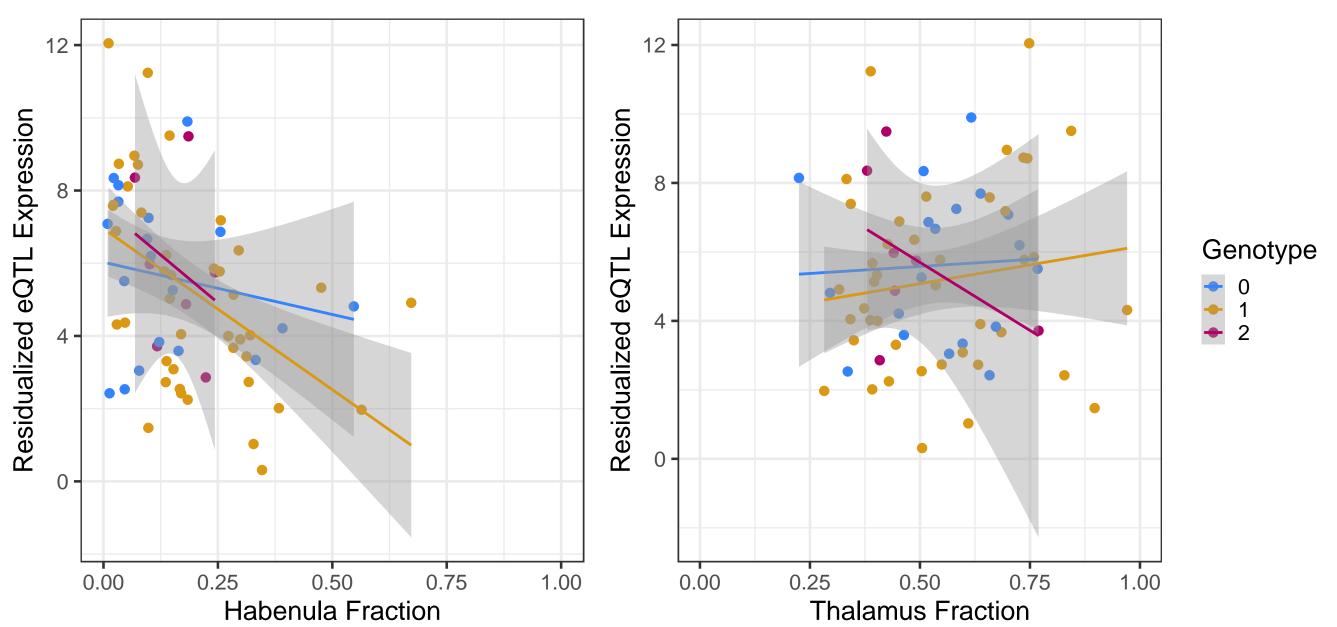
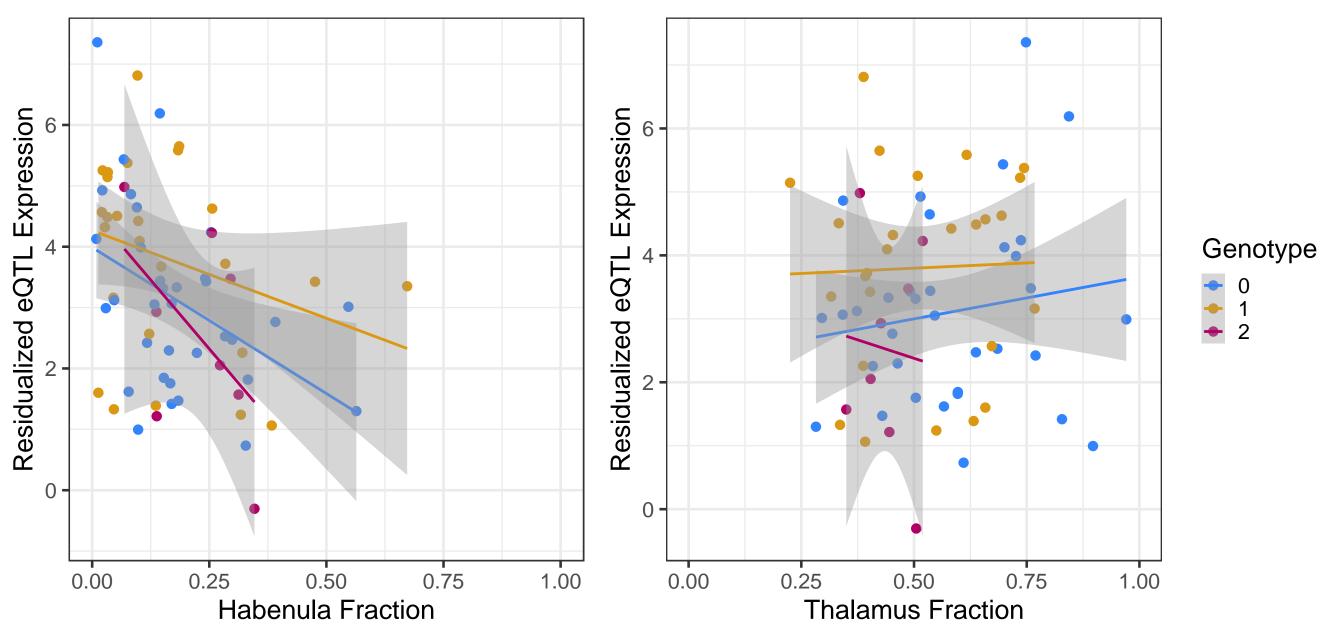
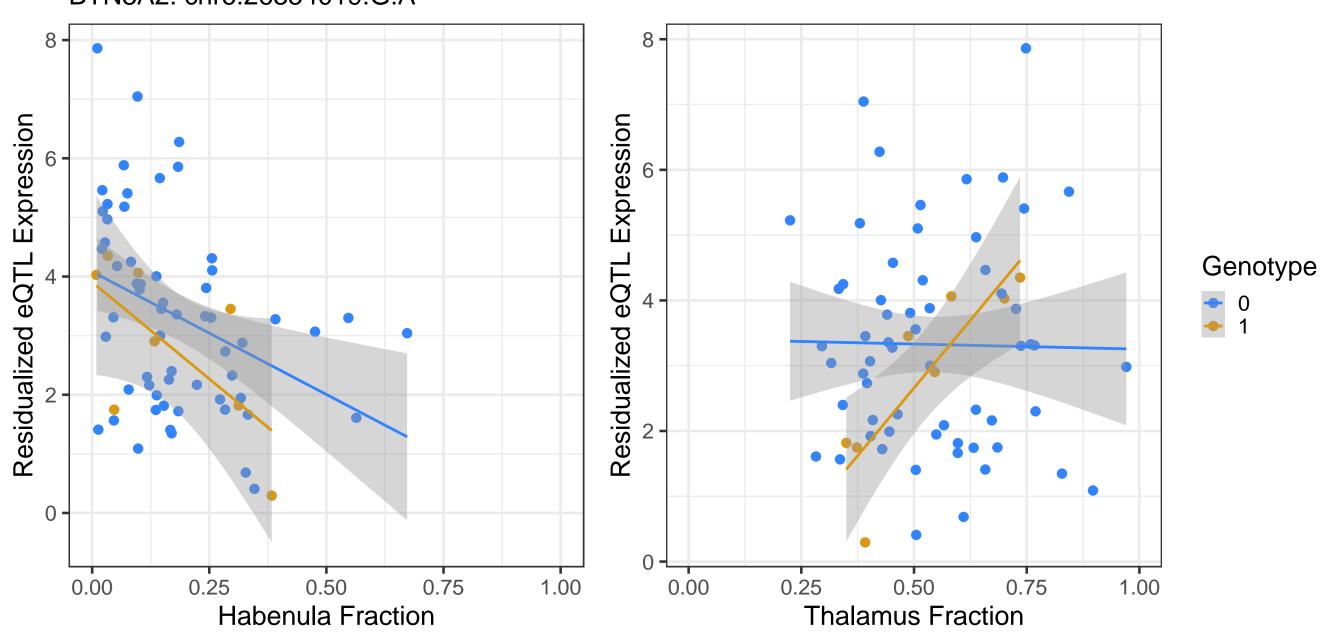
NEK4: chr3:52550699:A:G



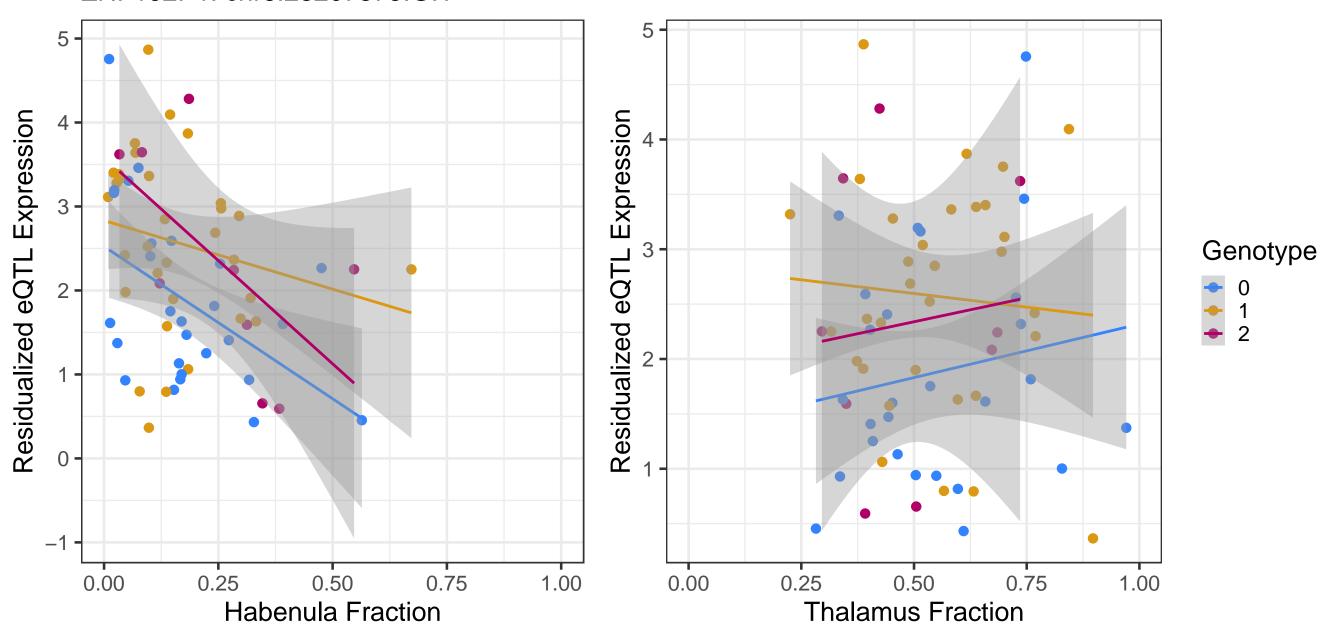
HIST1H4D: chr6:26183874:G:A



BTN3A2: chr6:26354019:G:A



ZNF192P1: chr6:28207375:G:T



ZNF603P: chr6:28128502:A:G 2 -Expression Genotype Residualized eQTI -2 **--4** -

0.00

0.50

Thalamus Fraction

0.25

0.75

1.00

Residualized eQTL Expression

0.25

0.00

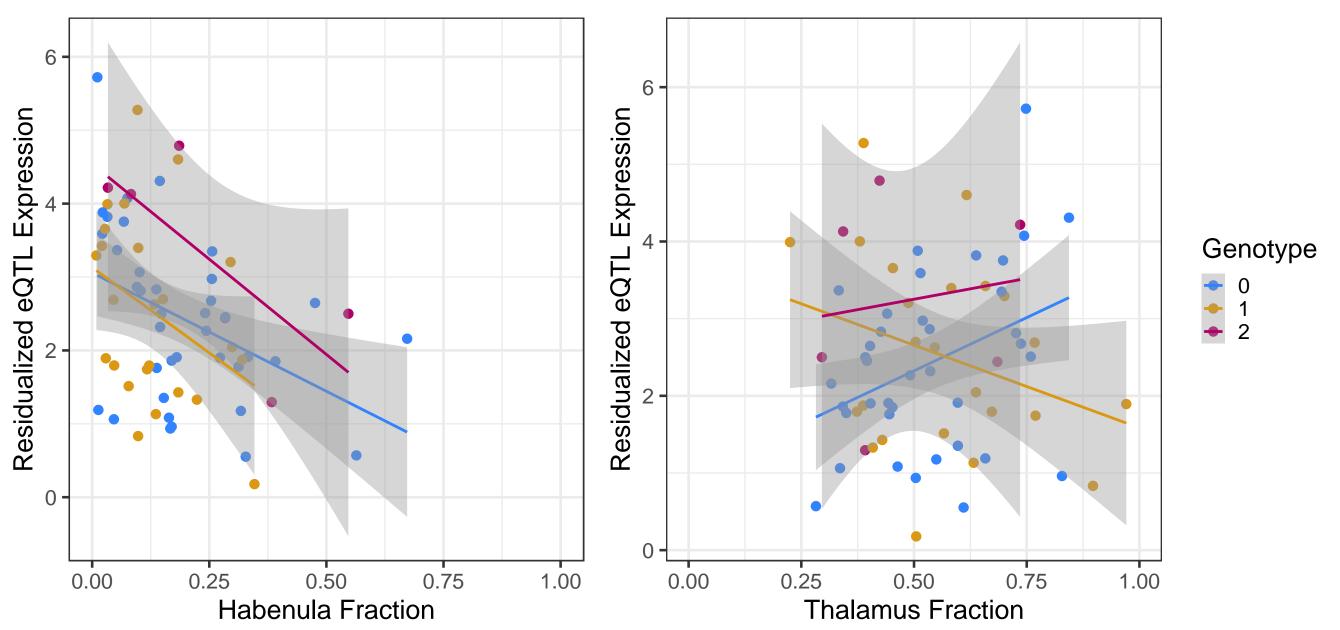
0.50

Habenula Fraction

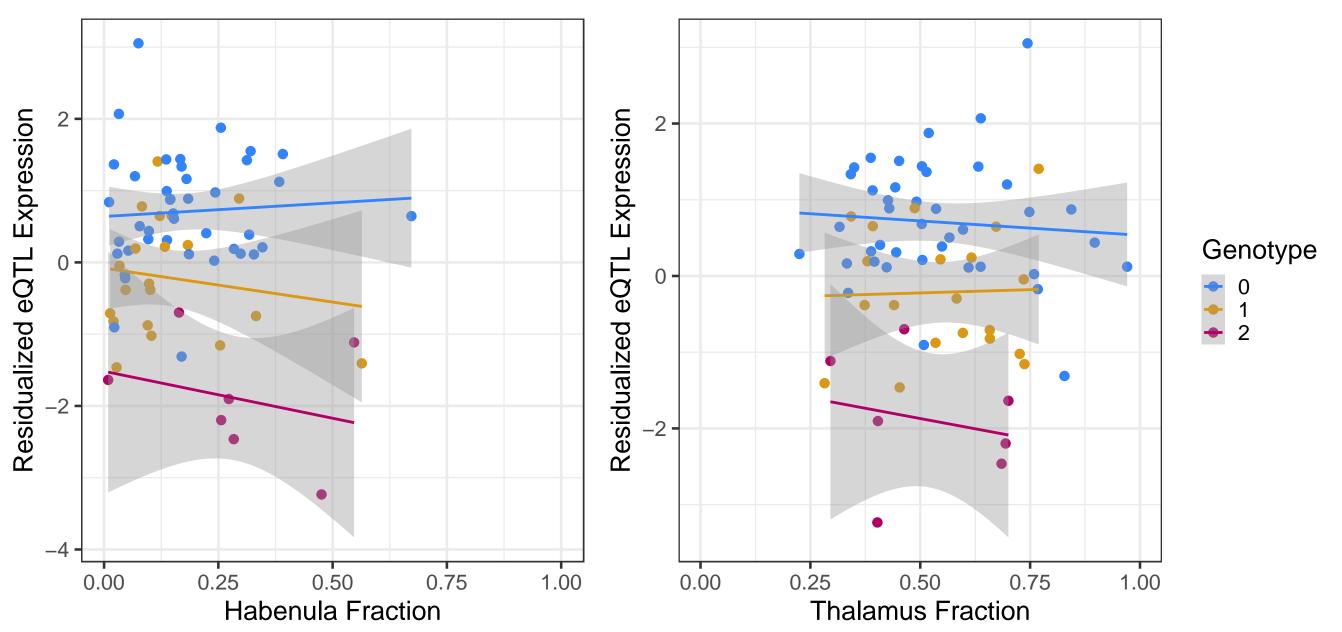
0.75

1.00

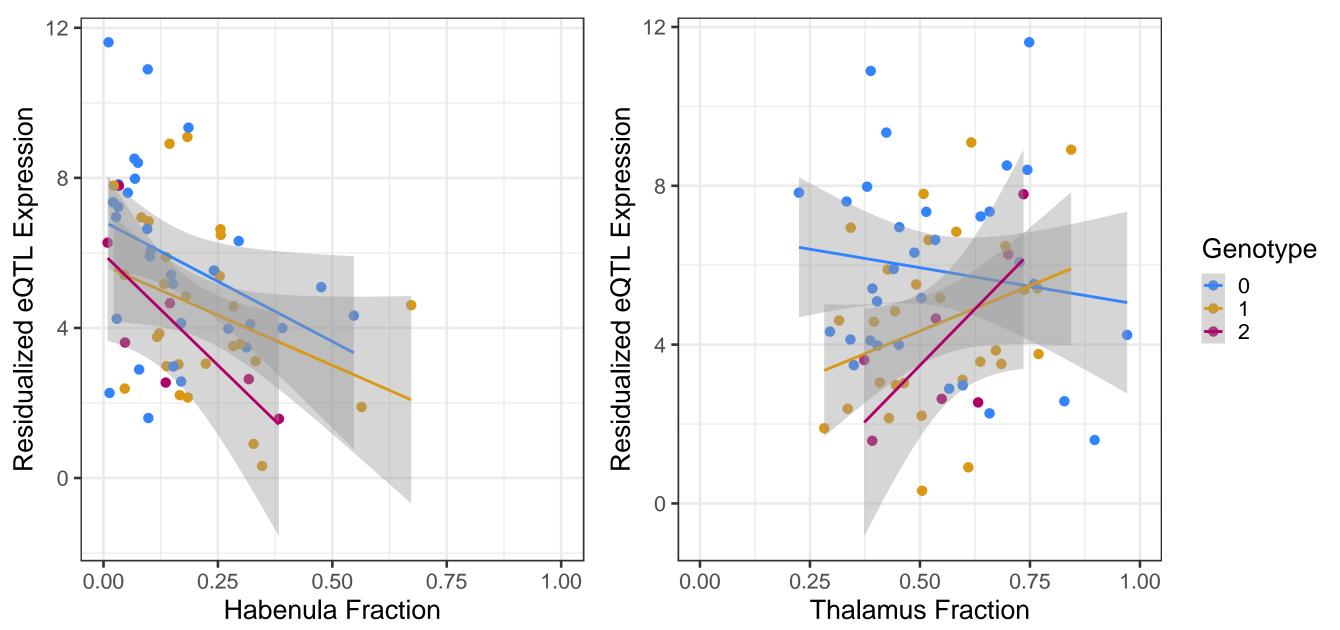
ZSCAN23: chr6:28273214:T:G



HLA-K: chr6:29861474:A:C

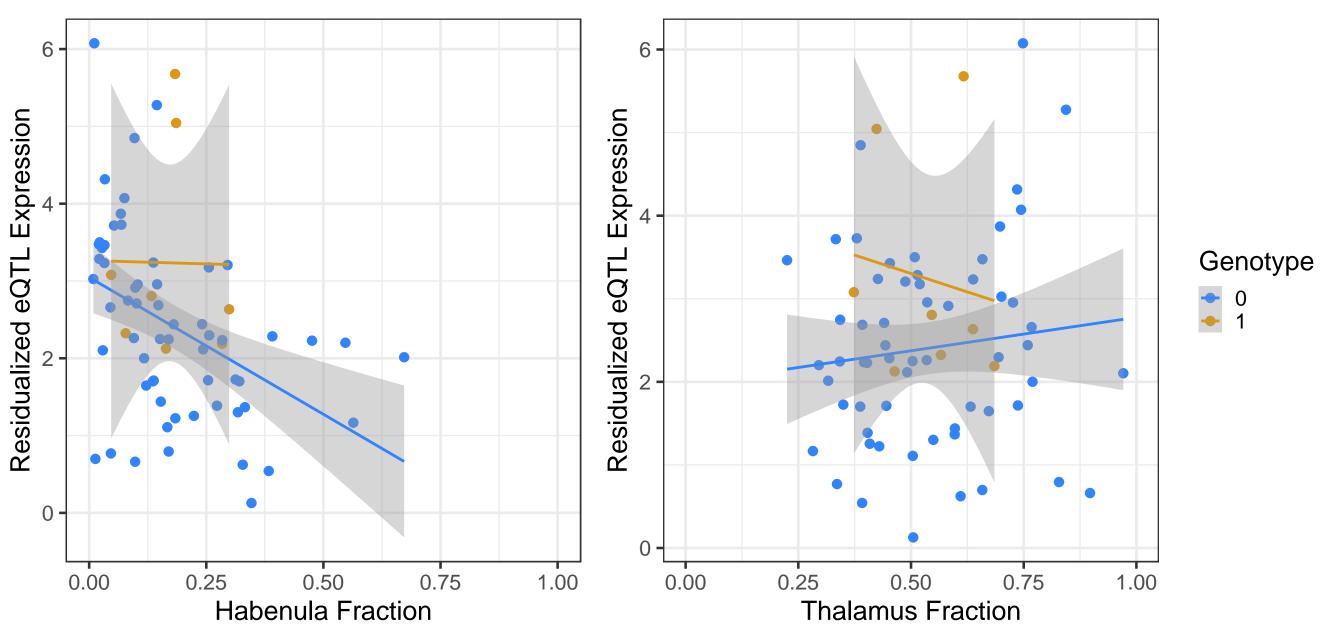


VARS2: chr6:30934095:C:G

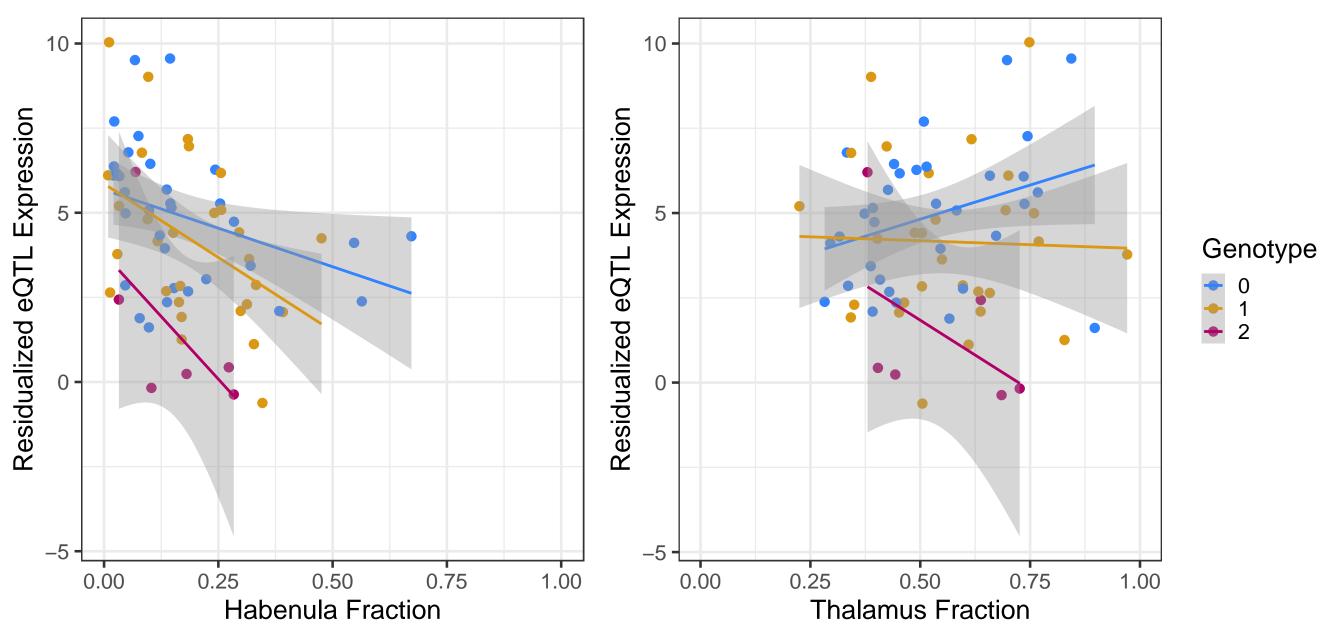


PSORS1C1: chr6:31025536:C:T 20 -0.0 -Residualized eQTL Expression Residualized eQTL Expression 10 --2.5 **-**Genotype 0 --5.0 **-**-10 **-**−7.5 **-**0.75 1.00 0.50 0.50 0.00 0.25 0.75 1.00 0.00 0.25 Habenula Fraction **Thalamus Fraction**

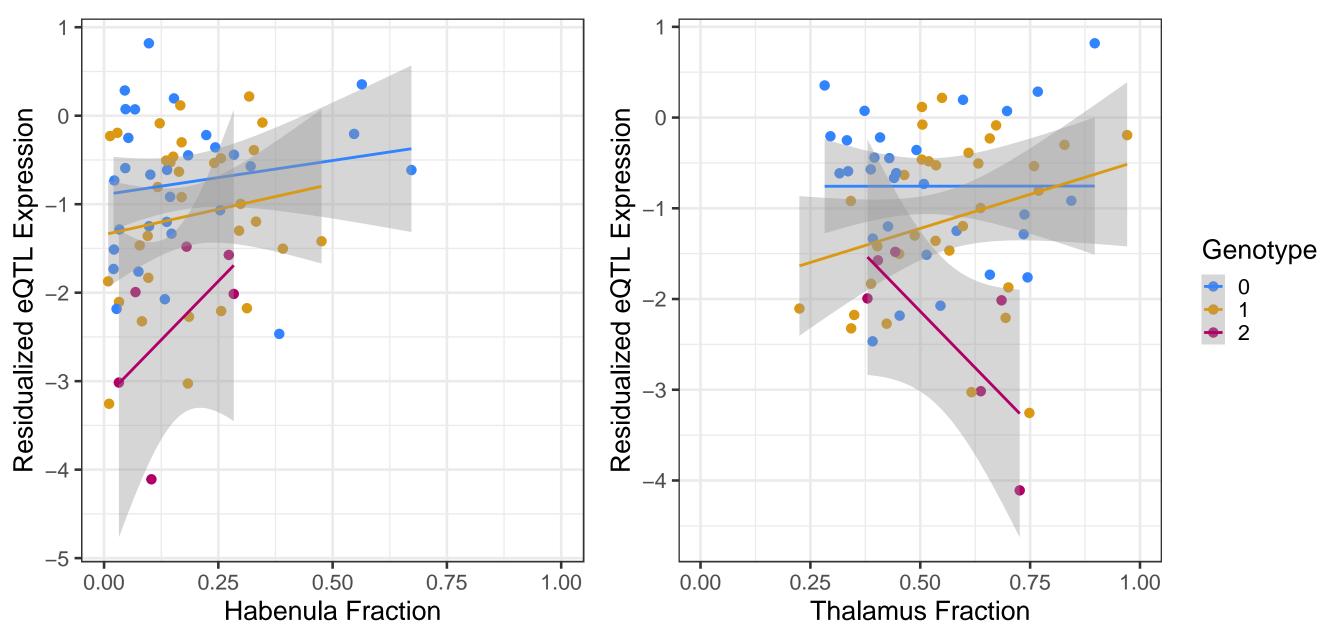
HLA-DMA: chr6:32895181:C:G



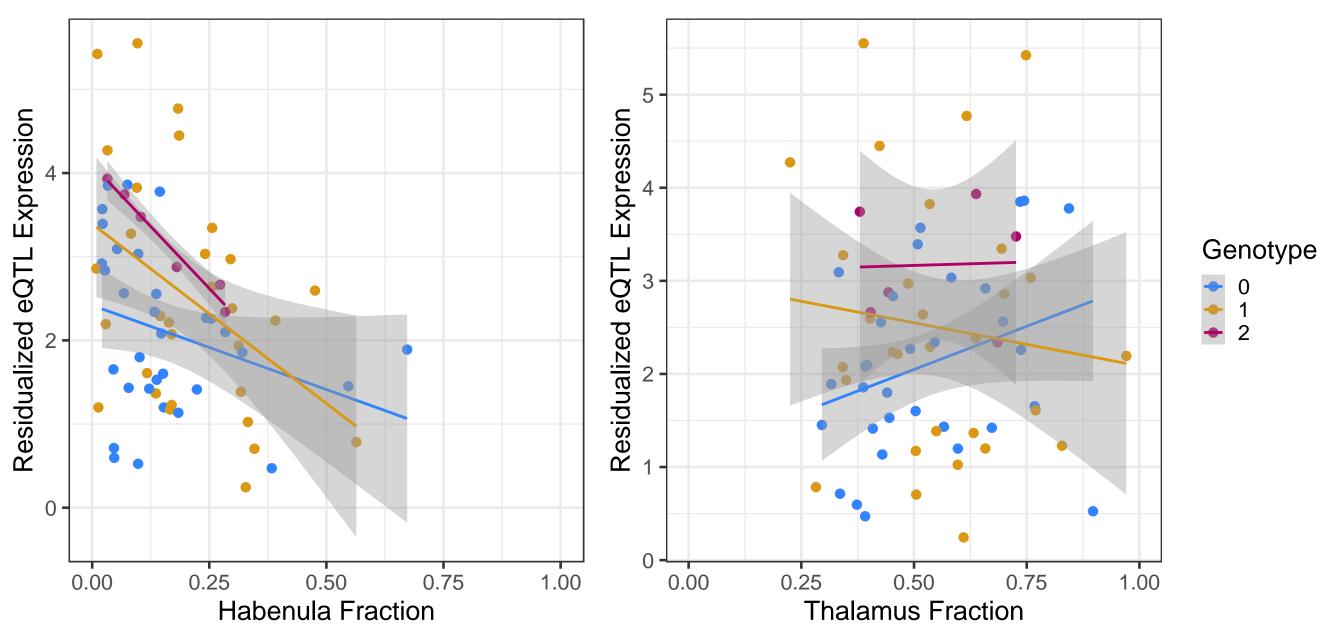
LRRC37A4P: chr17:45701314:T:A



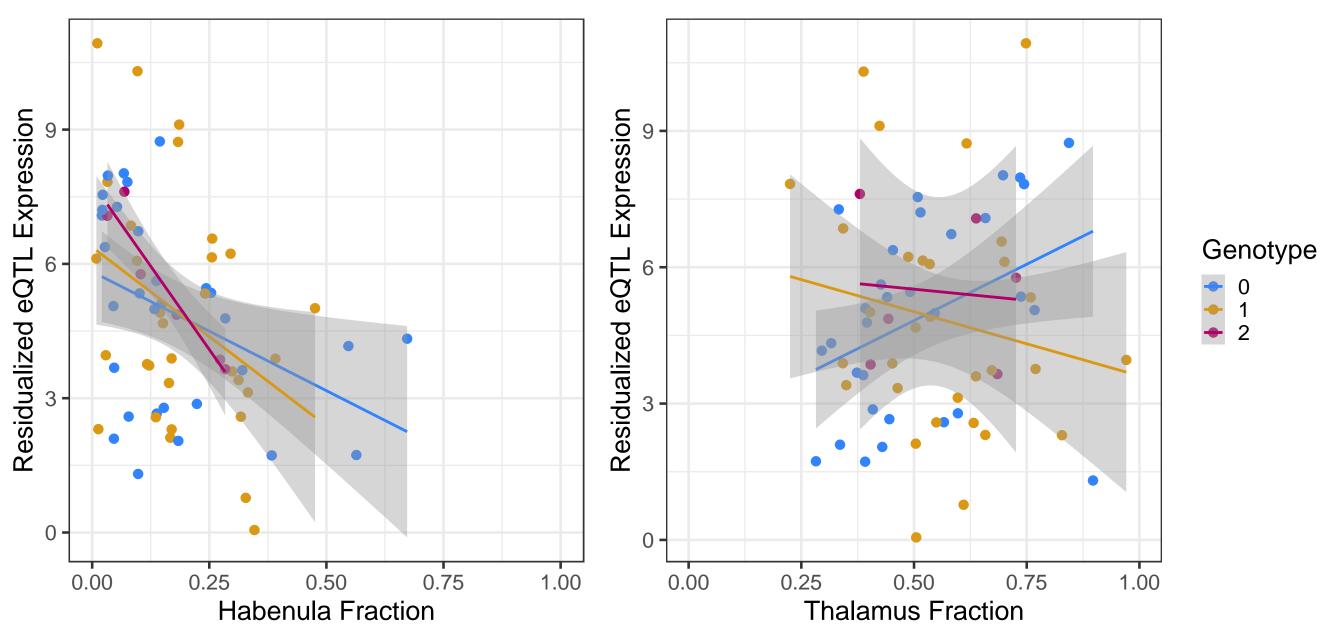
DND1P1: chr17:45910839:A:G



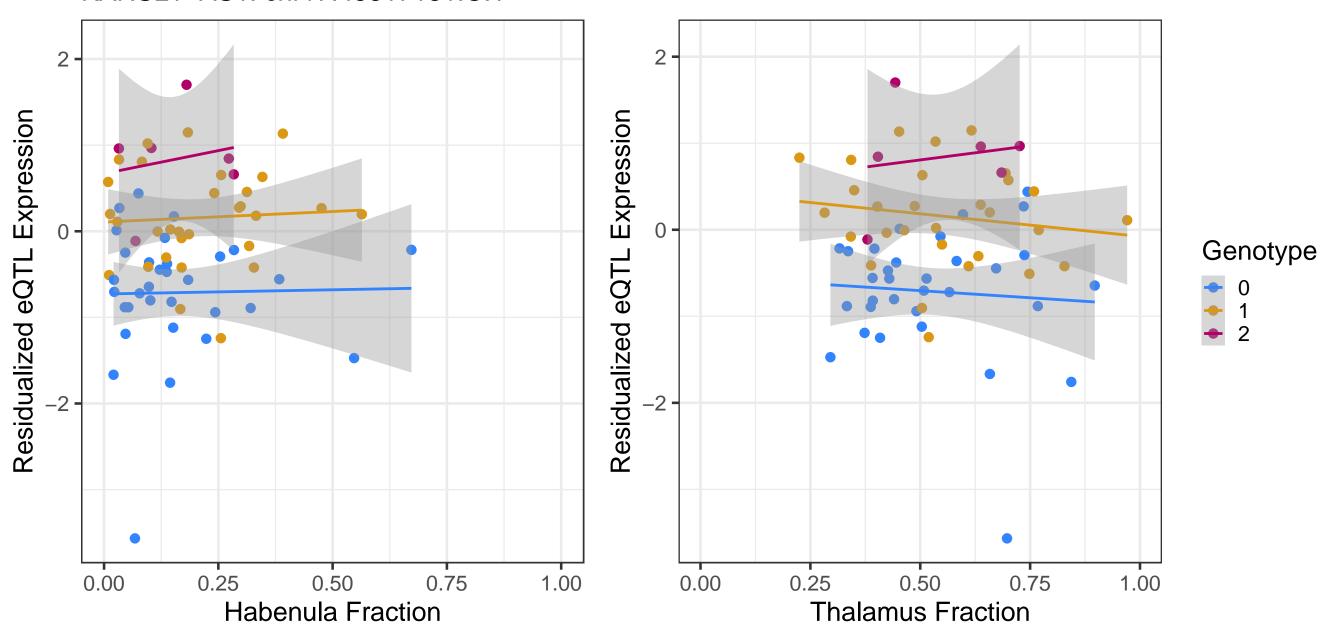
RP11-707O23.5: chr17:45817181:C:T



CRHR1-IT1: chr17:45887173:A:G



KANSL1-AS1: chr17:45817181:C:T



LRRC37A: chr17:45801942:A:G

