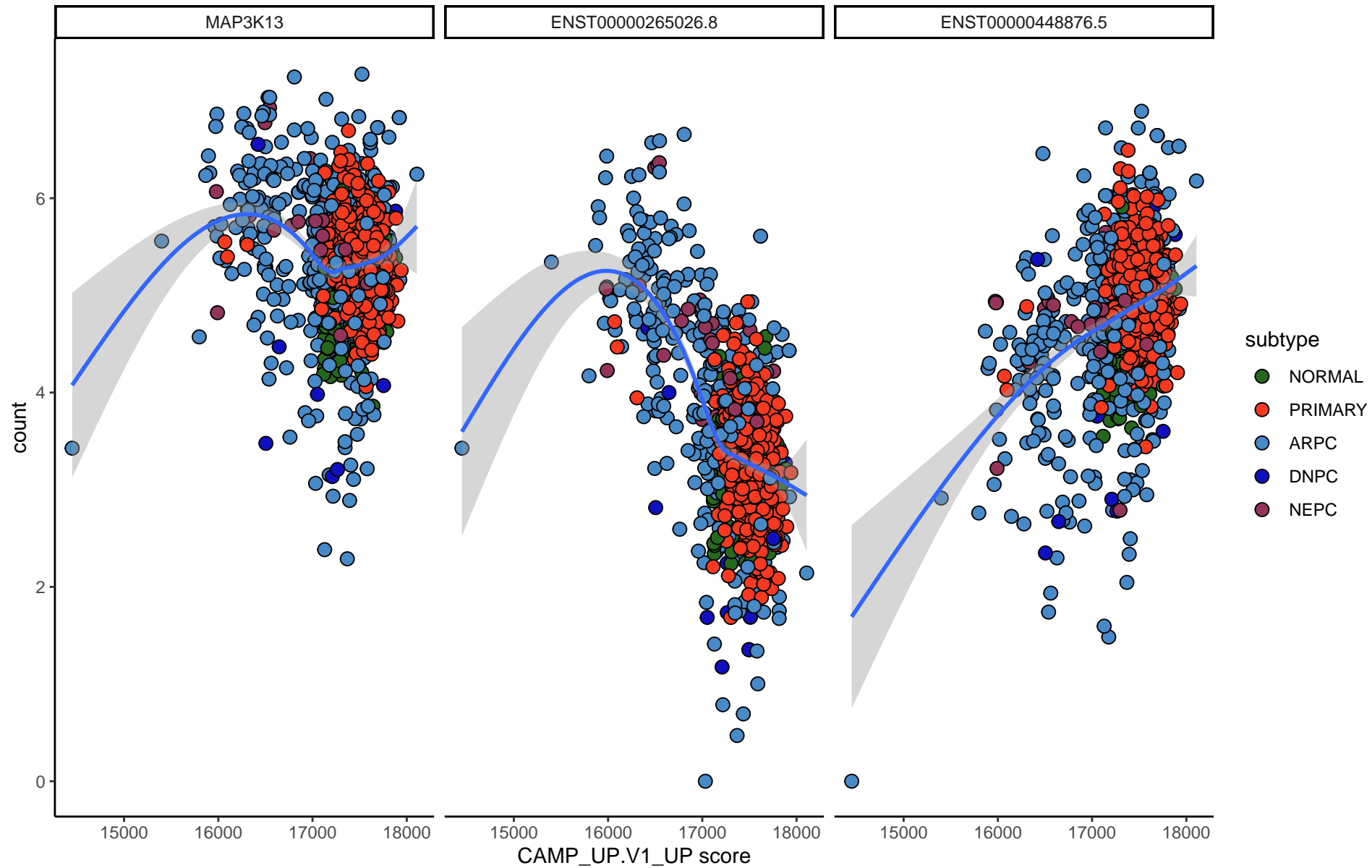


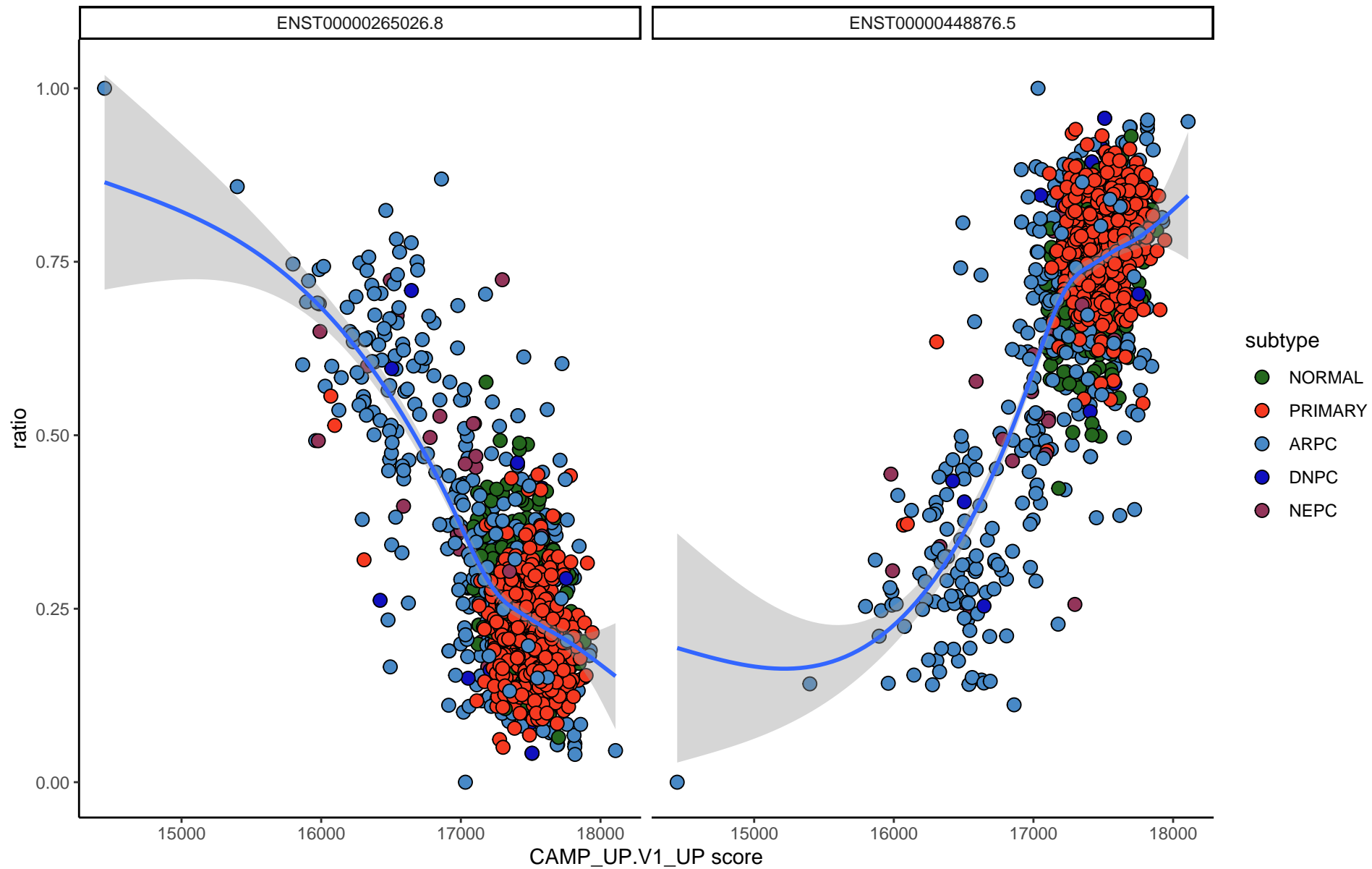
MAP3K13
Delta corr= 1.46
Gene corr= -0.14



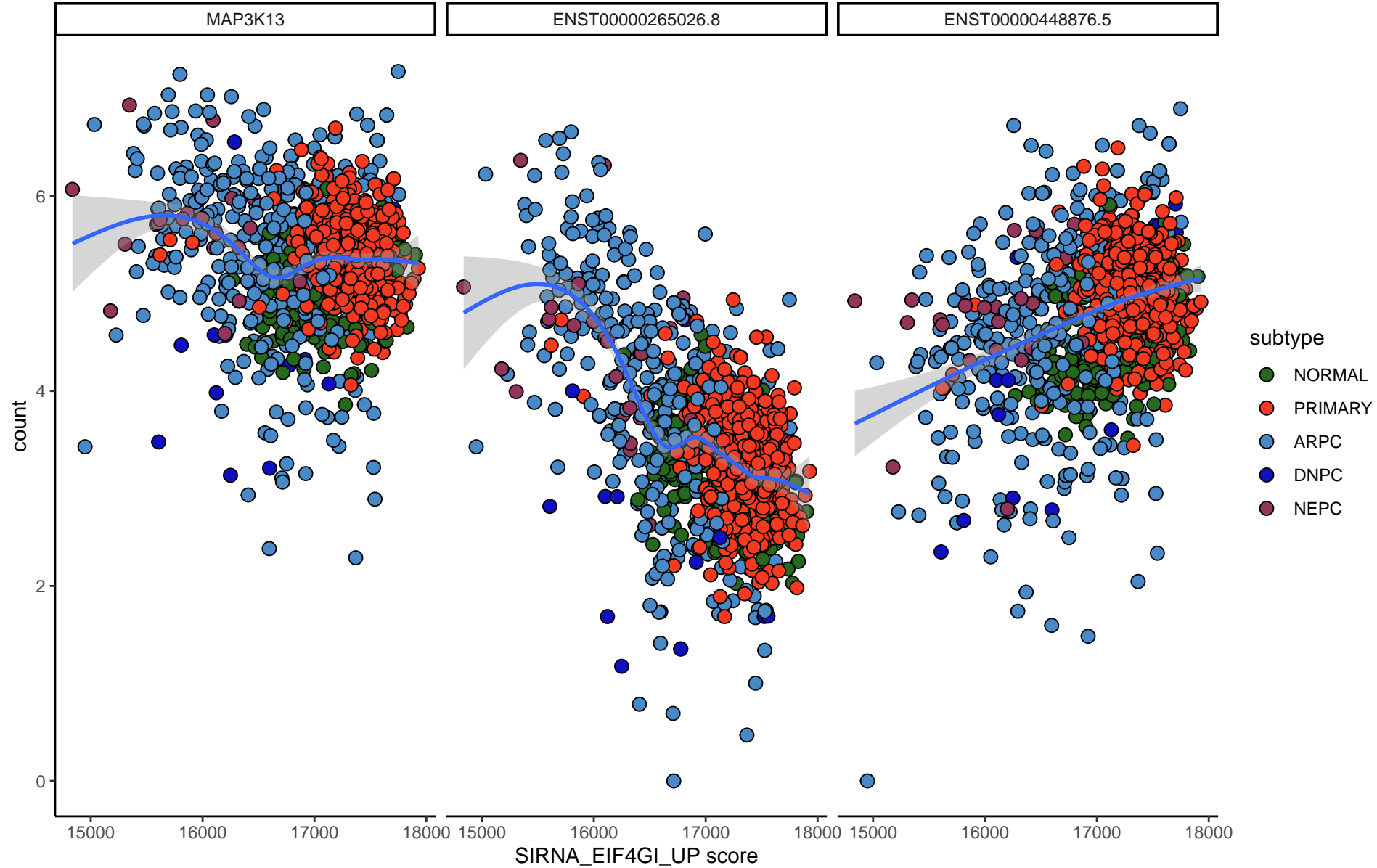
CAMP_UP.V1_UP

ENST00000265026.8 – MAP3K13–201 PCC = -0.72

ENST00000448876.5 – MAP3K13–212 PCC = 0.75



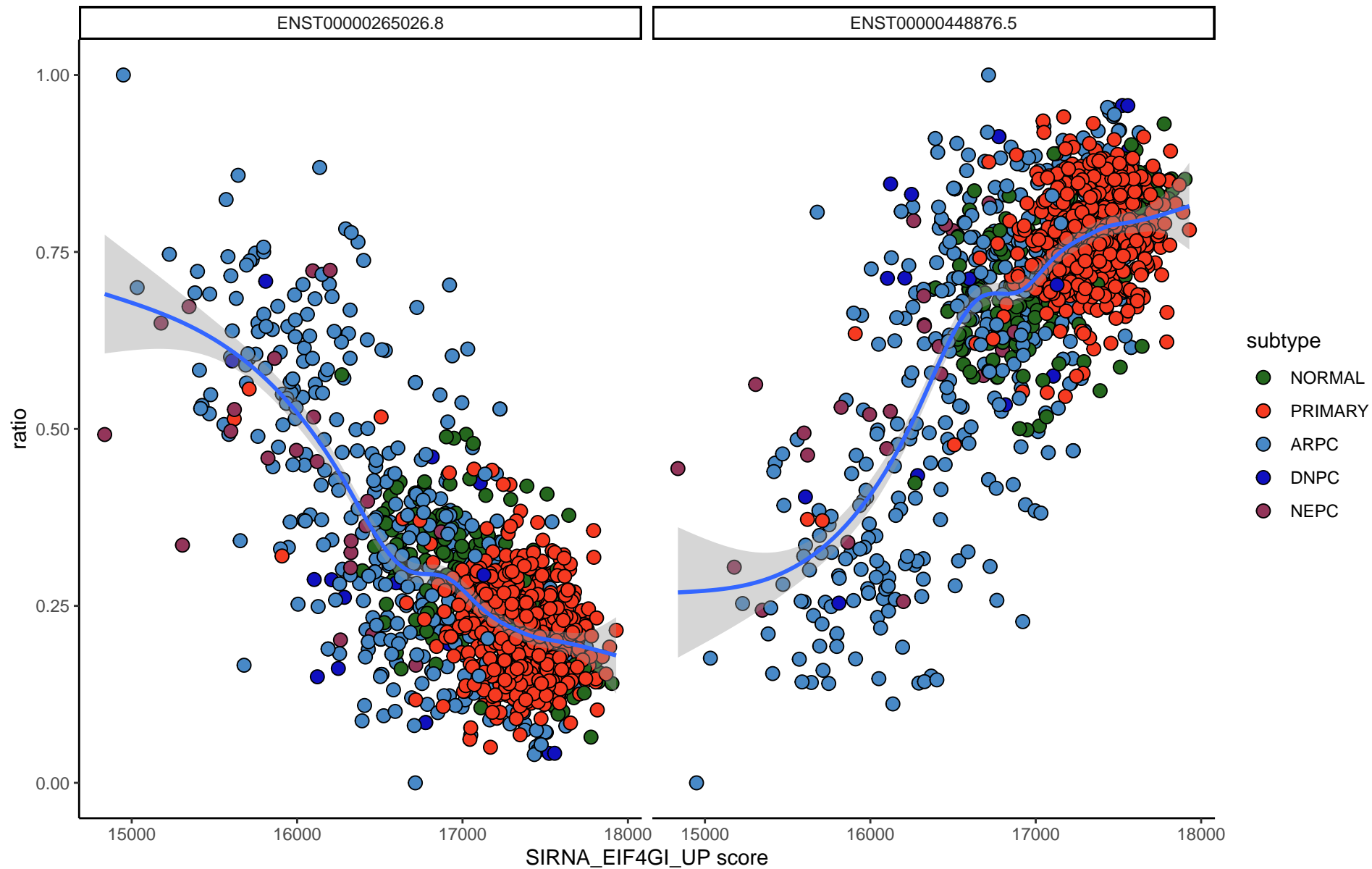
MAP3K13
Delta corr= 1.43
Gene corr= -0.12



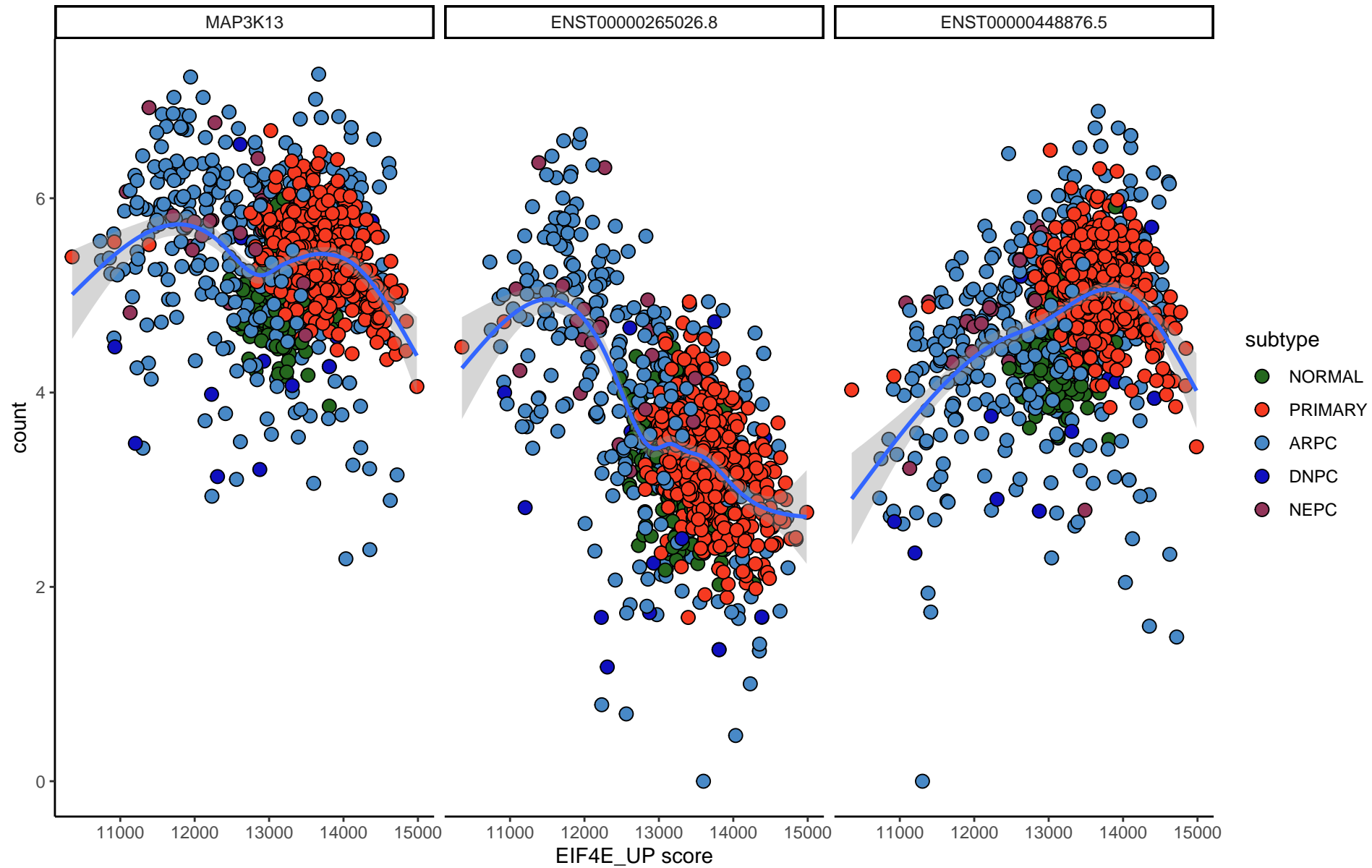
SIRNA_EIF4GI_UP

ENST00000265026.8 – MAP3K13–201 PCC = -0.7

ENST00000448876.5 – MAP3K13–212 PCC = 0.72



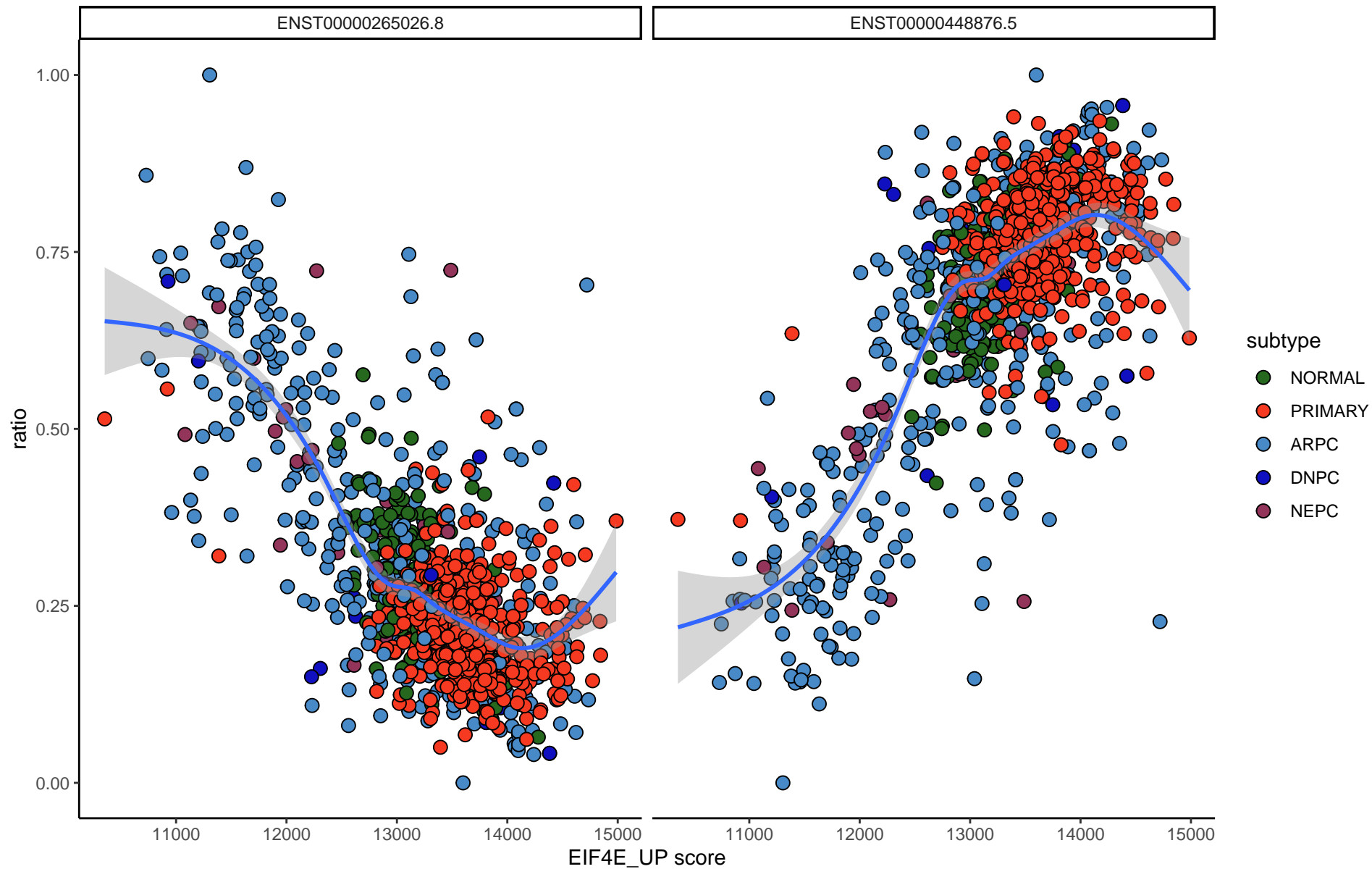
MAP3K13
Delta corr= 1.4
Gene corr= -0.14



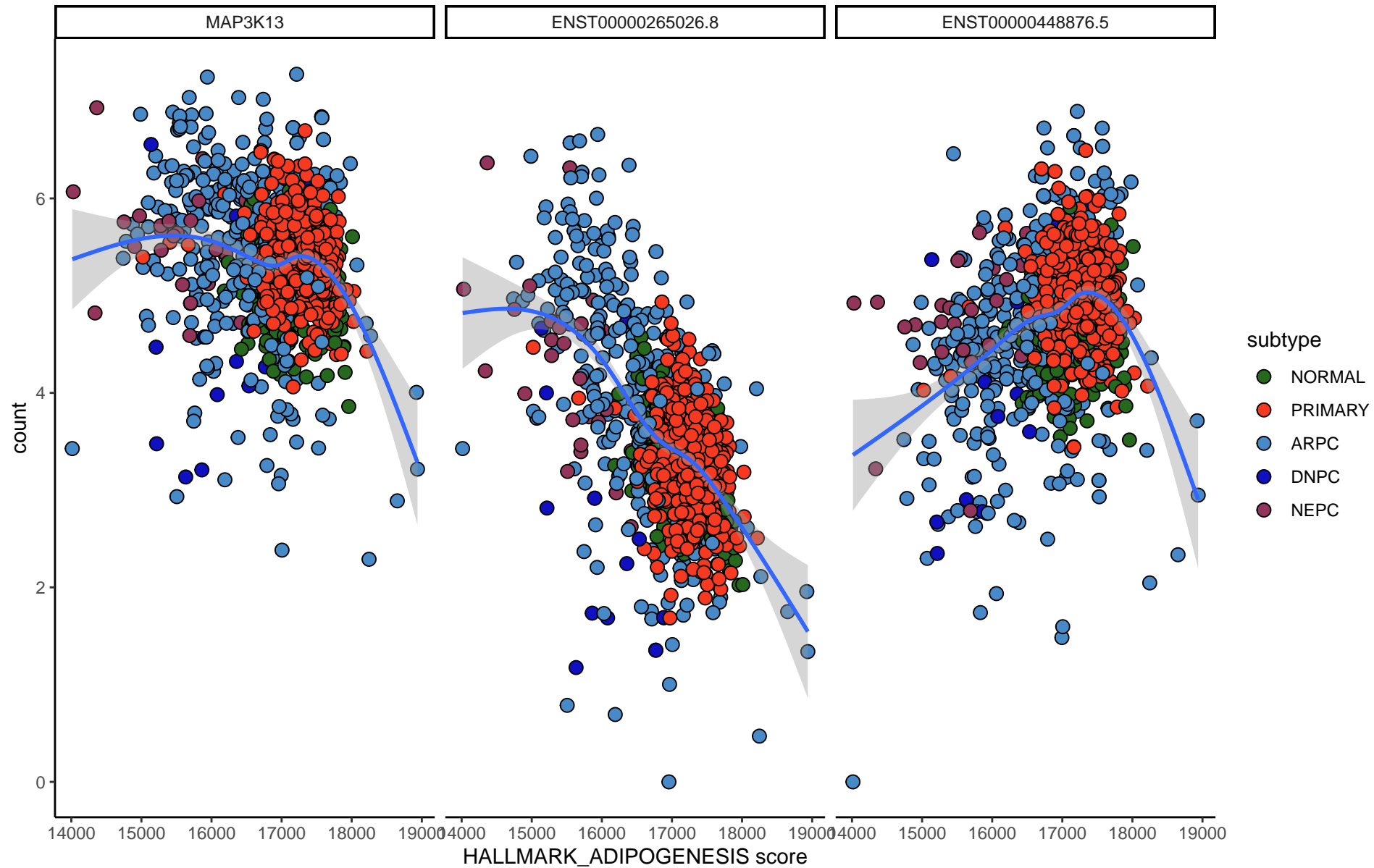
EIF4E_UP

ENST00000265026.8 – MAP3K13–201 PCC = -0.68

ENST00000448876.5 – MAP3K13–212 PCC = 0.72



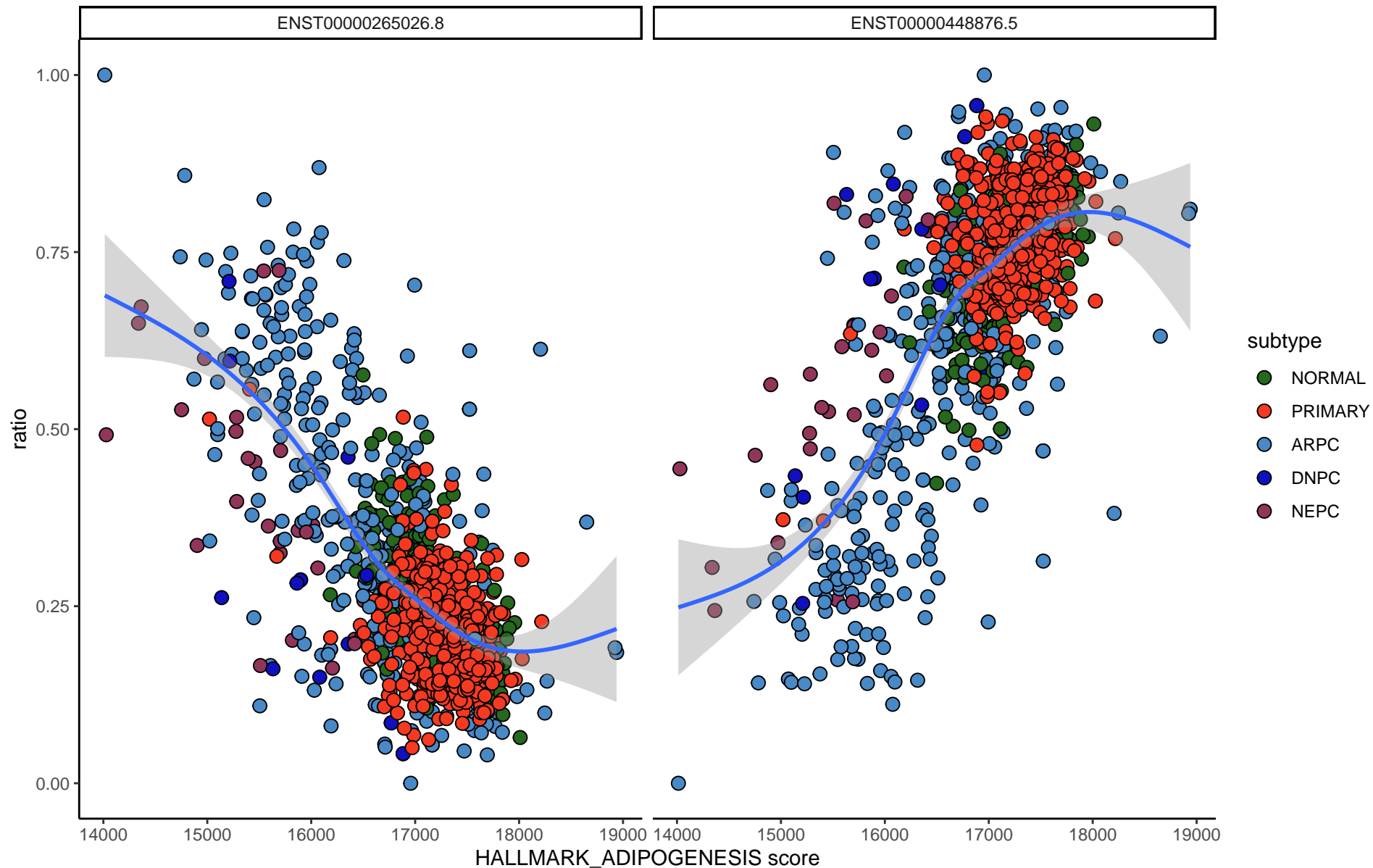
MAP3K13
Delta corr= 1.38
Gene corr= -0.17



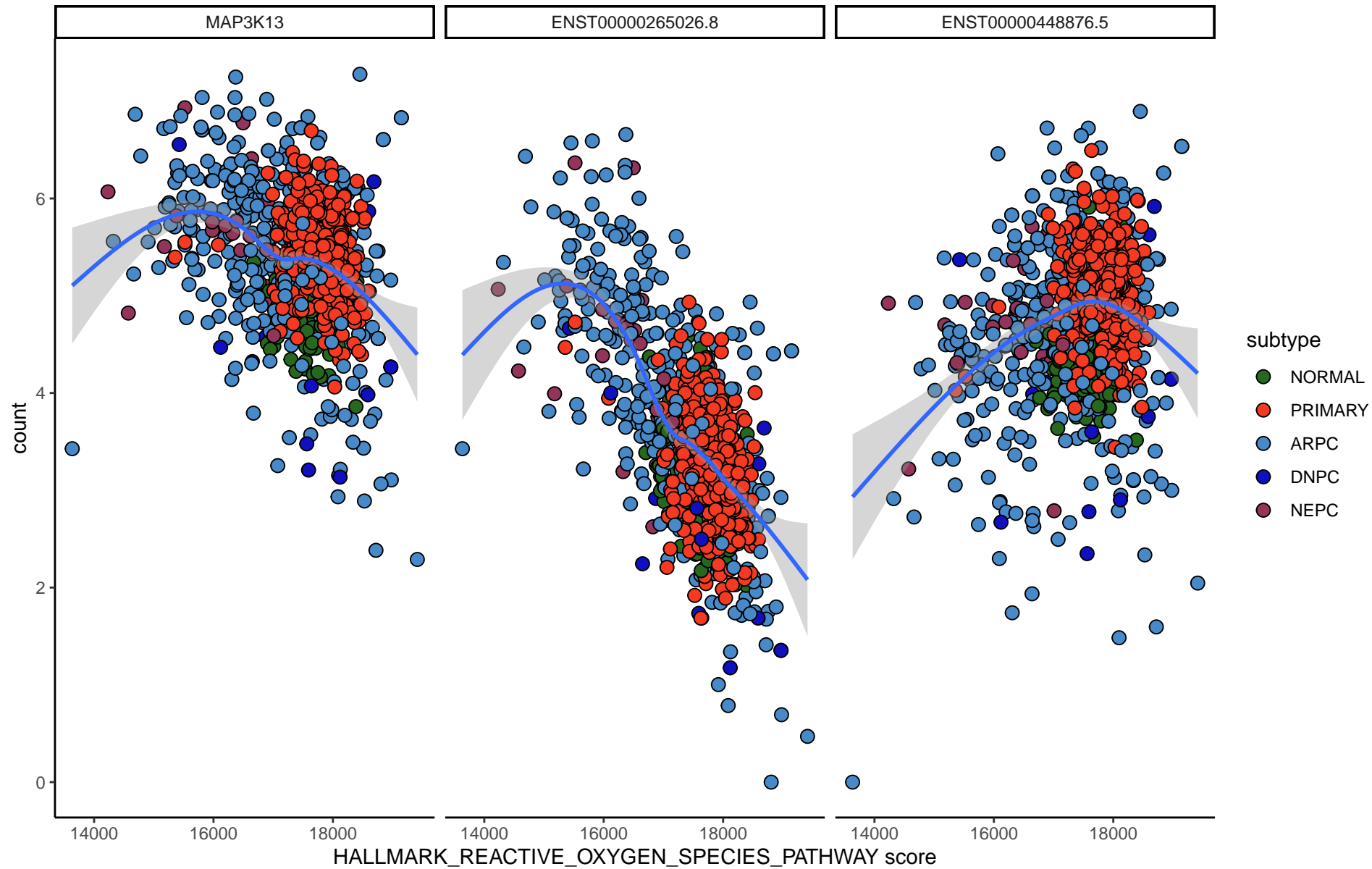
HALLMARK_ADIPOGENESIS

ENST00000265026.8 – MAP3K13–201 PCC = -0.68

ENST00000448876.5 – MAP3K13–212 PCC = 0.71



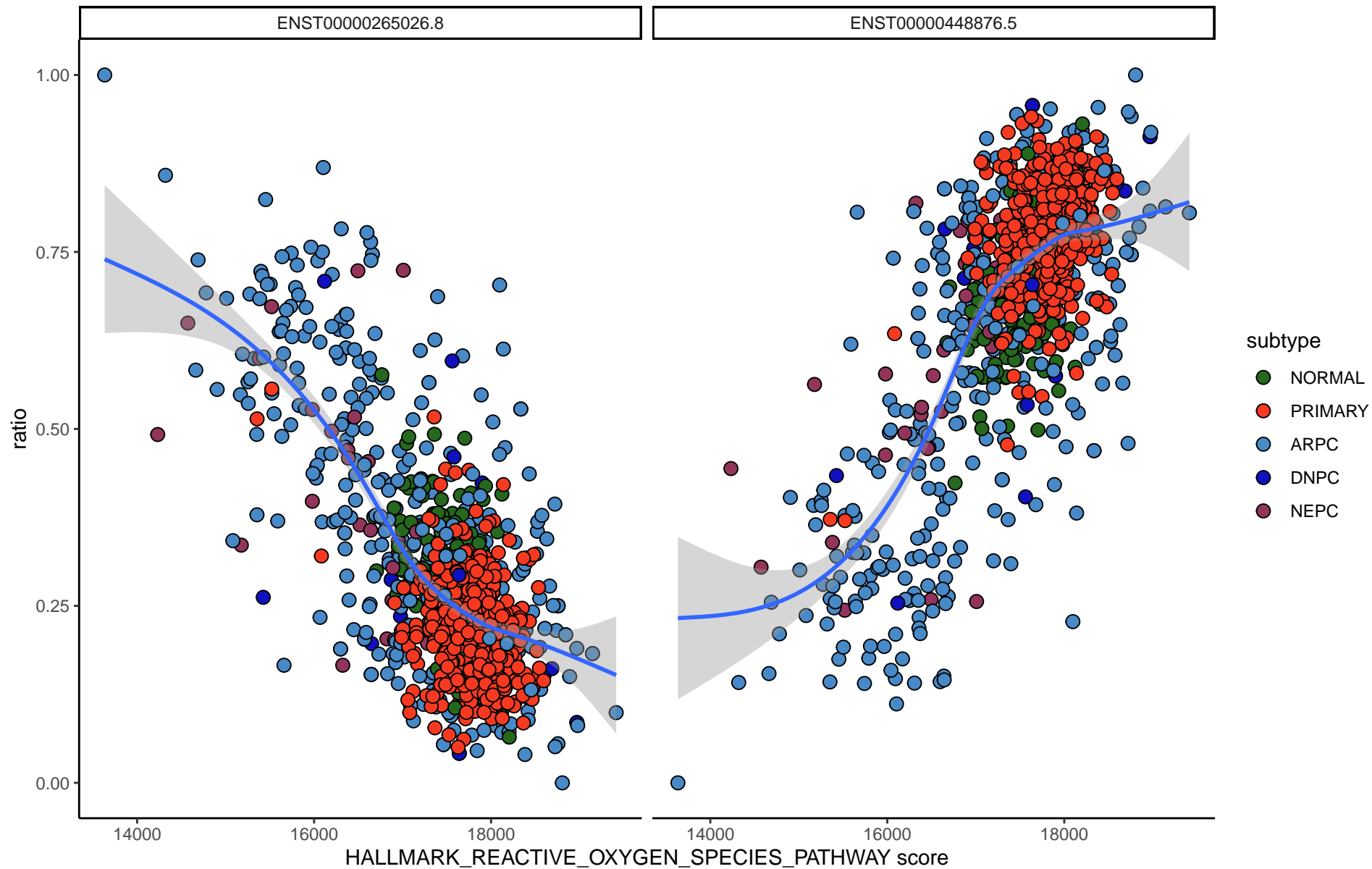
MAP3K13
Delta corr= 1.38
Gene corr= -0.29



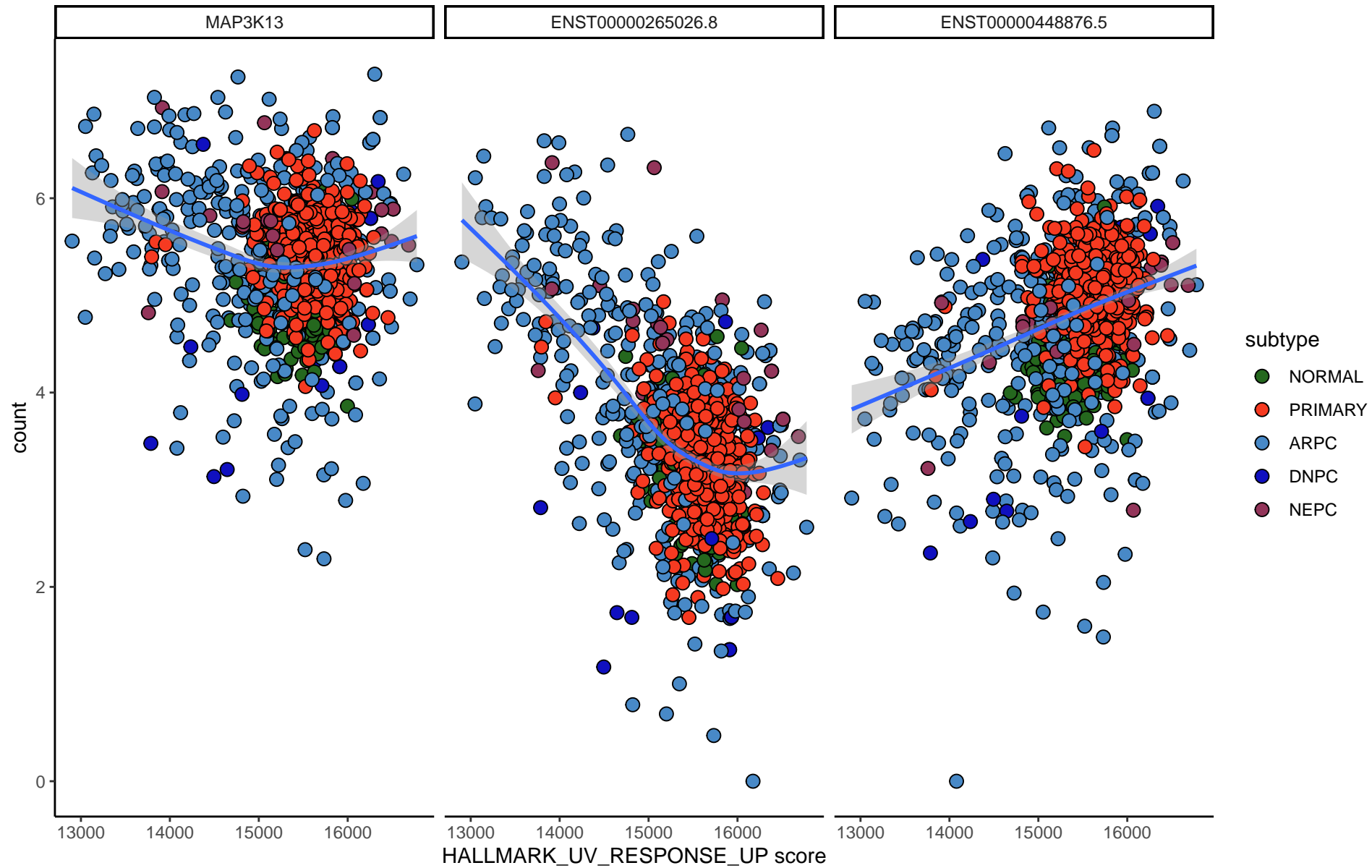
HALLMARK_REACTIVE_OXYGEN_SPECIES_PATHWAY

ENST00000265026.8 – MAP3K13–201 PCC = -0.67

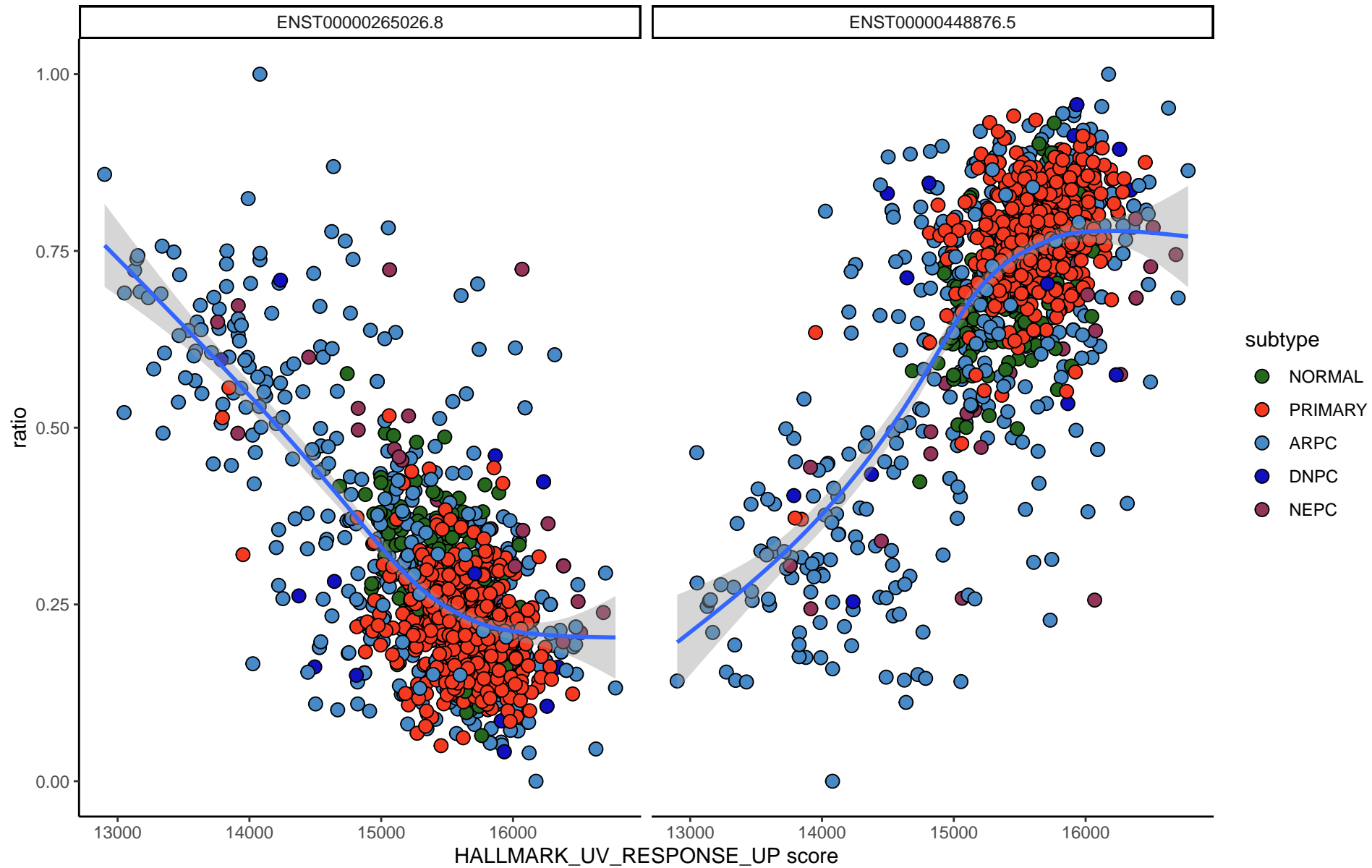
ENST00000448876.5 – MAP3K13–212 PCC = 0.71



MAP3K13
Delta corr= 1.36
Gene corr= -0.13



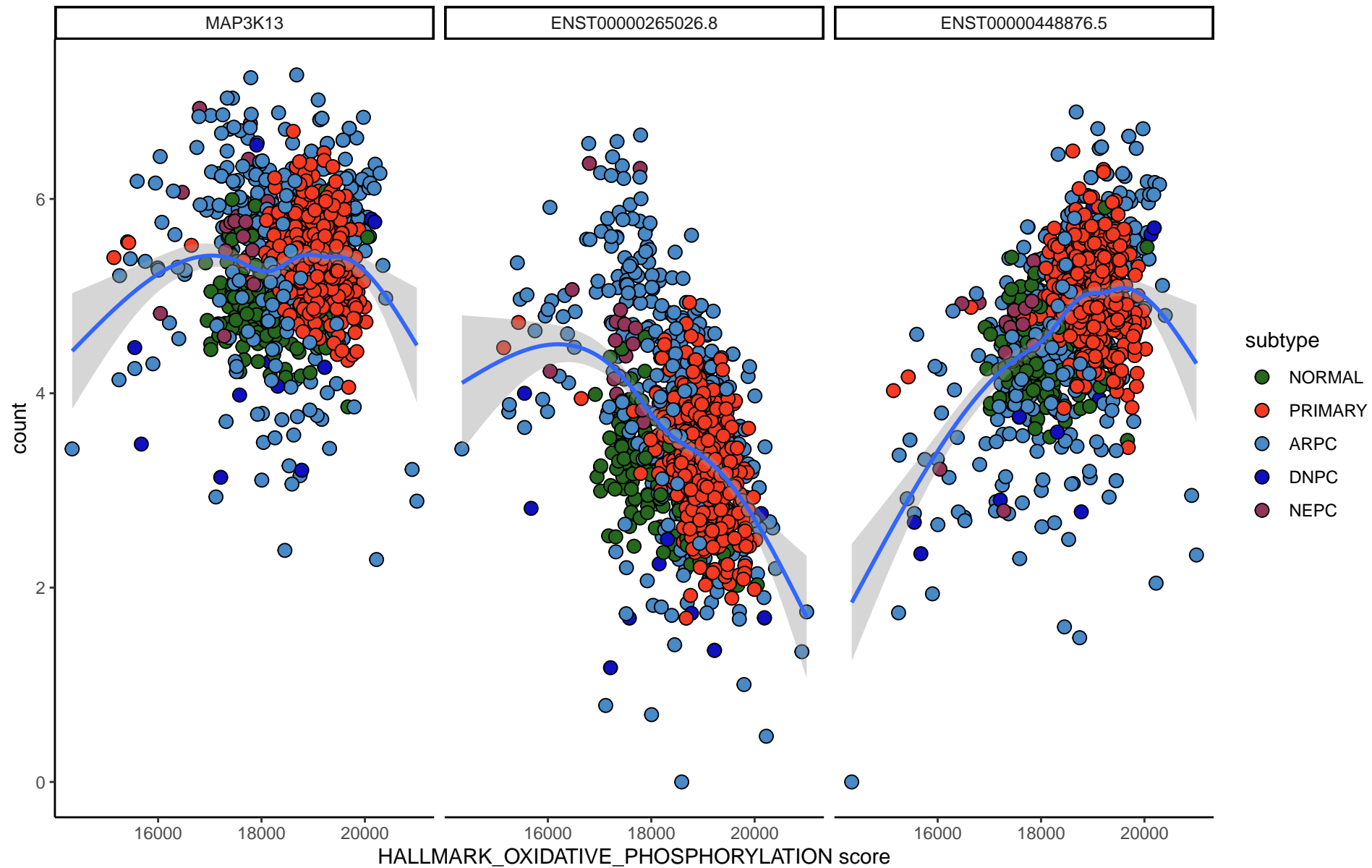
HALLMARK_UV_RESPONSE_UP
ENST00000265026.8 – MAP3K13–201 PCC = -0.67
ENST00000448876.5 – MAP3K13–212 PCC = 0.69



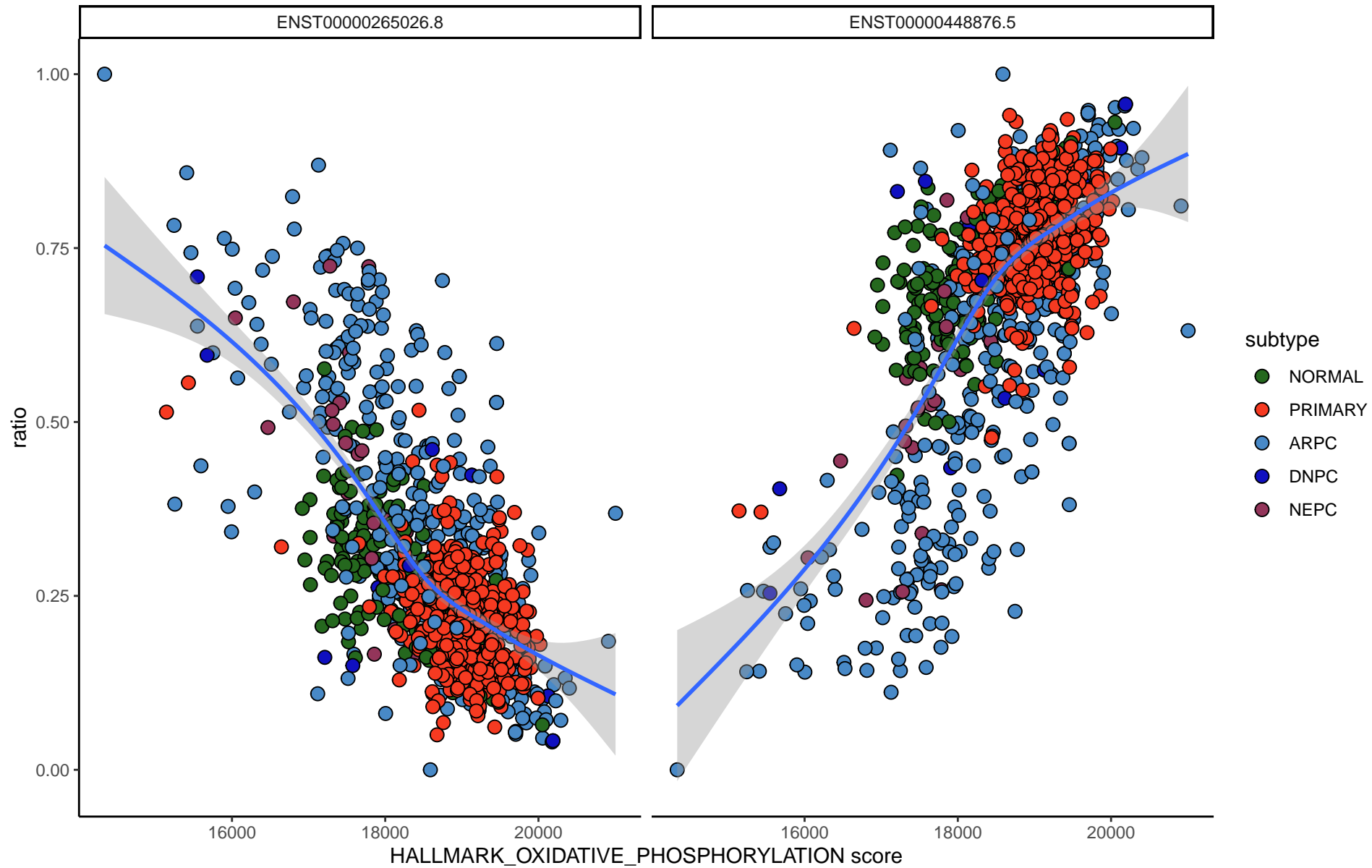
MAP3K13

Delta corr= 1.35

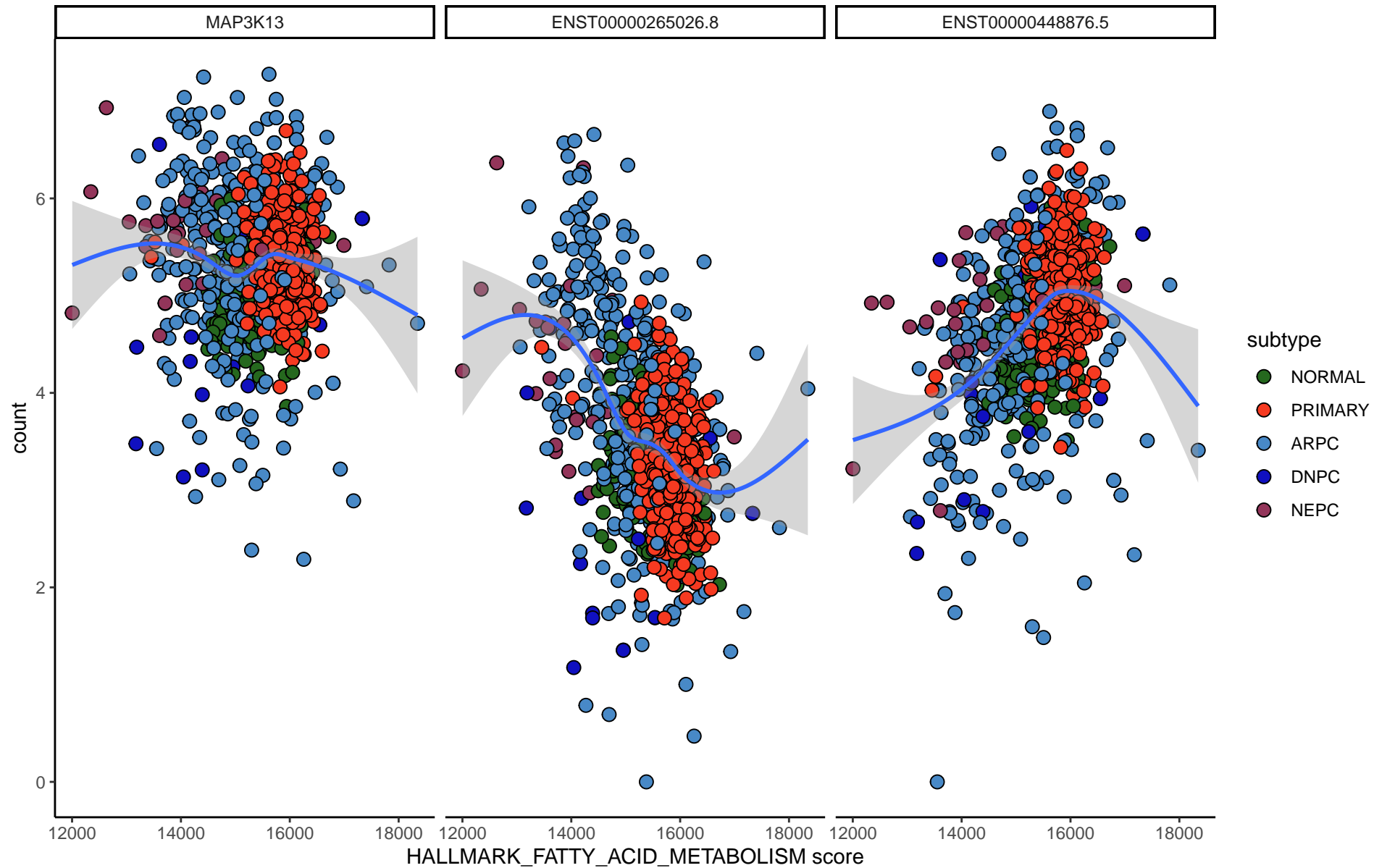
Gene corr= 0.05



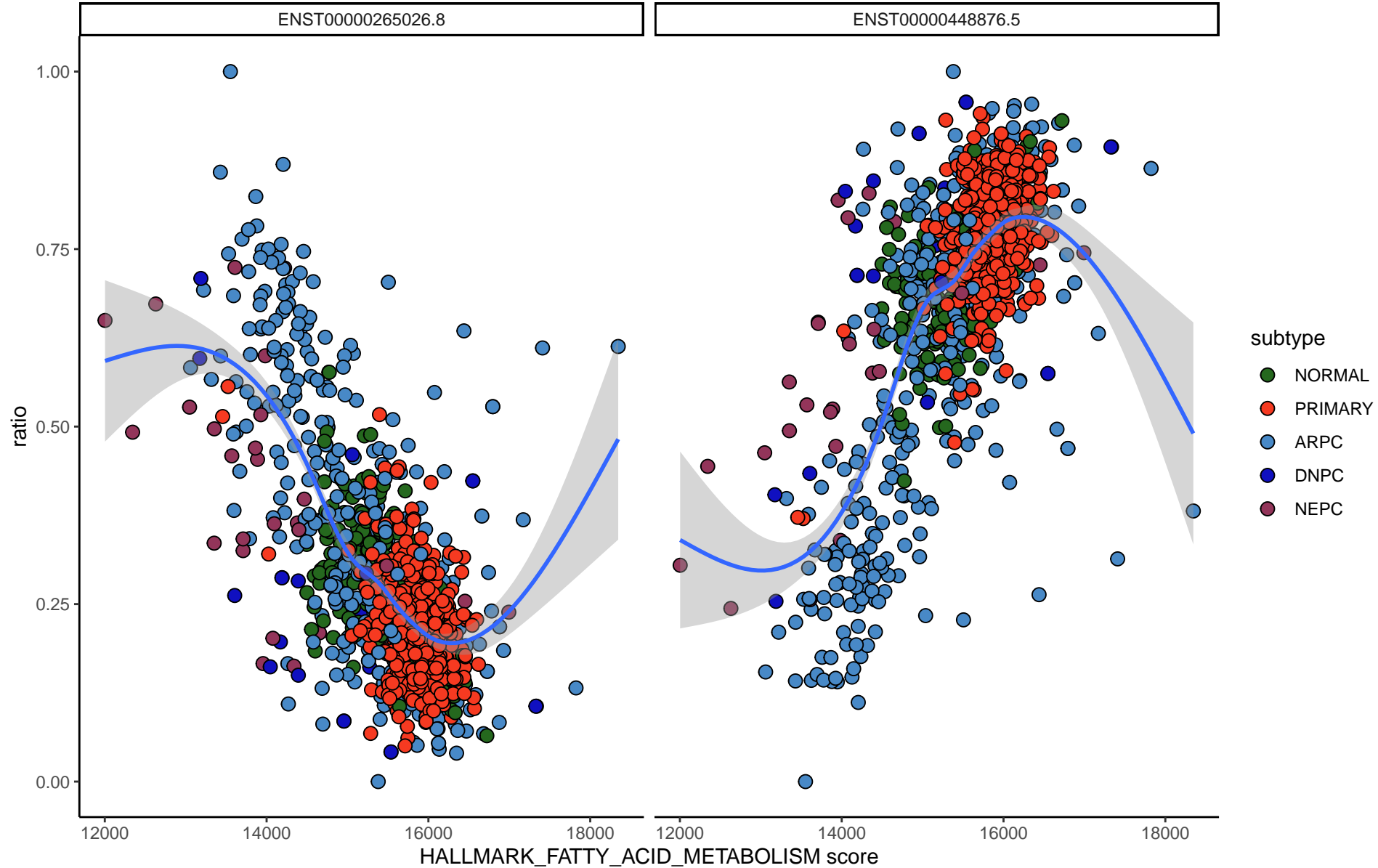
HALLMARK_OXIDATIVE_PHOSPHORYLATION
ENST00000265026.8 – MAP3K13–201 PCC = -0.67
ENST00000448876.5 – MAP3K13–212 PCC = 0.69



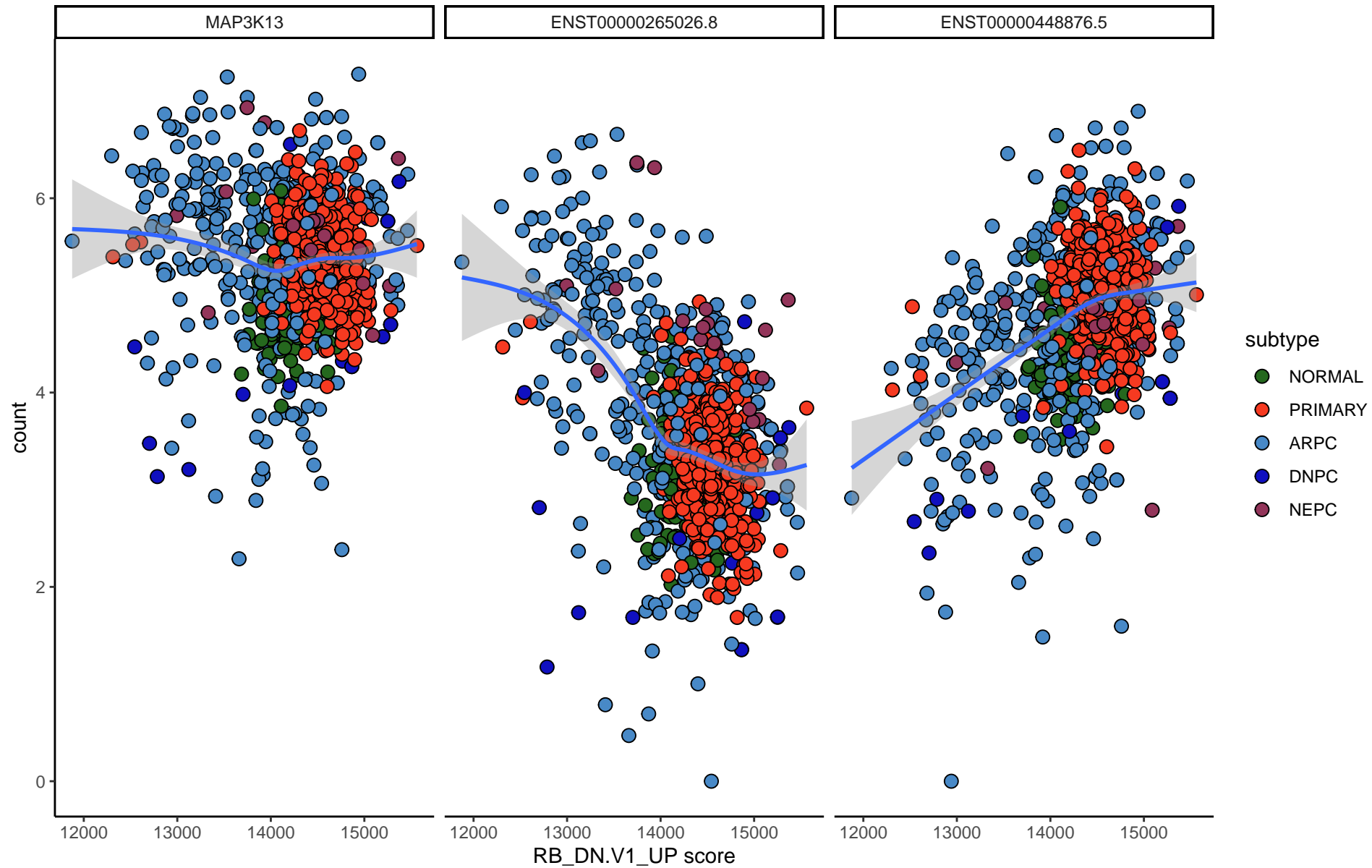
MAP3K13
Delta corr= 1.34
Gene corr= -0.02



HALLMARK_FATTY_ACID_METABOLISM
ENST00000265026.8 – MAP3K13–201 PCC = -0.66
ENST00000448876.5 – MAP3K13–212 PCC = 0.68



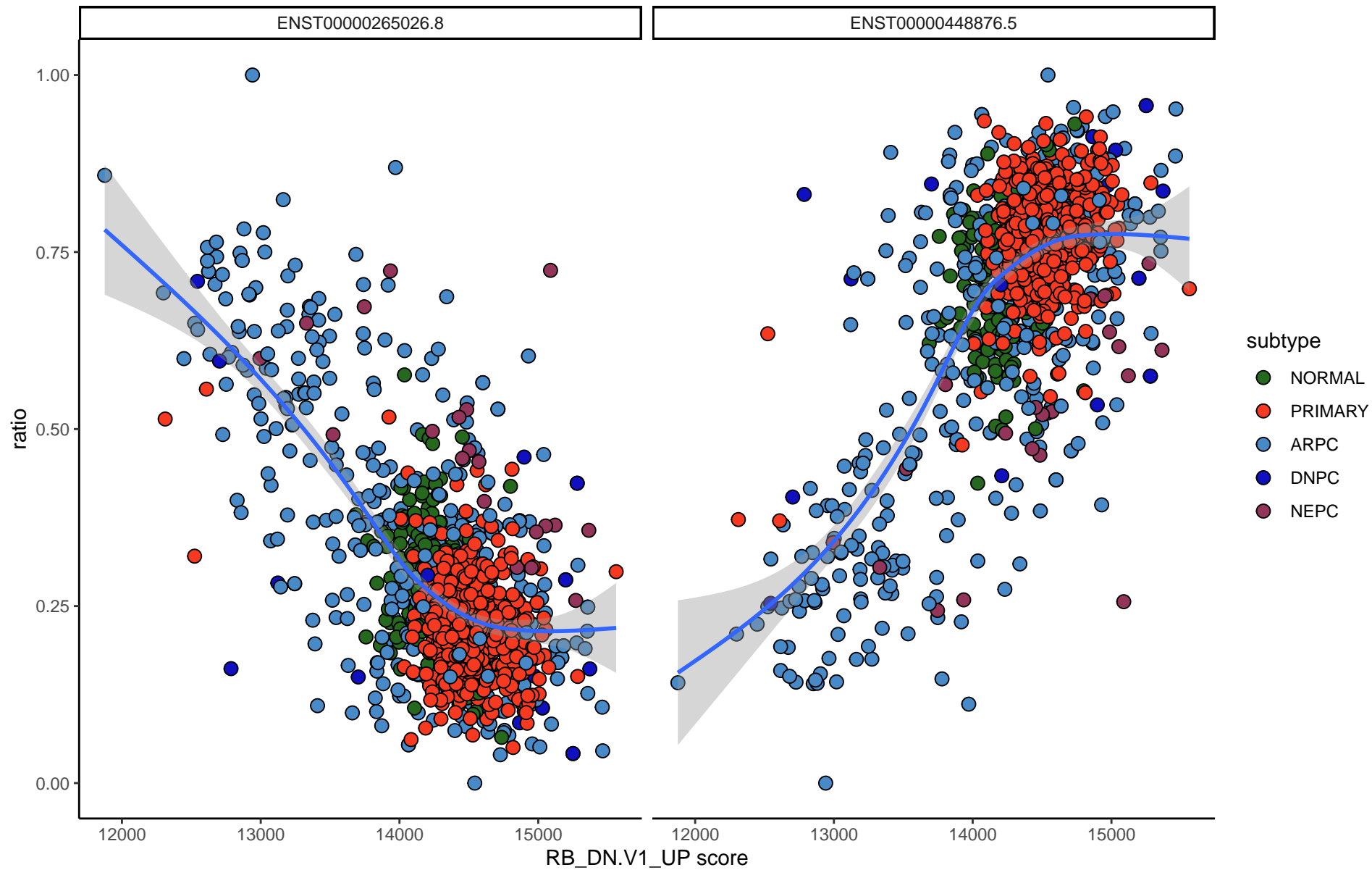
MAP3K13
Delta corr= 1.31
Gene corr= -0.05



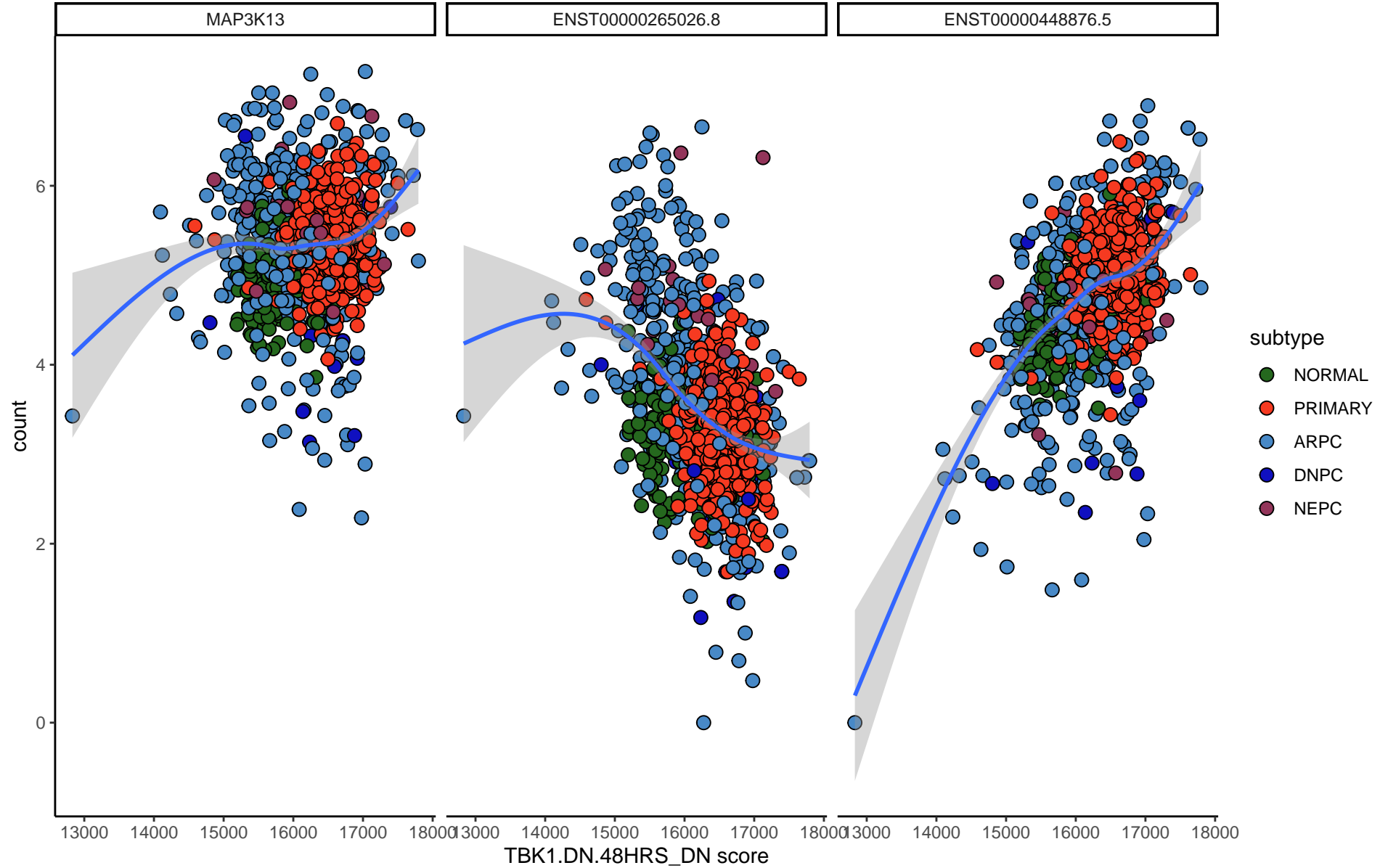
RB_DN.V1_UP

ENST00000265026.8 – MAP3K13–201 PCC = -0.64

ENST00000448876.5 – MAP3K13–212 PCC = 0.67



MAP3K13
Delta corr= 1.3
Gene corr= 0.11



TBK1.DN.48HRS_DN

ENST00000265026.8 – MAP3K13–201 PCC = -0.64

ENST00000448876.5 – MAP3K13–212 PCC = 0.66

