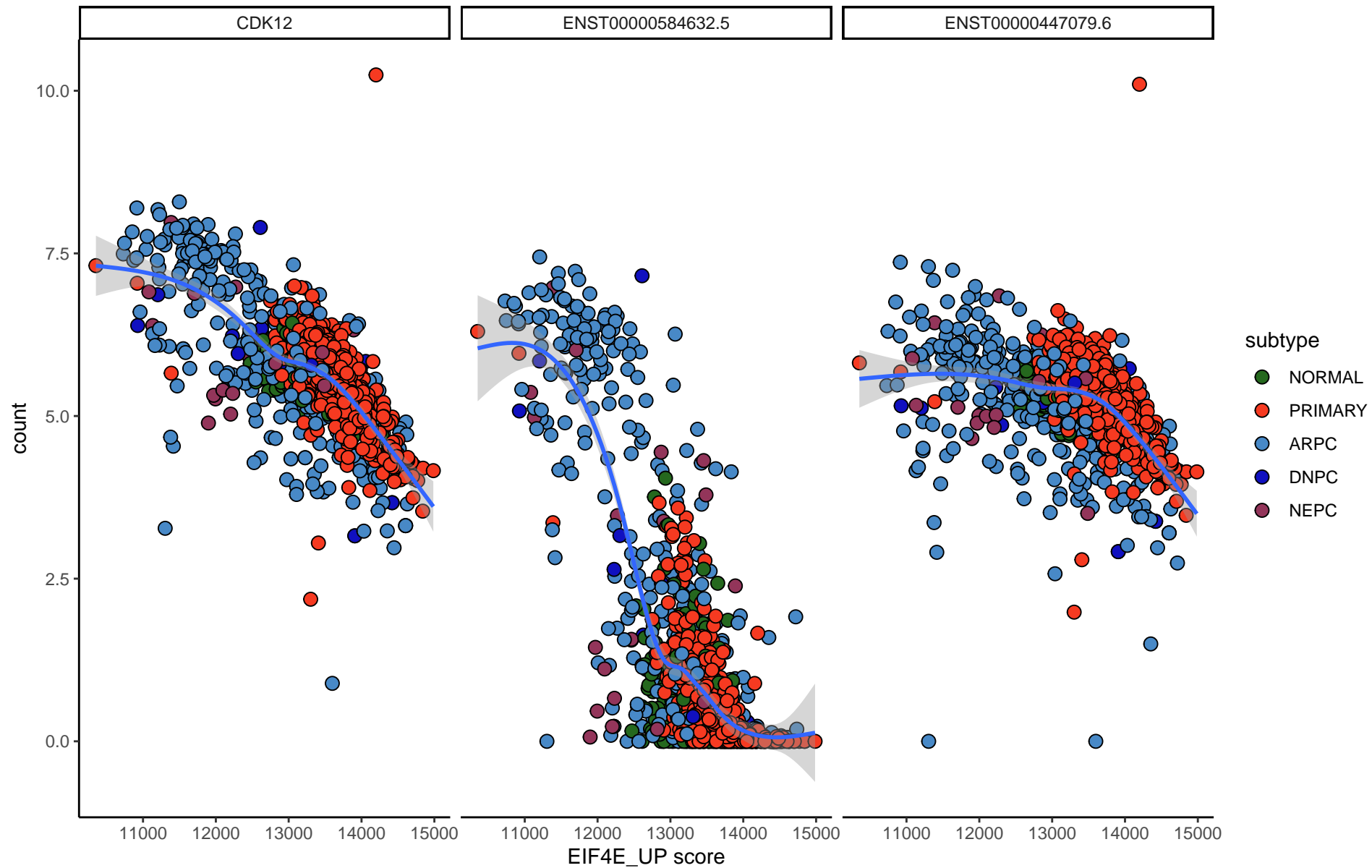


CDK12

Delta corr= 1.47

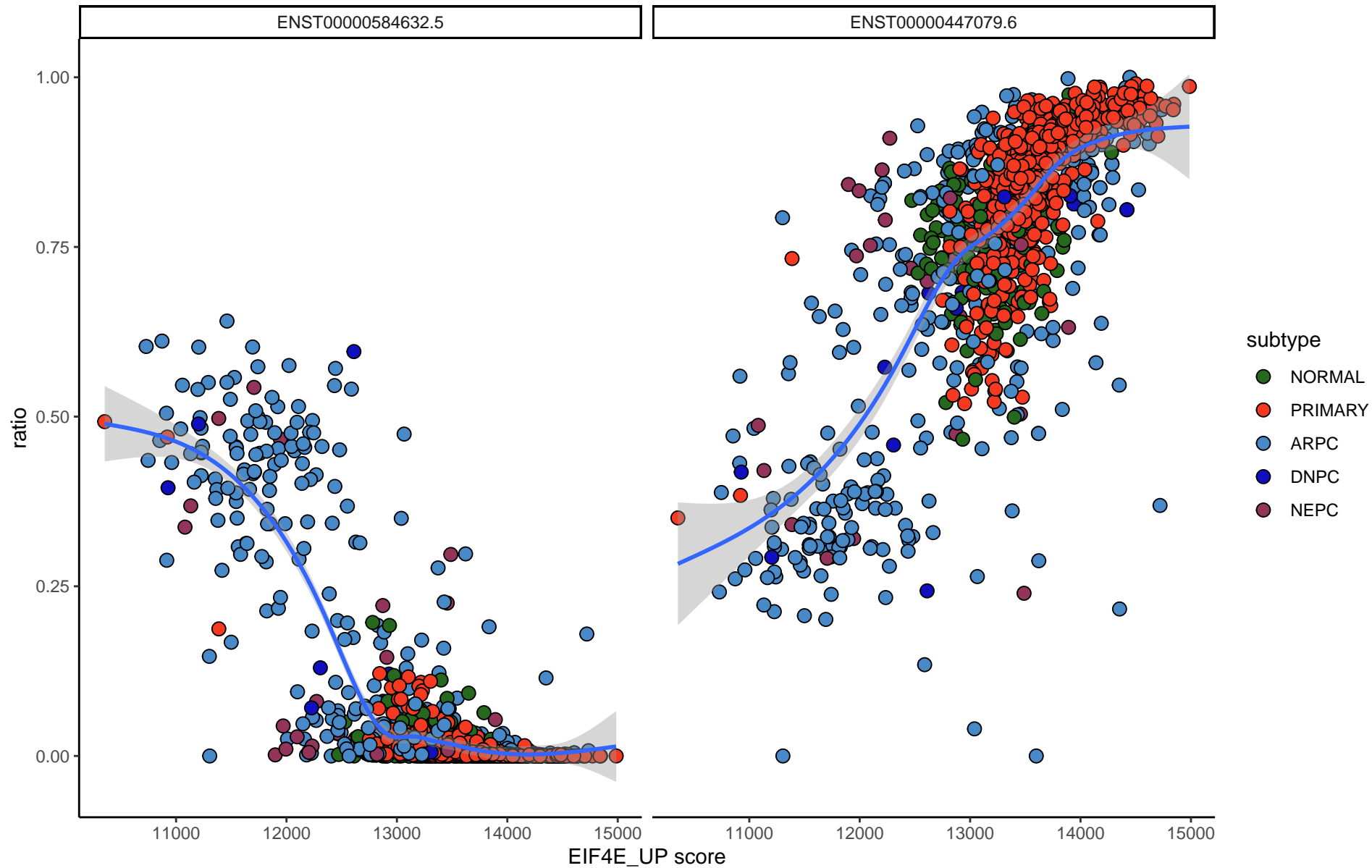
Gene corr= -0.68



EIF4E_UP

ENST00000584632.5 – CDK12–209 PCC = -0.74

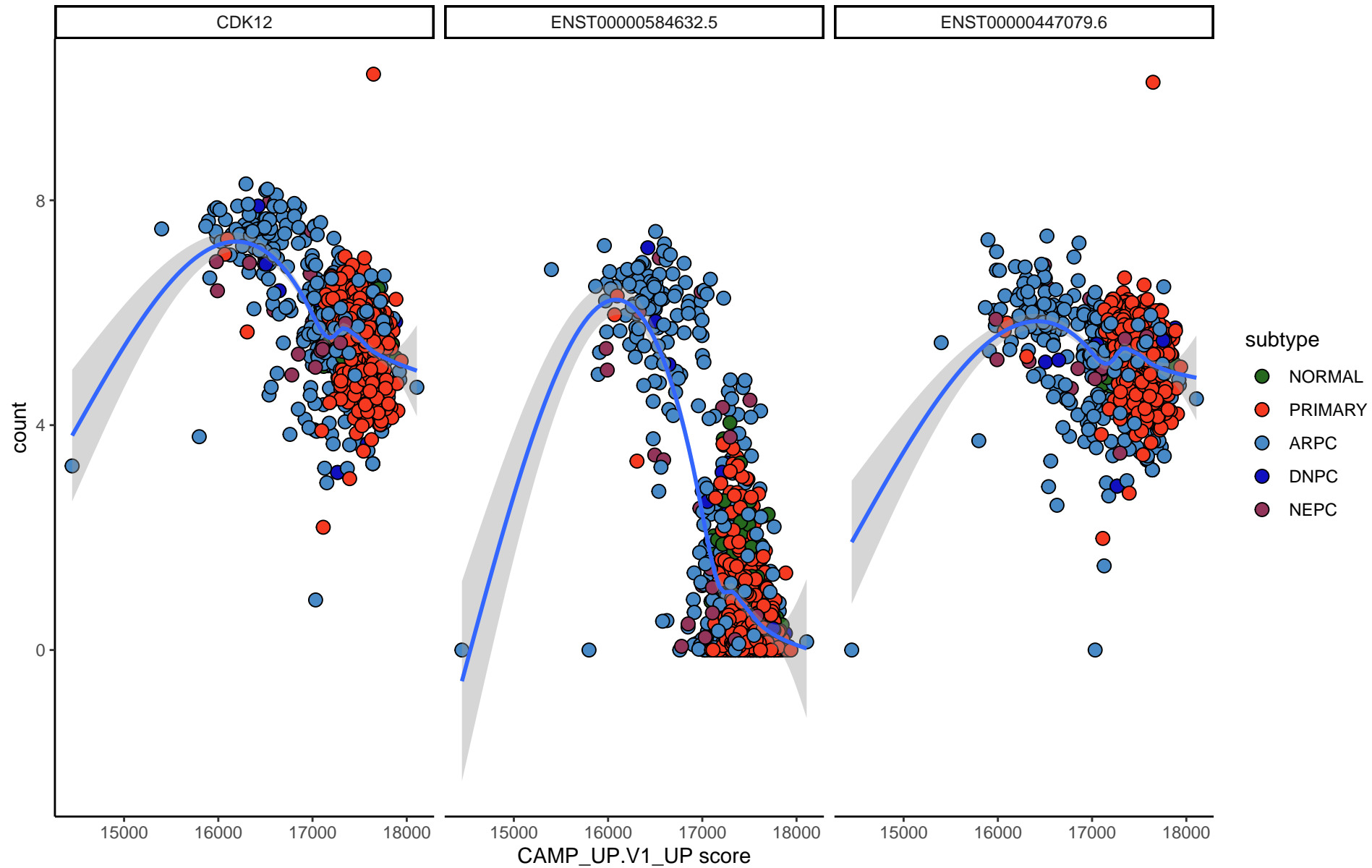
ENST00000447079.6 – CDK12–202 PCC = 0.73



CDK12

Delta corr= 1.39

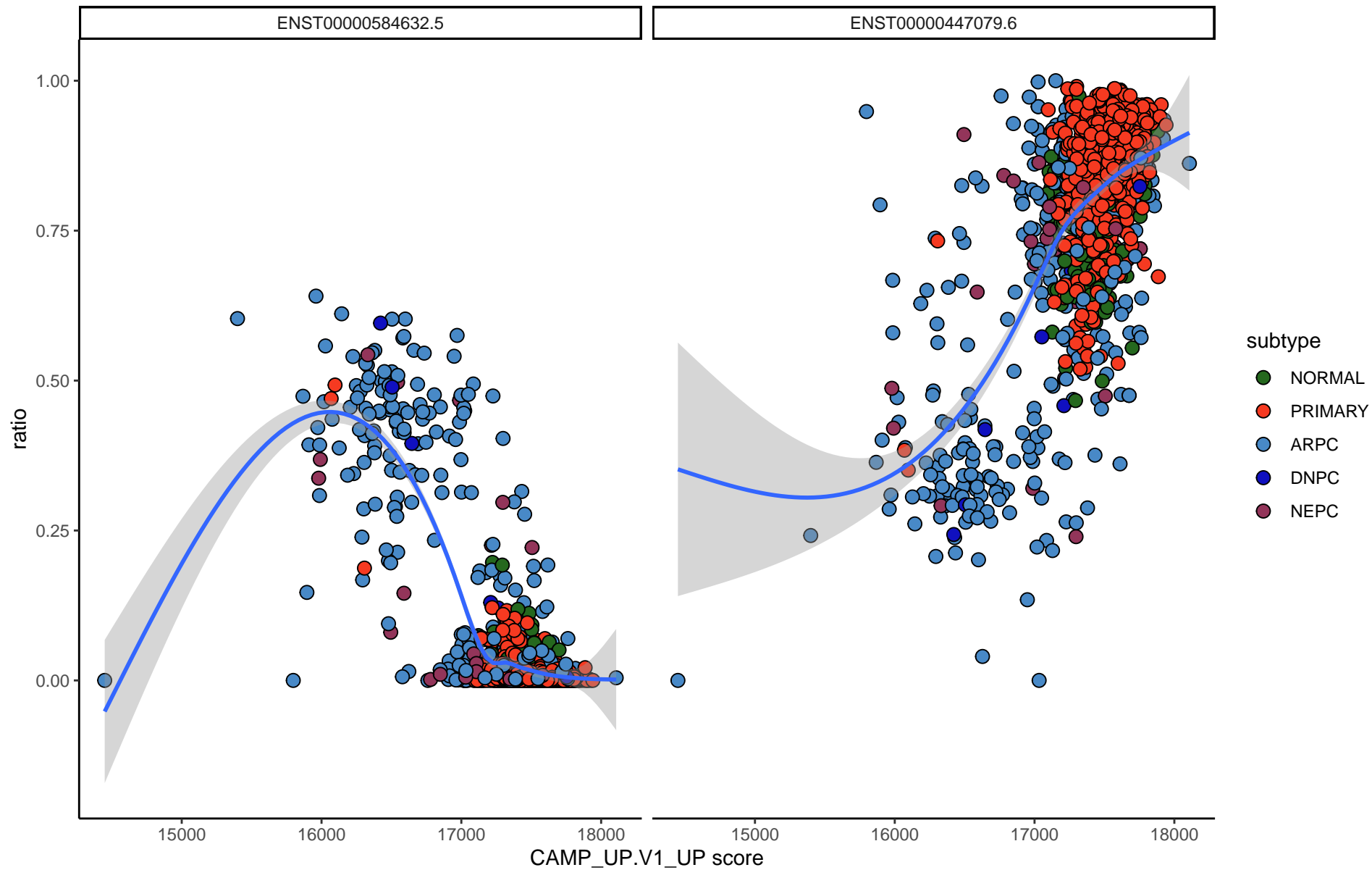
Gene corr= -0.51



CAMP_UP.V1_UP

ENST00000584632.5 – CDK12–209 PCC = -0.74

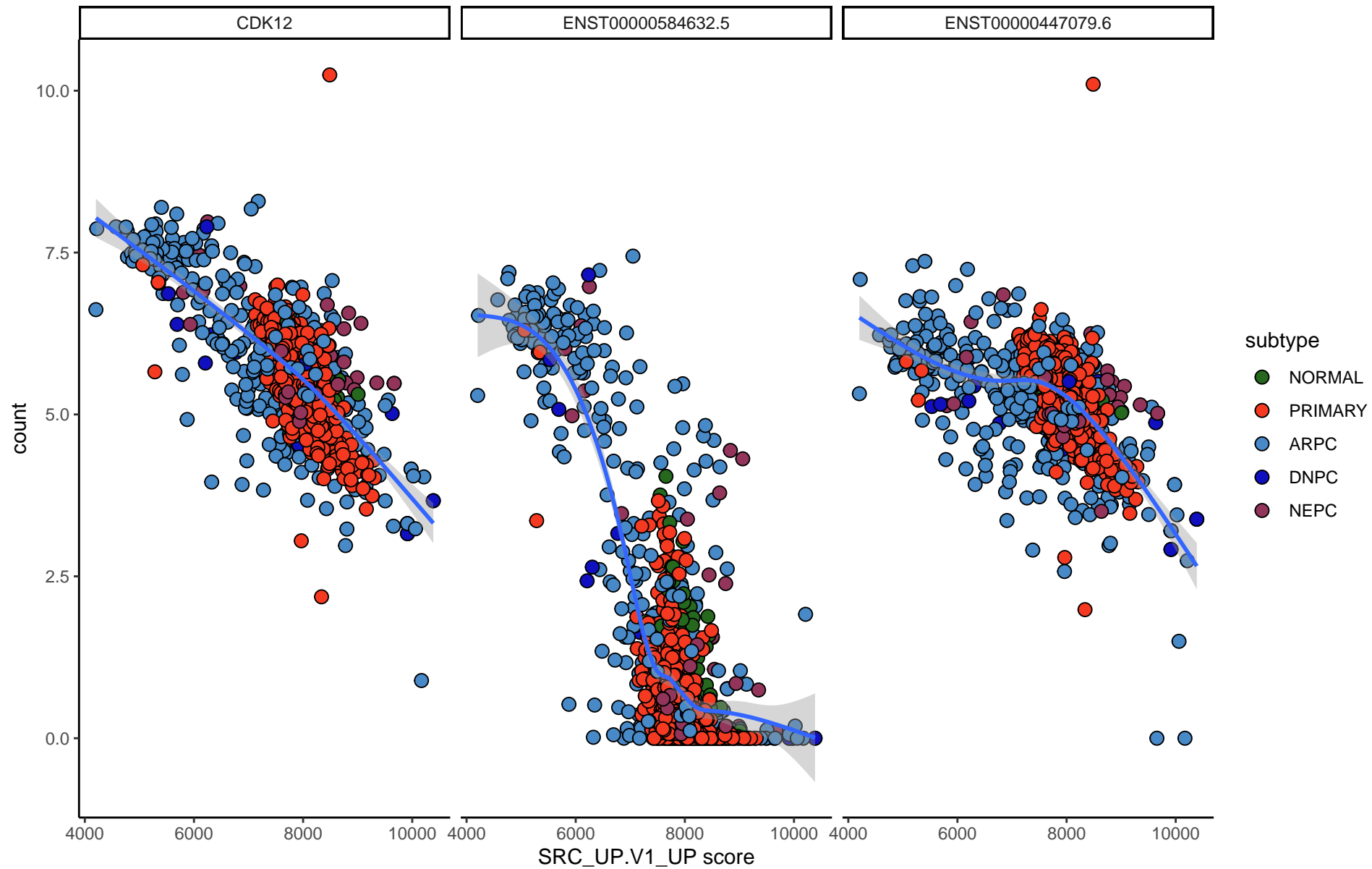
ENST00000447079.6 – CDK12–202 PCC = 0.65



CDK12

Delta corr= 1.36

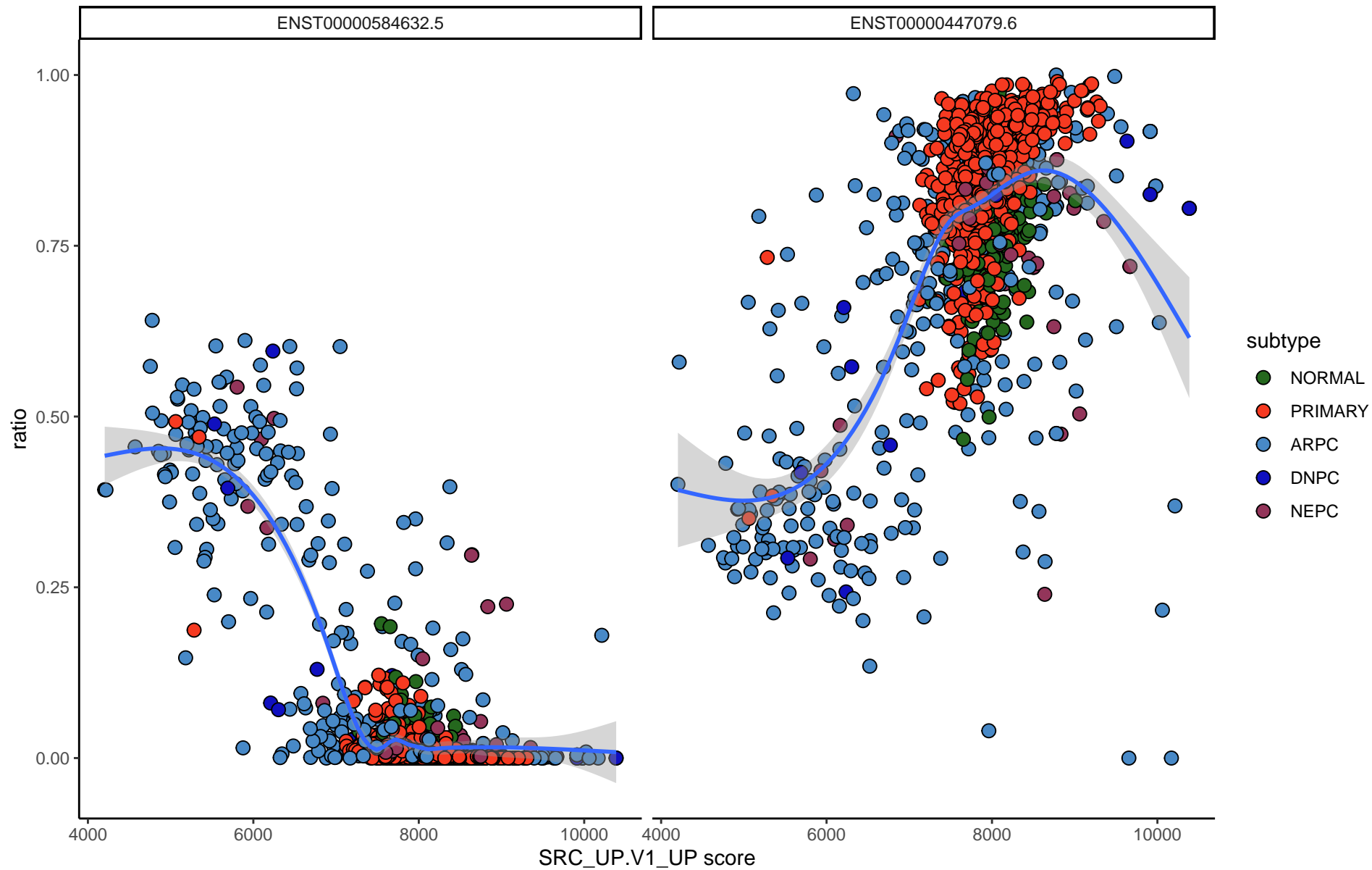
Gene corr= -0.73



SRC_UP.V1_UP

ENST00000584632.5 – CDK12–209 PCC = -0.76

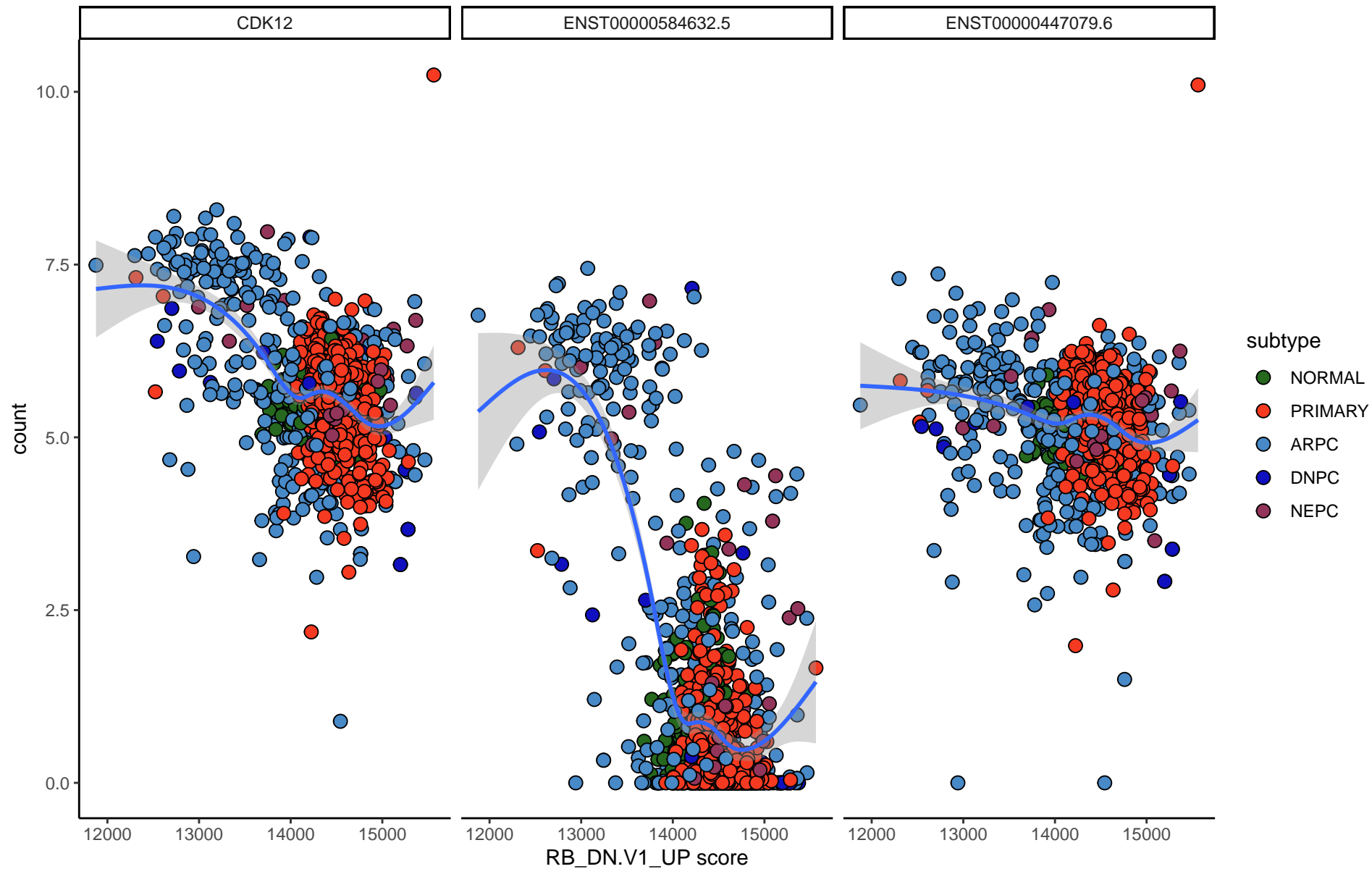
ENST00000447079.6 – CDK12–202 PCC = 0.6



CDK12

Delta corr= 1.31

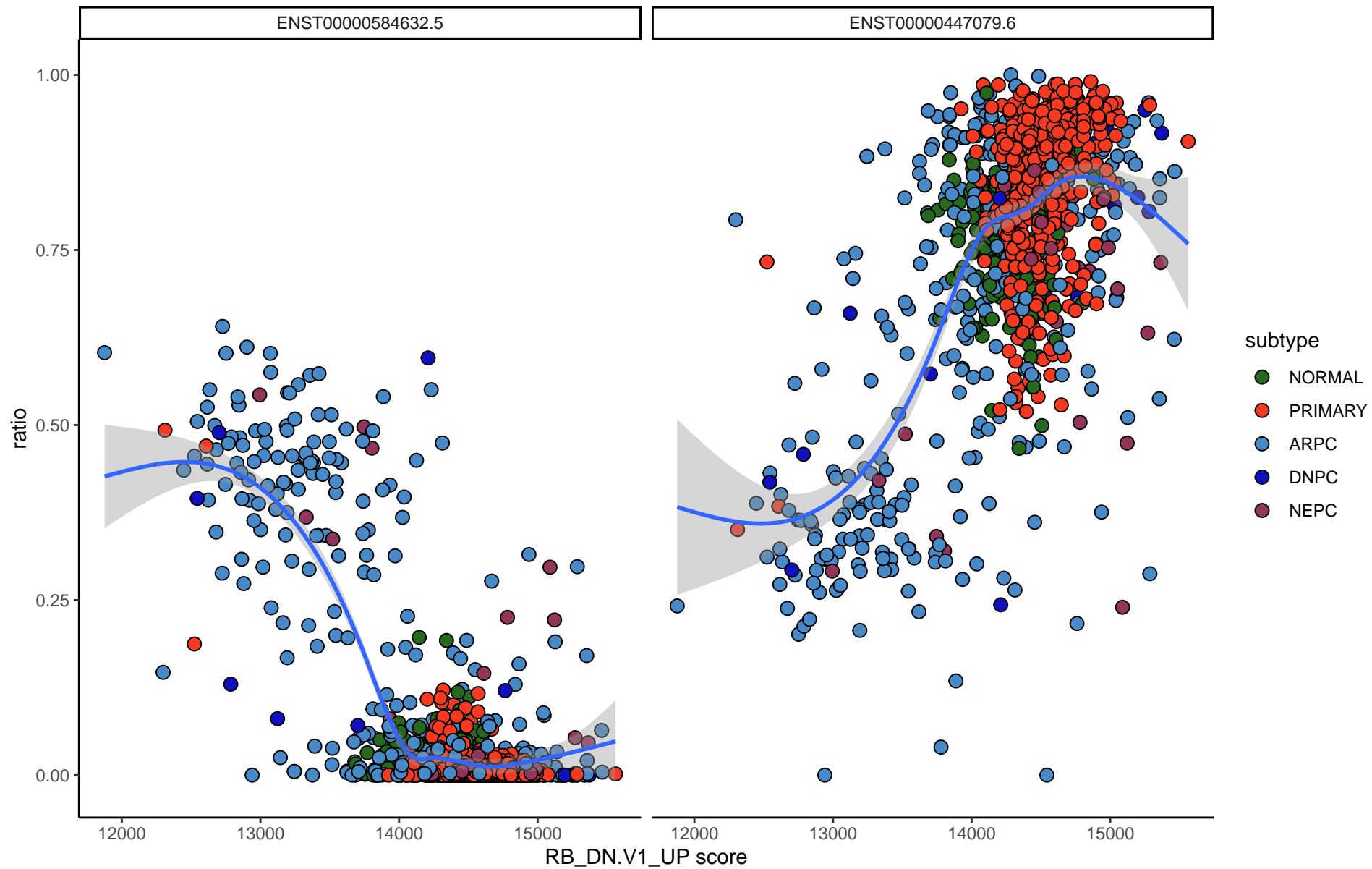
Gene corr= -0.47



RB_DN.V1_UP

ENST00000584632.5 – CDK12–209 PCC = -0.7

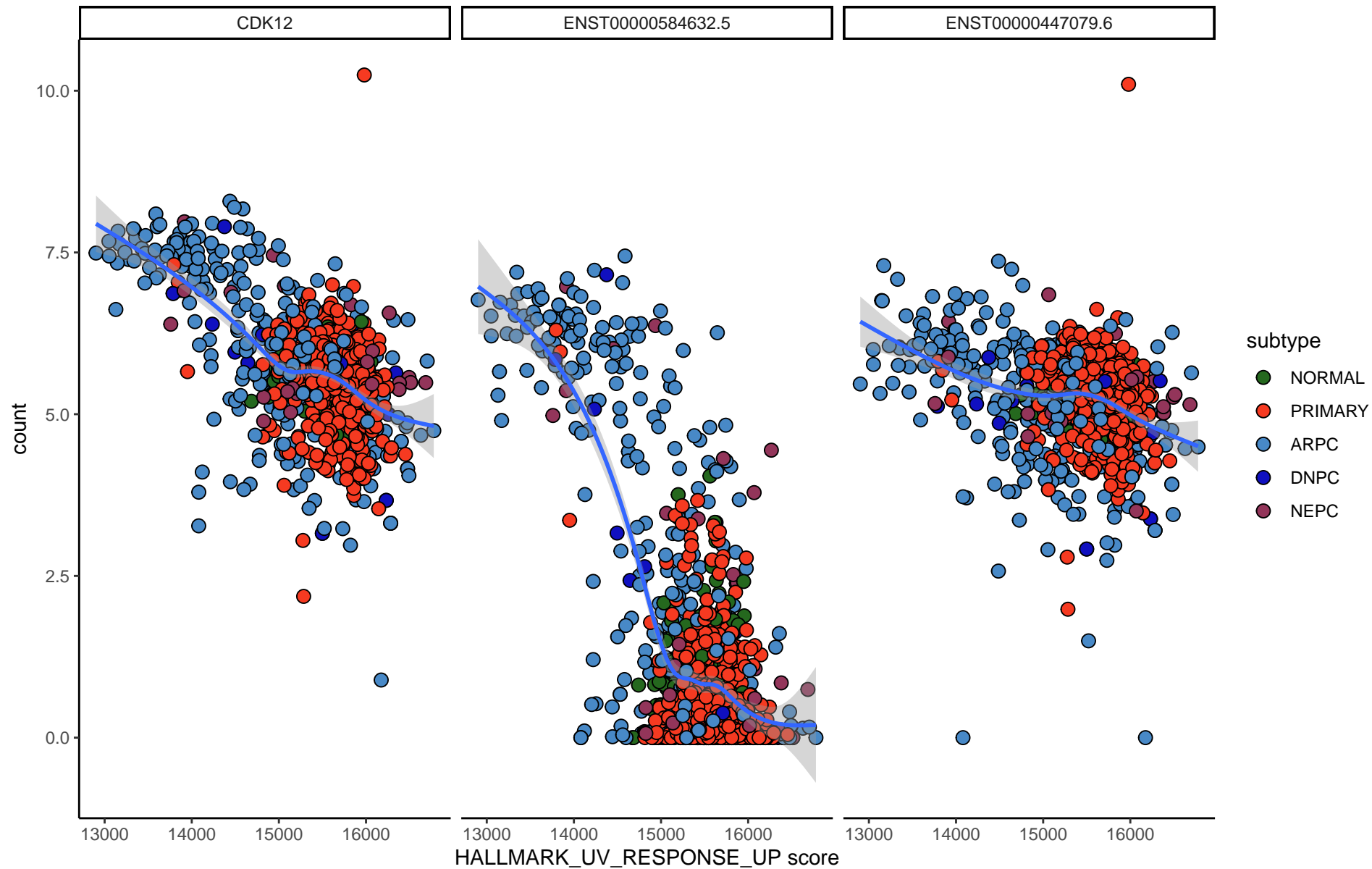
ENST00000447079.6 – CDK12–202 PCC = 0.61



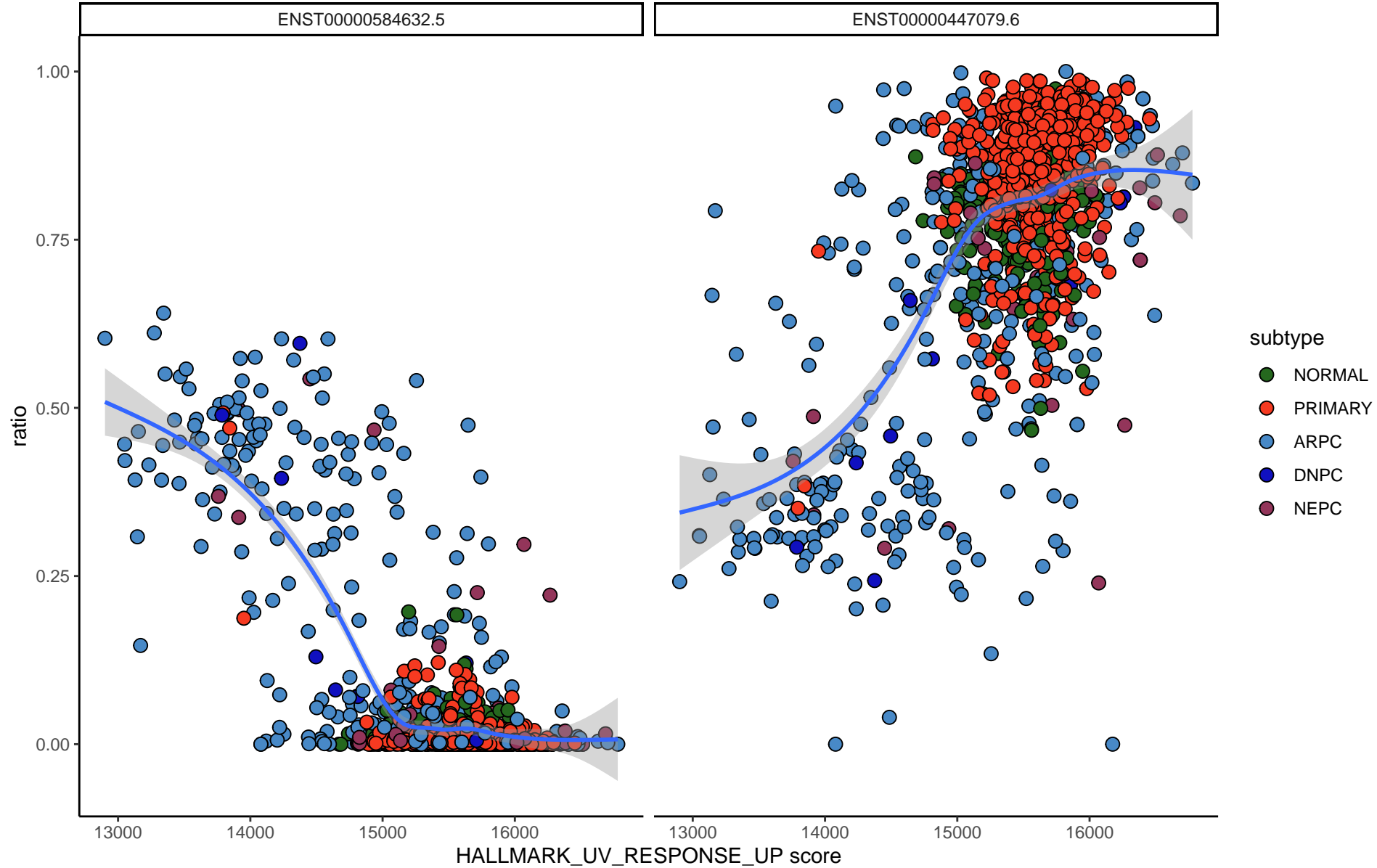
CDK12

Delta corr= 1.31

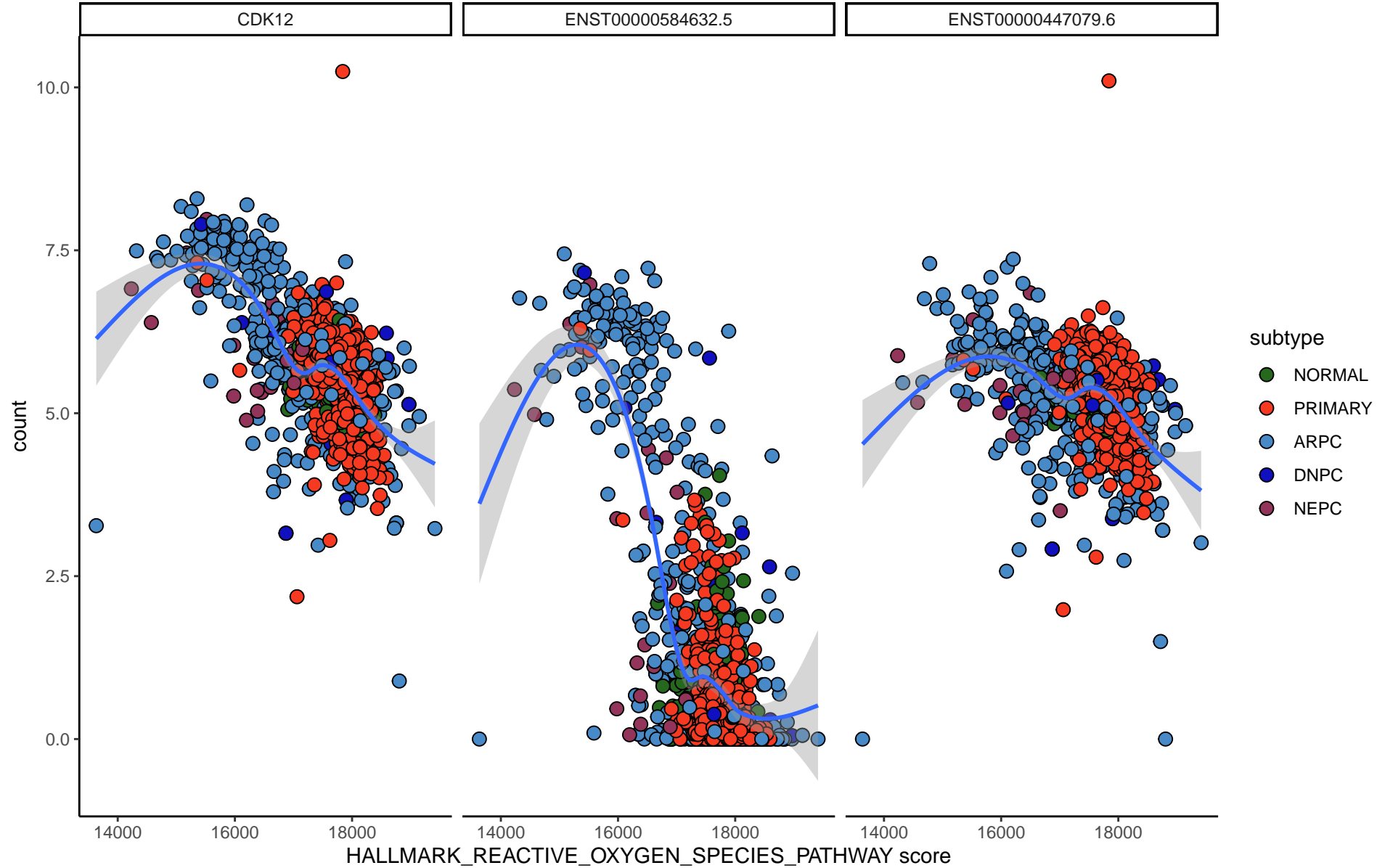
Gene corr= -0.55



HALLMARK_UV_RESPONSE_UP
ENST00000584632.5 – CDK12-209 PCC = -0.71
ENST00000447079.6 – CDK12-202 PCC = 0.59



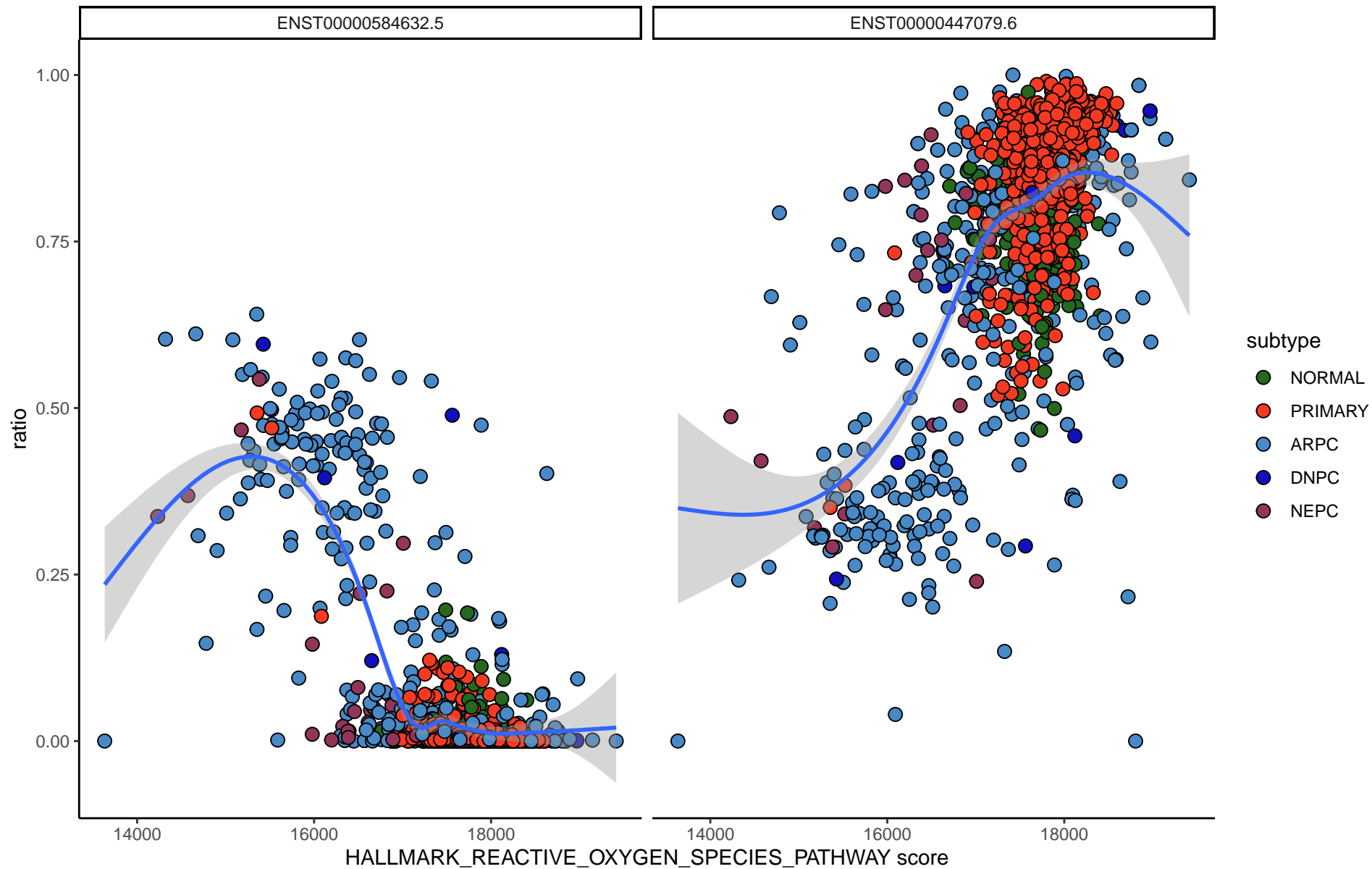
CDK12
Delta corr= 1.3
Gene corr= -0.6



HALLMARK_REACTIVE_OXYGEN_SPECIES_PATHWAY

ENST00000584632.5 – CDK12–209 PCC = -0.69

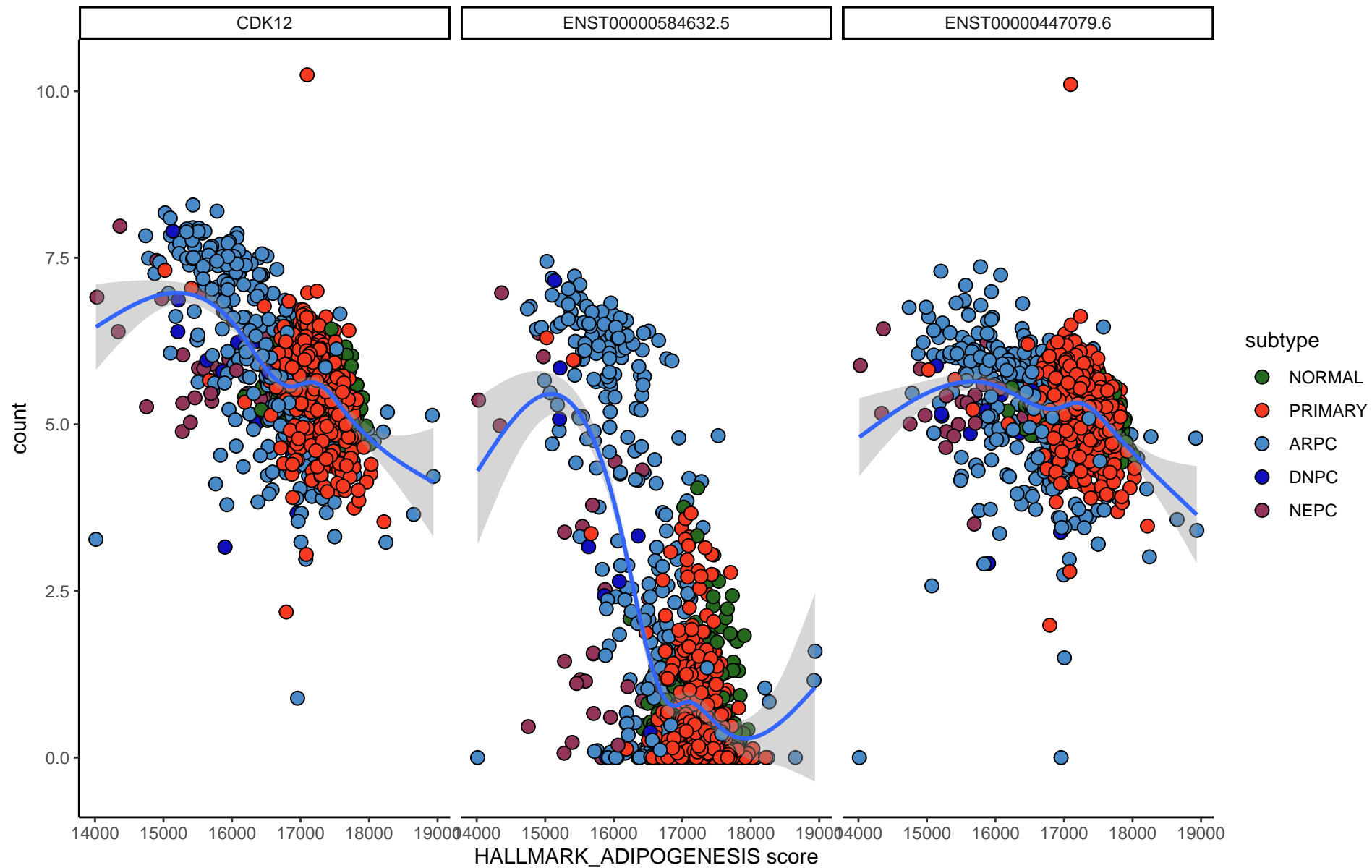
ENST00000447079.6 – CDK12–202 PCC = 0.61



CDK12

Delta corr= 1.28

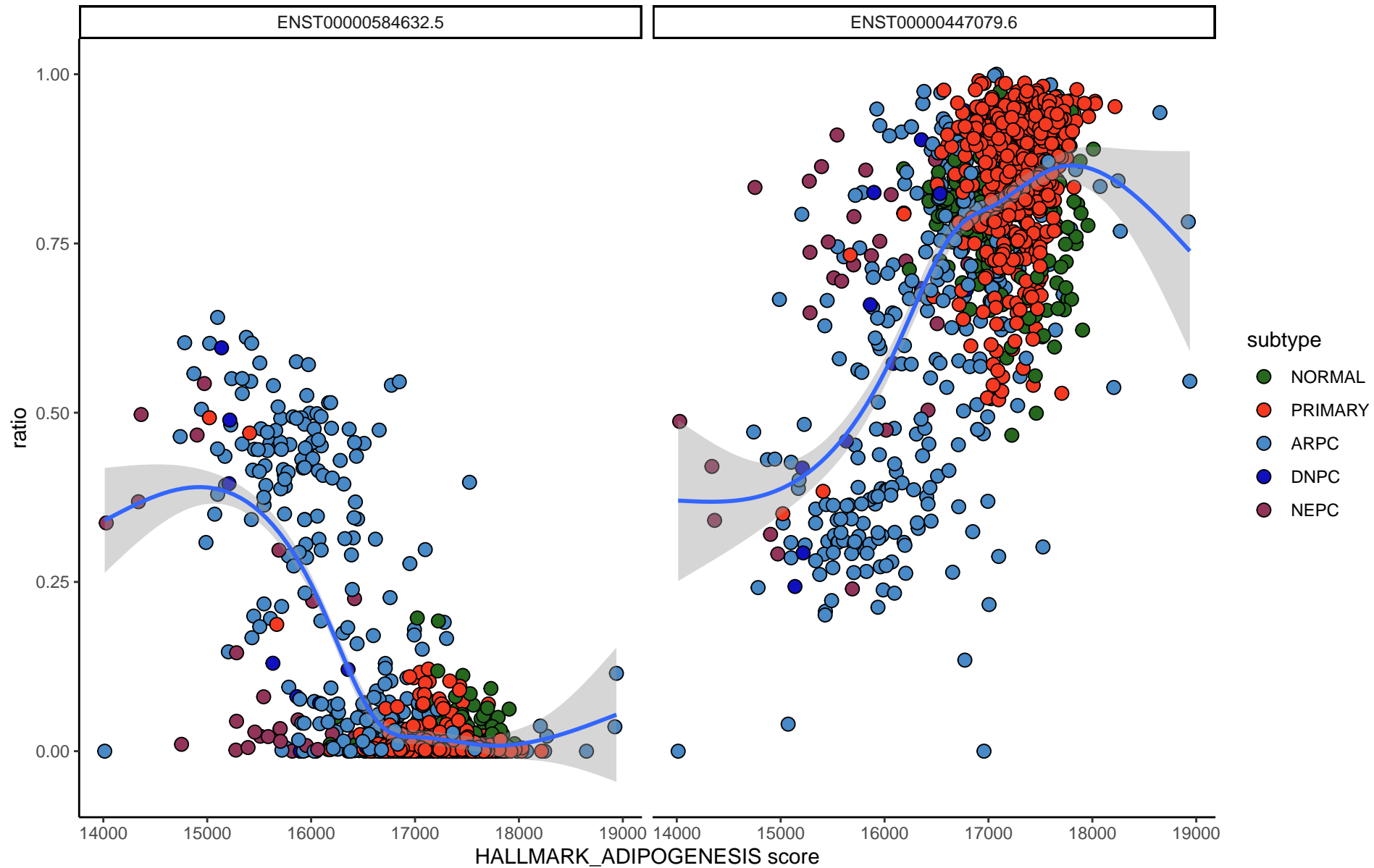
Gene corr= -0.51



HALLMARK_ADIPOGENESIS

ENST00000584632.5 – CDK12–209 PCC = -0.67

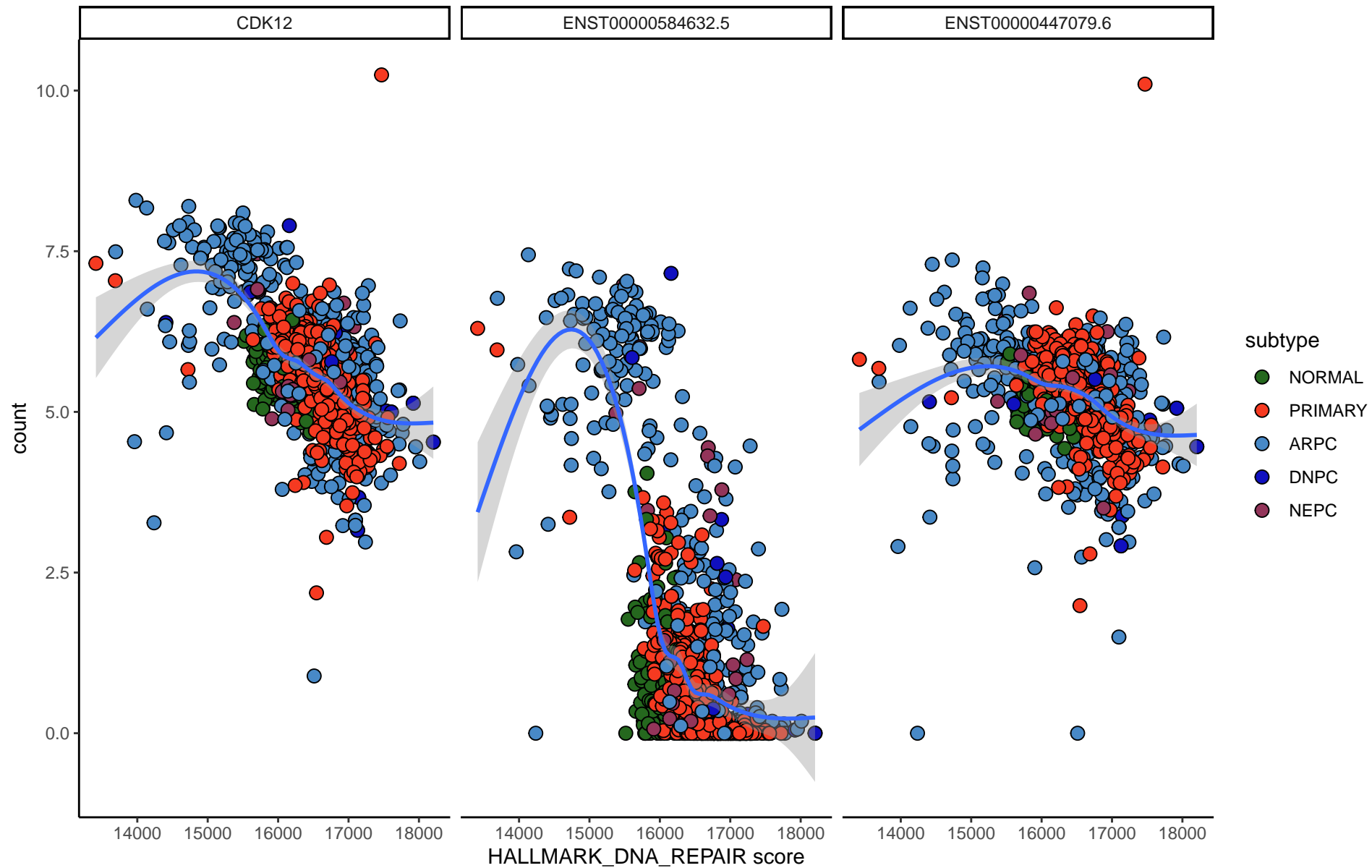
ENST00000447079.6 – CDK12–202 PCC = 0.62



CDK12

Delta corr= 1.28

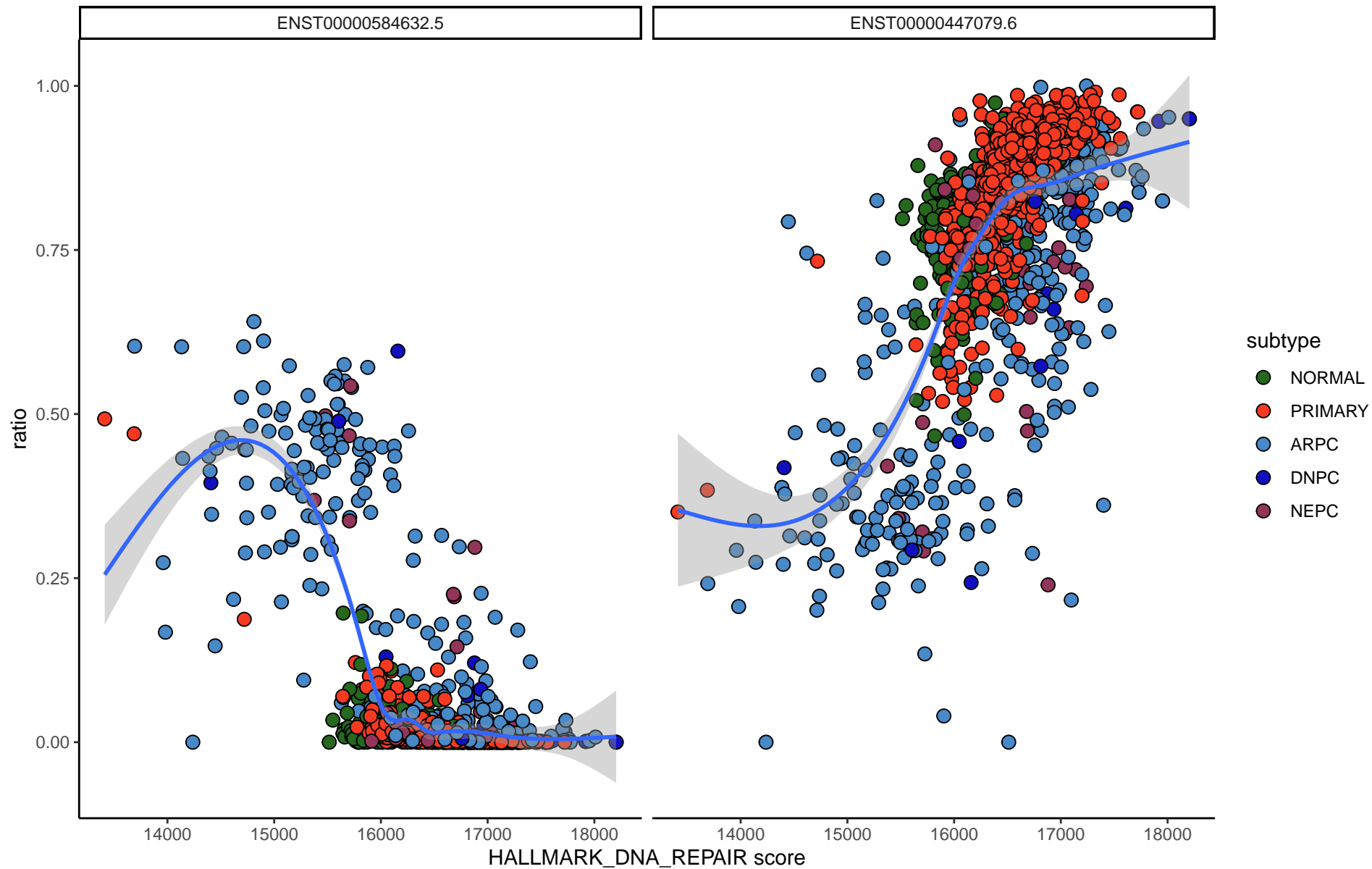
Gene corr= -0.58



HALLMARK_DNA_REPAIR

ENST00000584632.5 – CDK12–209 PCC = -0.64

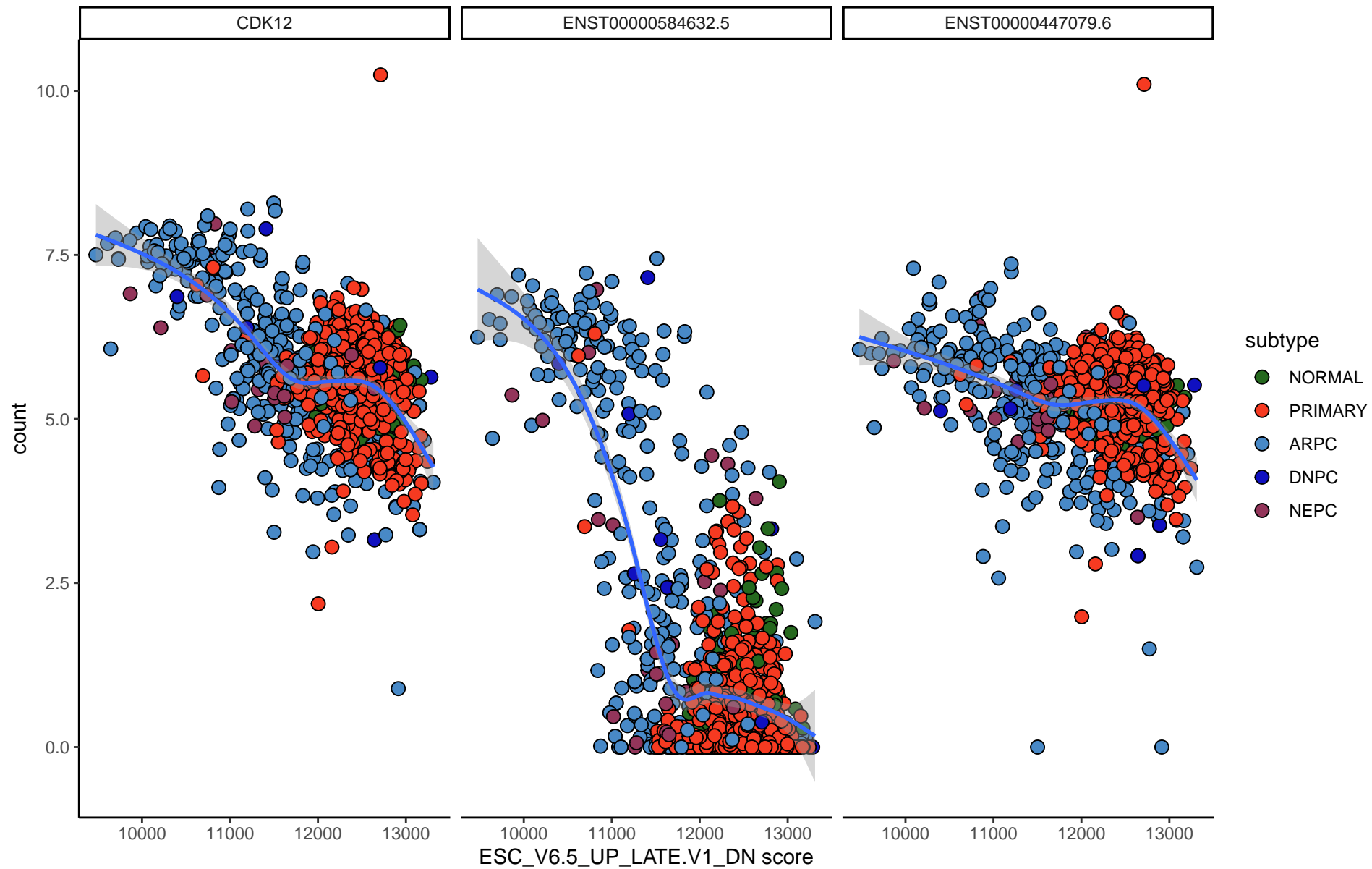
ENST00000447079.6 – CDK12–202 PCC = 0.64



CDK12

Delta corr= 1.28

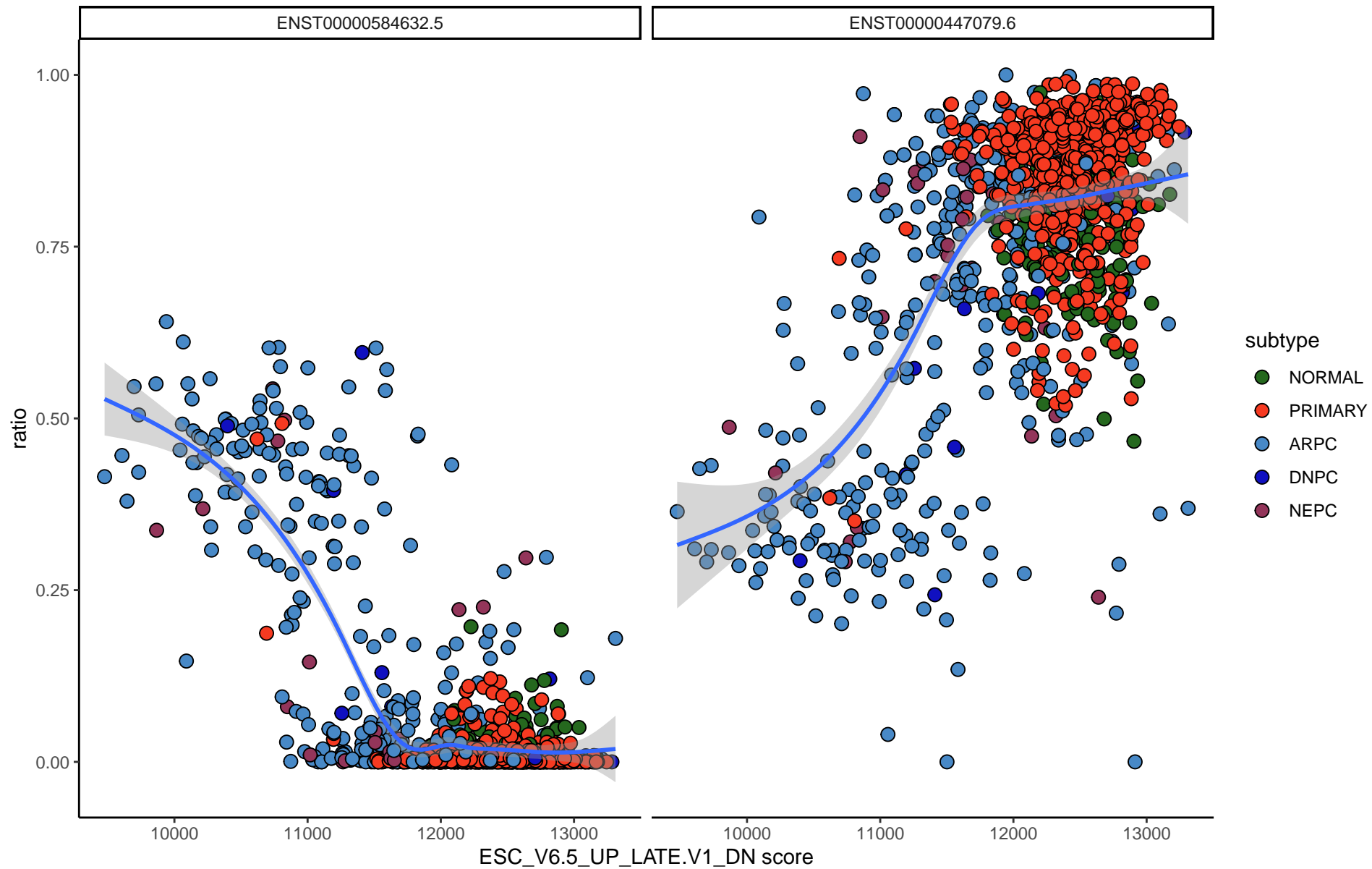
Gene corr= -0.54



ESC_V6.5_UP_LATE.V1_DN

ENST00000584632.5 – CDK12-209 PCC = -0.7

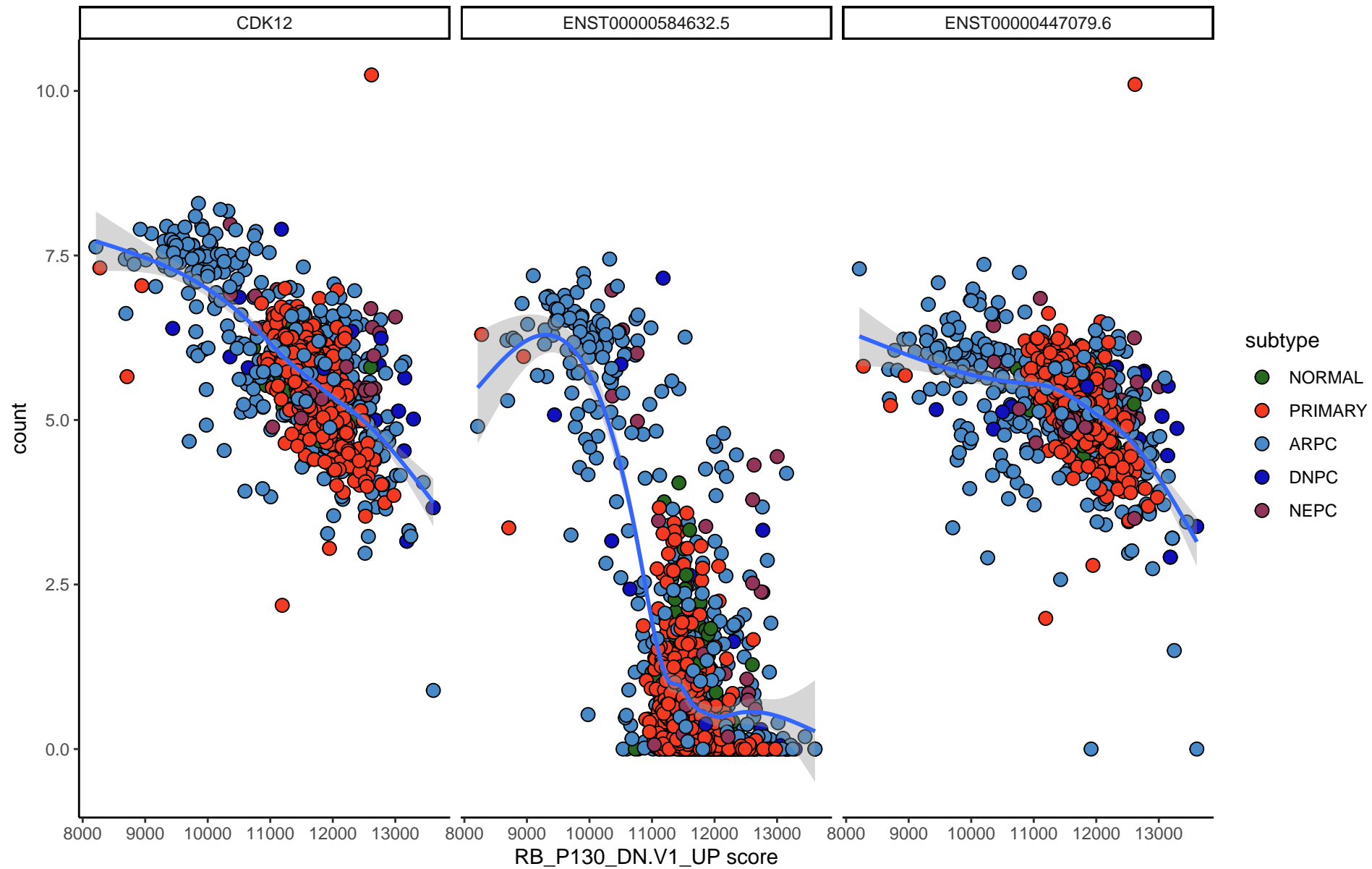
ENST00000447079.6 – CDK12-202 PCC = 0.58



CDK12

Delta corr= 1.27

Gene corr= -0.67



RB_P130_DN.V1_UP

ENST00000584632.5 – CDK12-209 PCC = -0.7

ENST00000447079.6 – CDK12-202 PCC = 0.57

