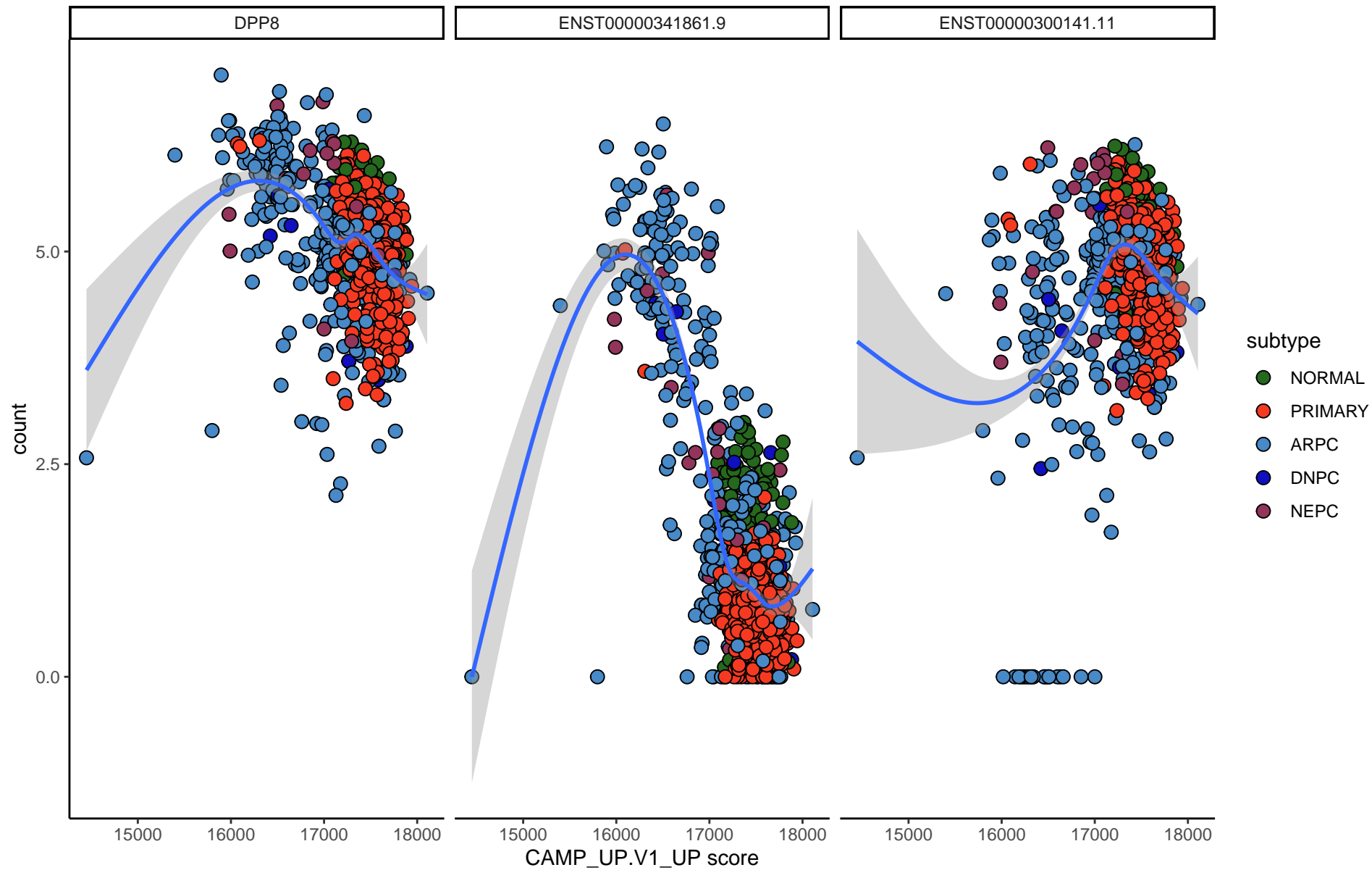


DPP8

Delta corr= 1.44

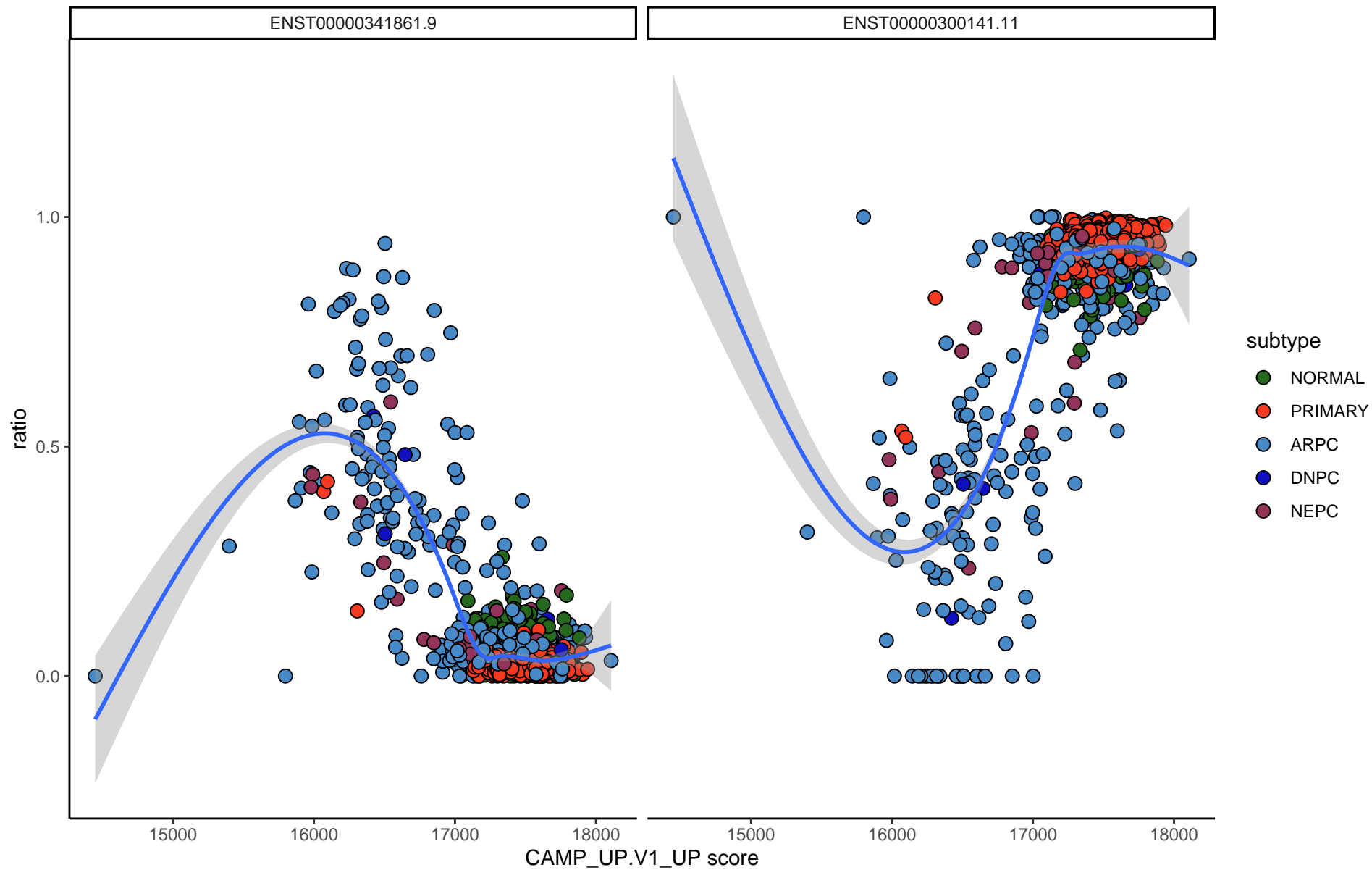
Gene corr= -0.36



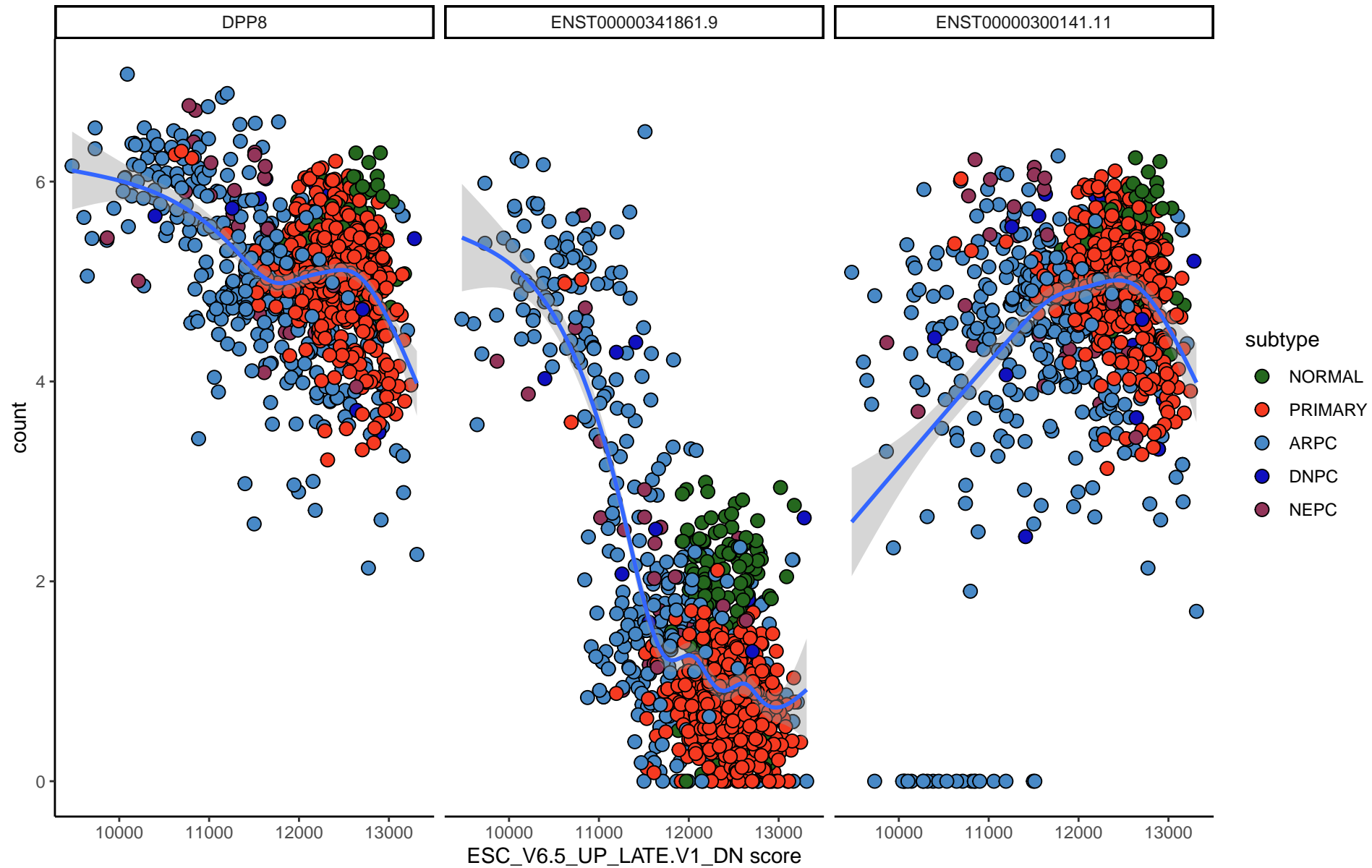
CAMP_UP.V1_UP

ENST00000341861.9 – DPP8–203 PCC = -0.71

ENST00000300141.11 – DPP8–201 PCC = 0.73



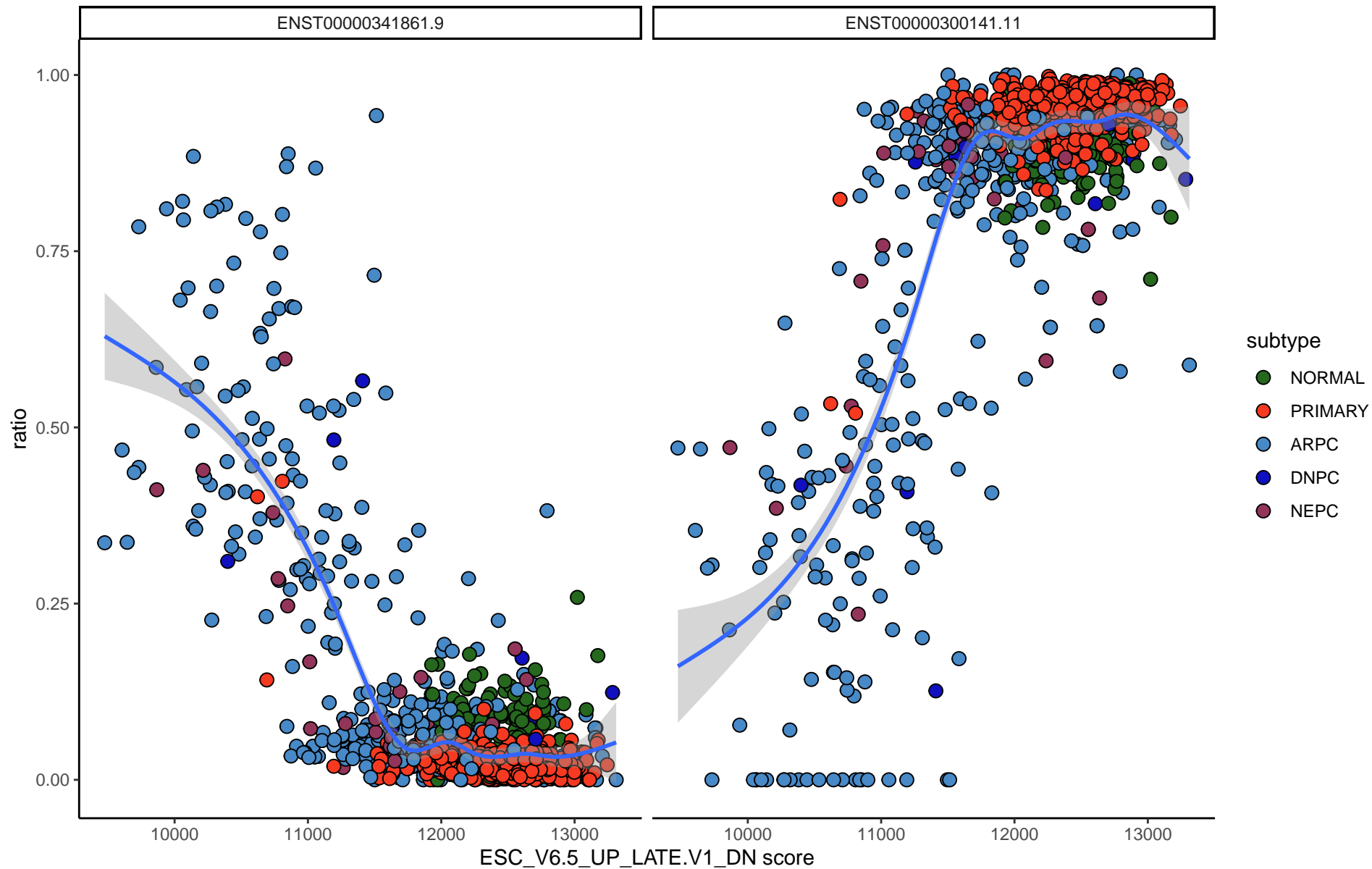
DPP8
Delta corr= 1.41
Gene corr= -0.35



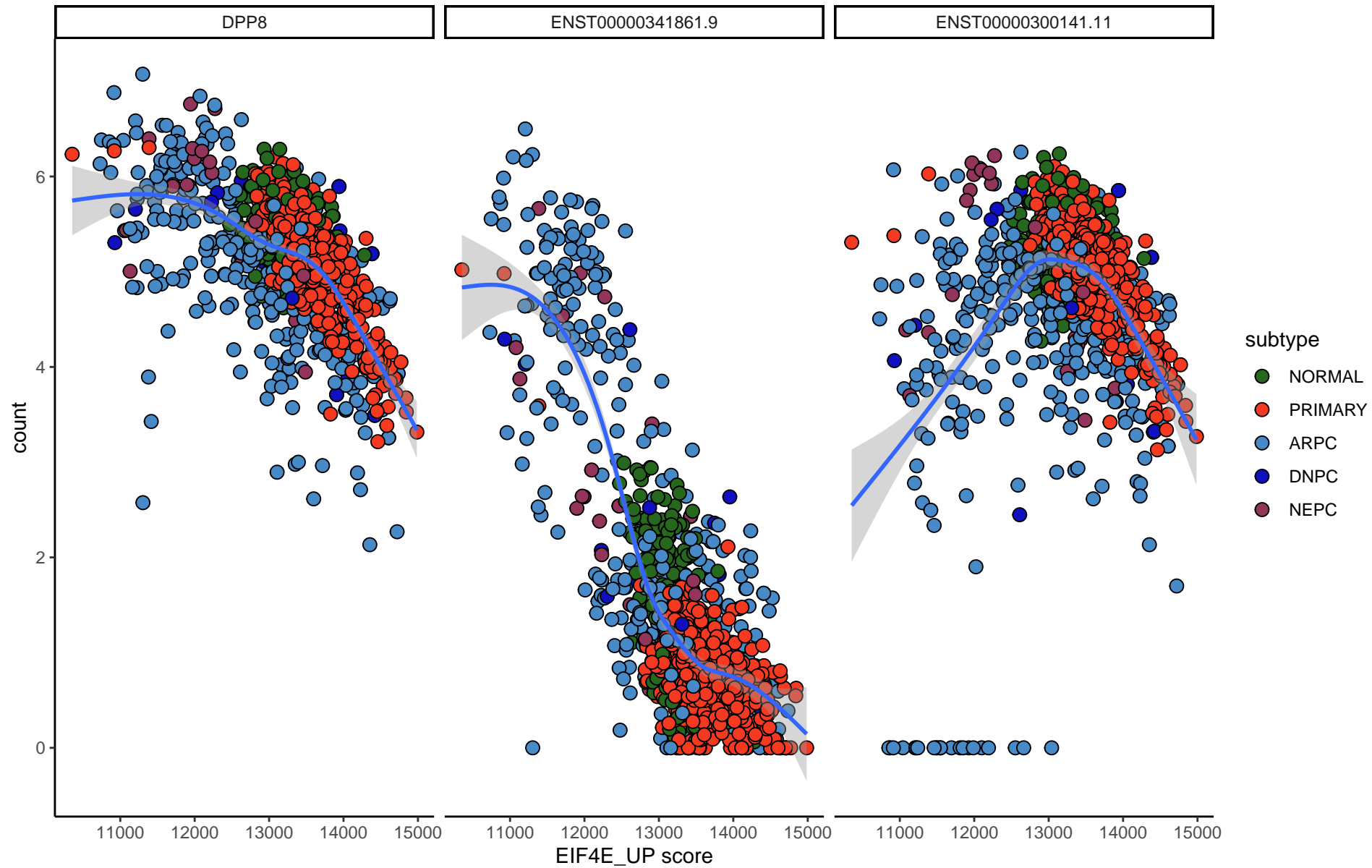
ESC_V6.5_UP_LATE.V1_DN

ENST00000341861.9 – DPP8–203 PCC = -0.7

ENST00000300141.11 – DPP8–201 PCC = 0.72



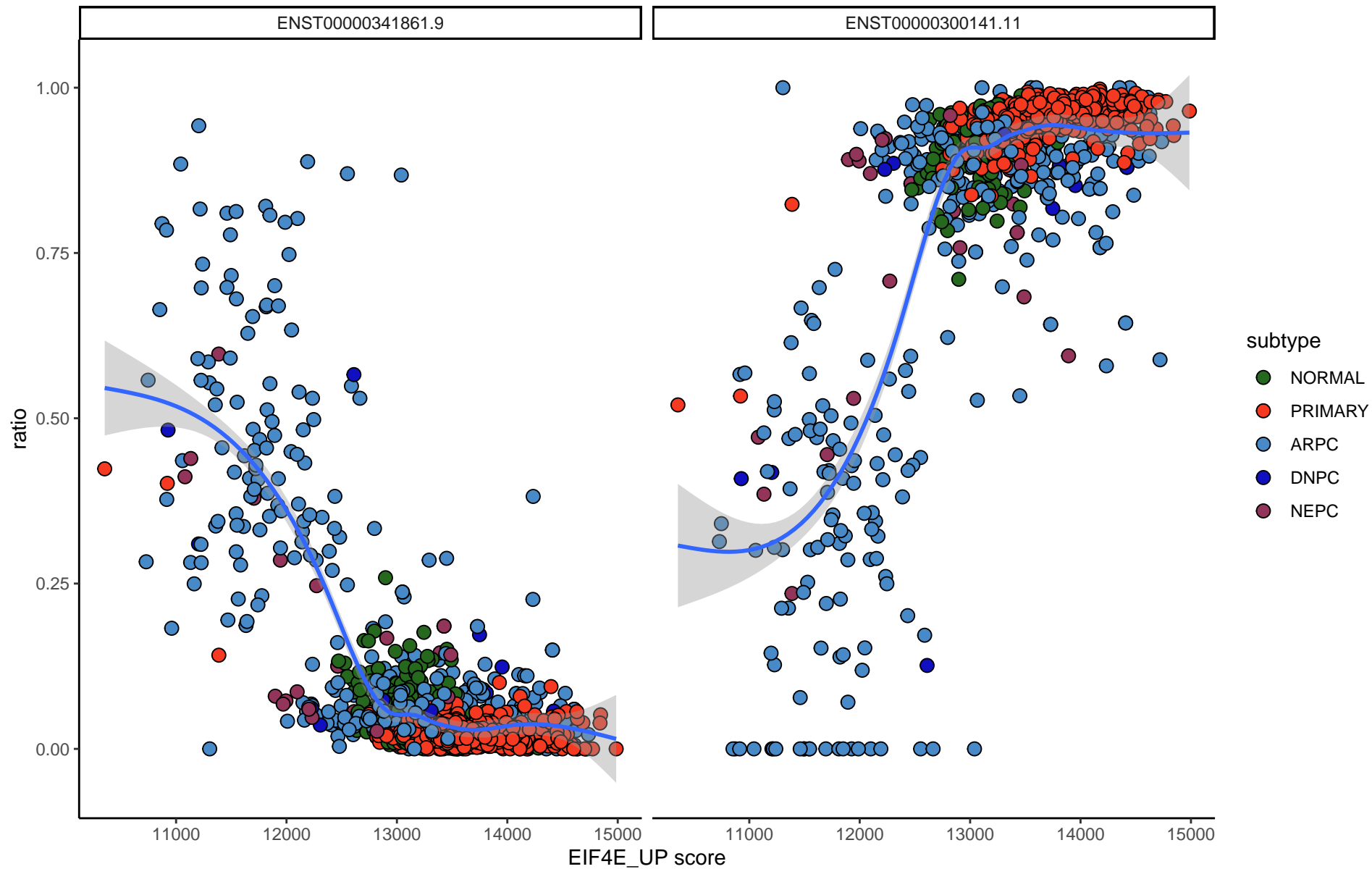
DPP8
Delta corr= 1.39
Gene corr= -0.58



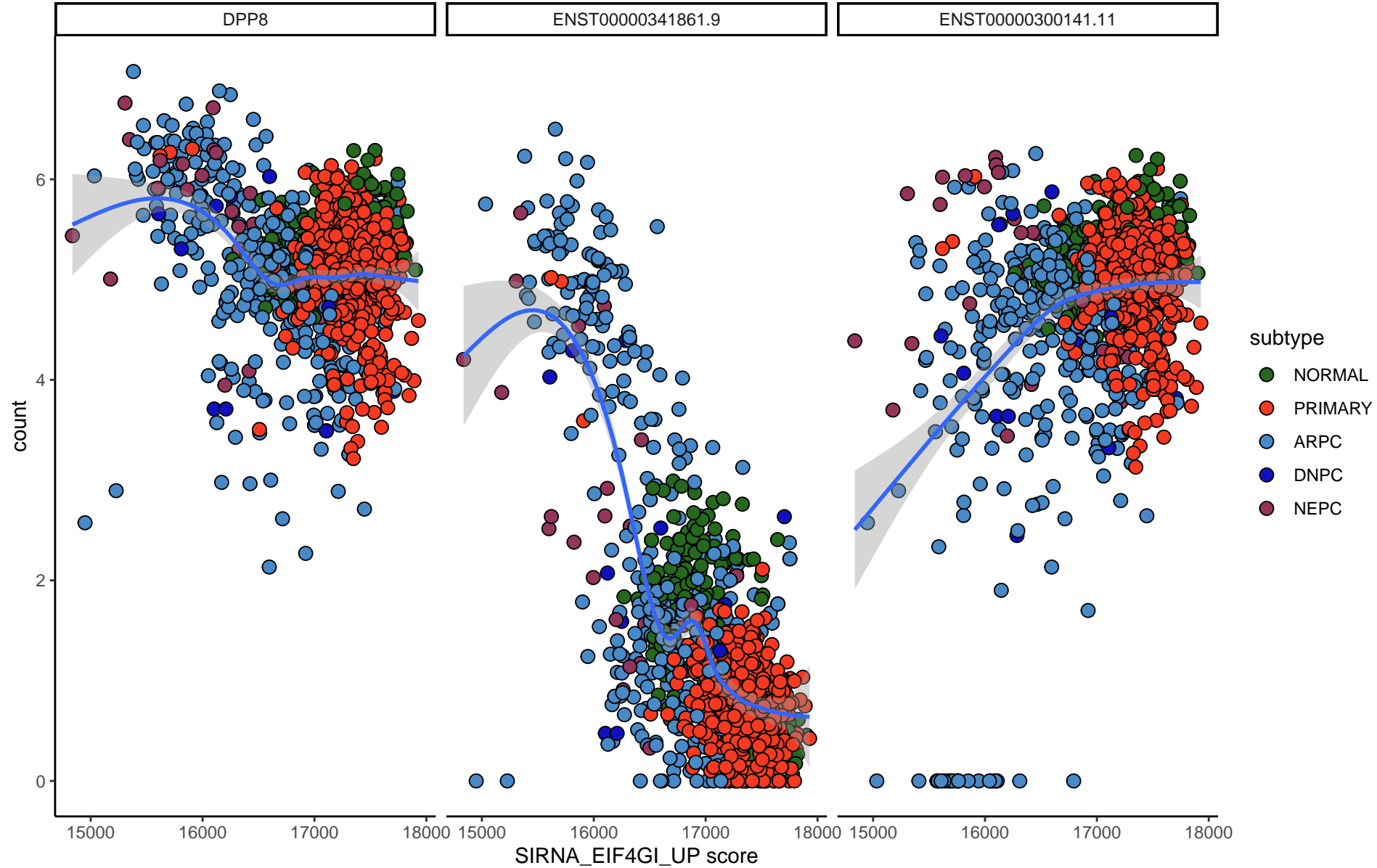
EIF4E_UP

ENST00000341861.9 – DPP8–203 PCC = -0.69

ENST00000300141.11 – DPP8–201 PCC = 0.7



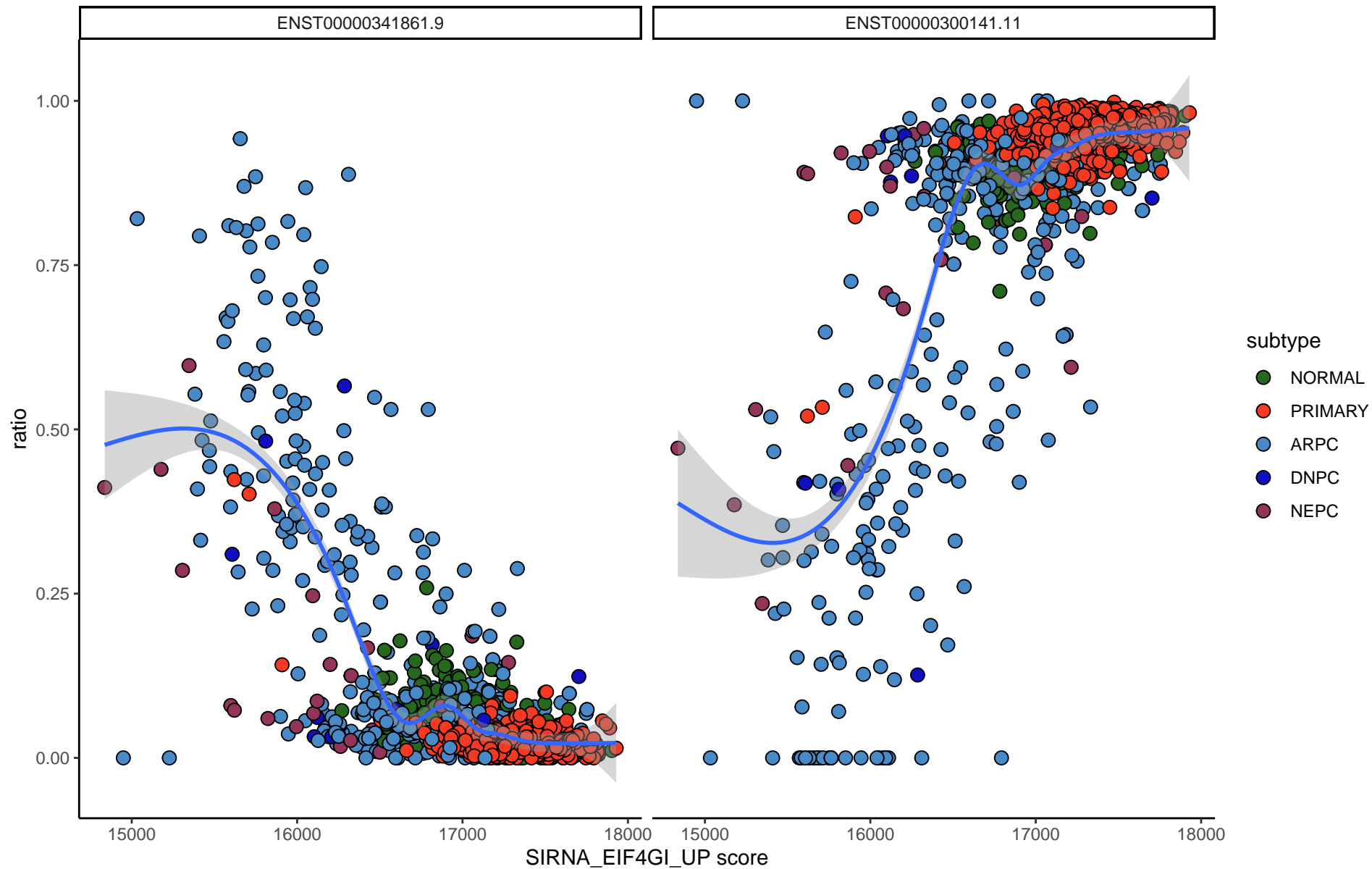
DPP8
Delta corr= 1.38
Gene corr= -0.25



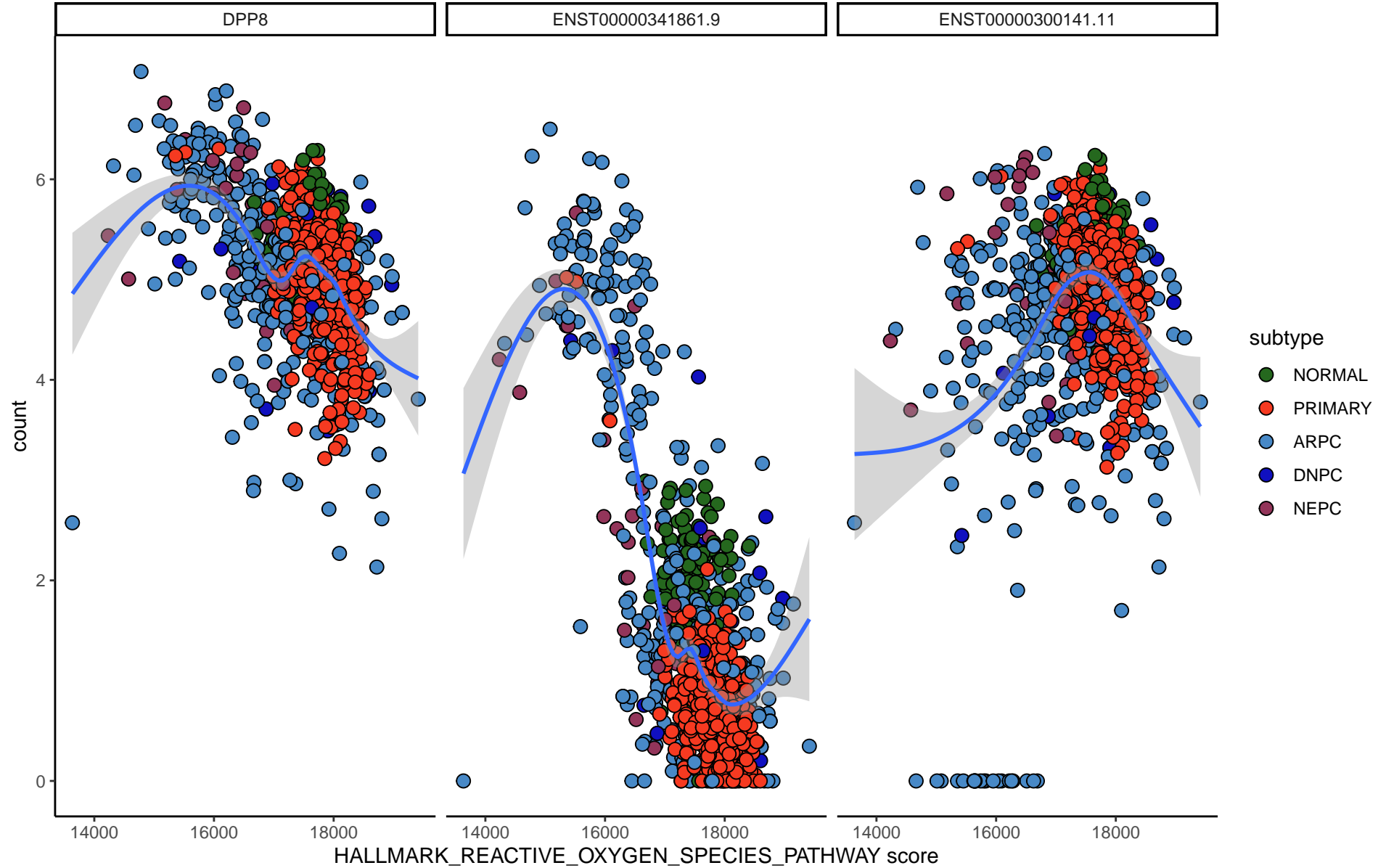
SIRNA_EIF4GI_UP

ENST00000341861.9 – DPP8–203 PCC = -0.69

ENST00000300141.11 – DPP8–201 PCC = 0.69



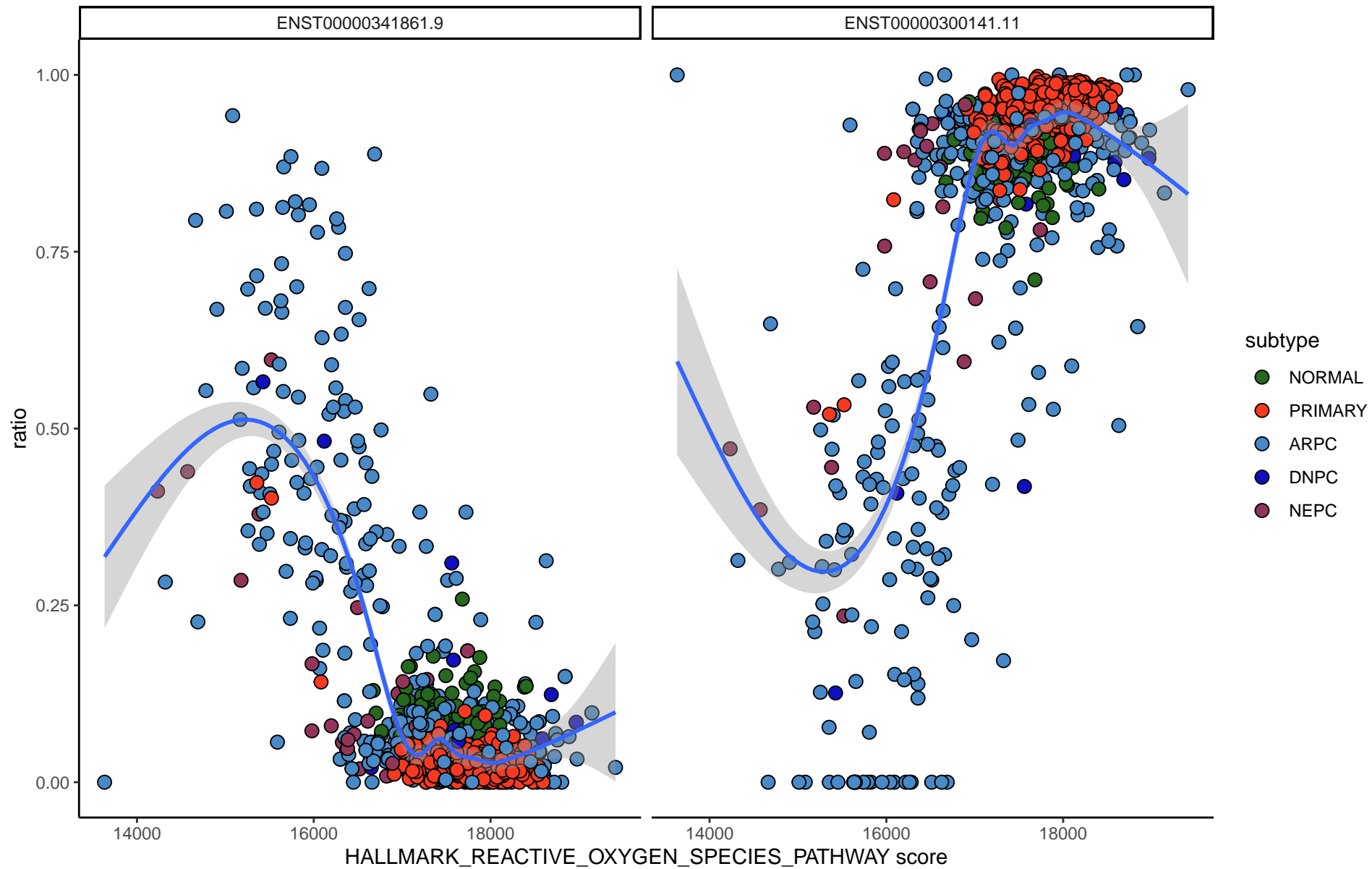
DPP8
Delta corr= 1.37
Gene corr= -0.42



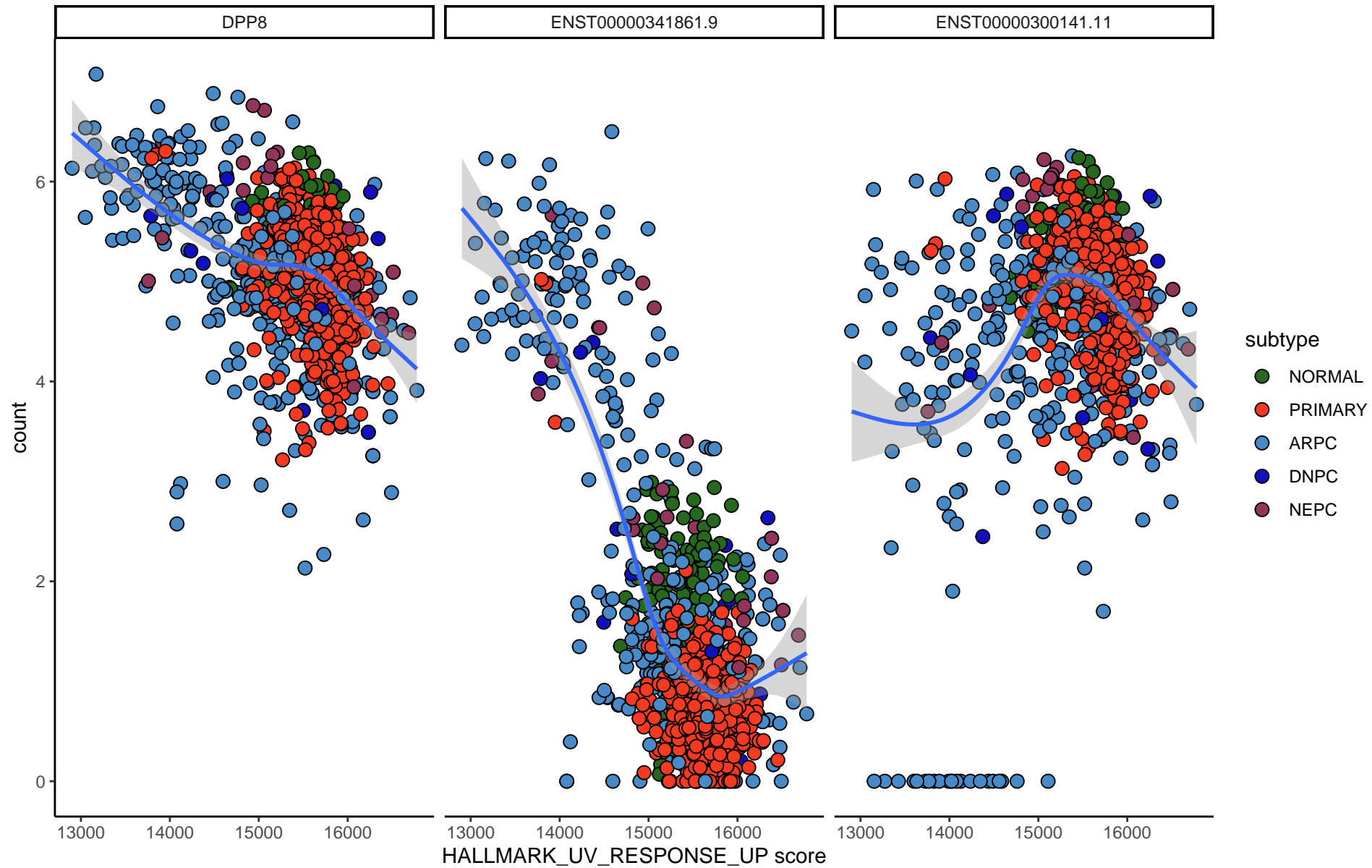
HALLMARK_REACTIVE_OXYGEN_SPECIES_PATHWAY

ENST00000341861.9 – DPP8–203 PCC = -0.68

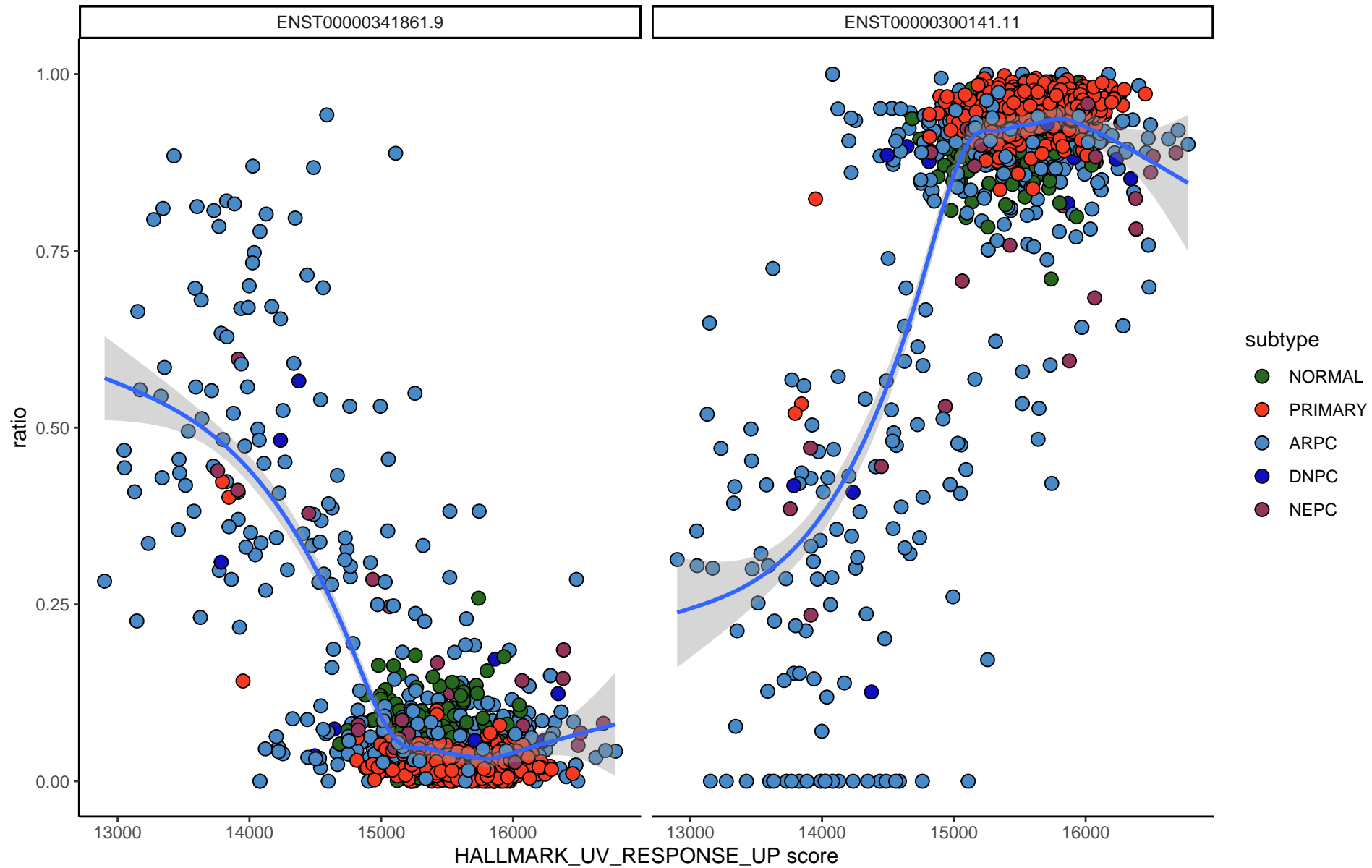
ENST00000300141.11 – DPP8–201 PCC = 0.69



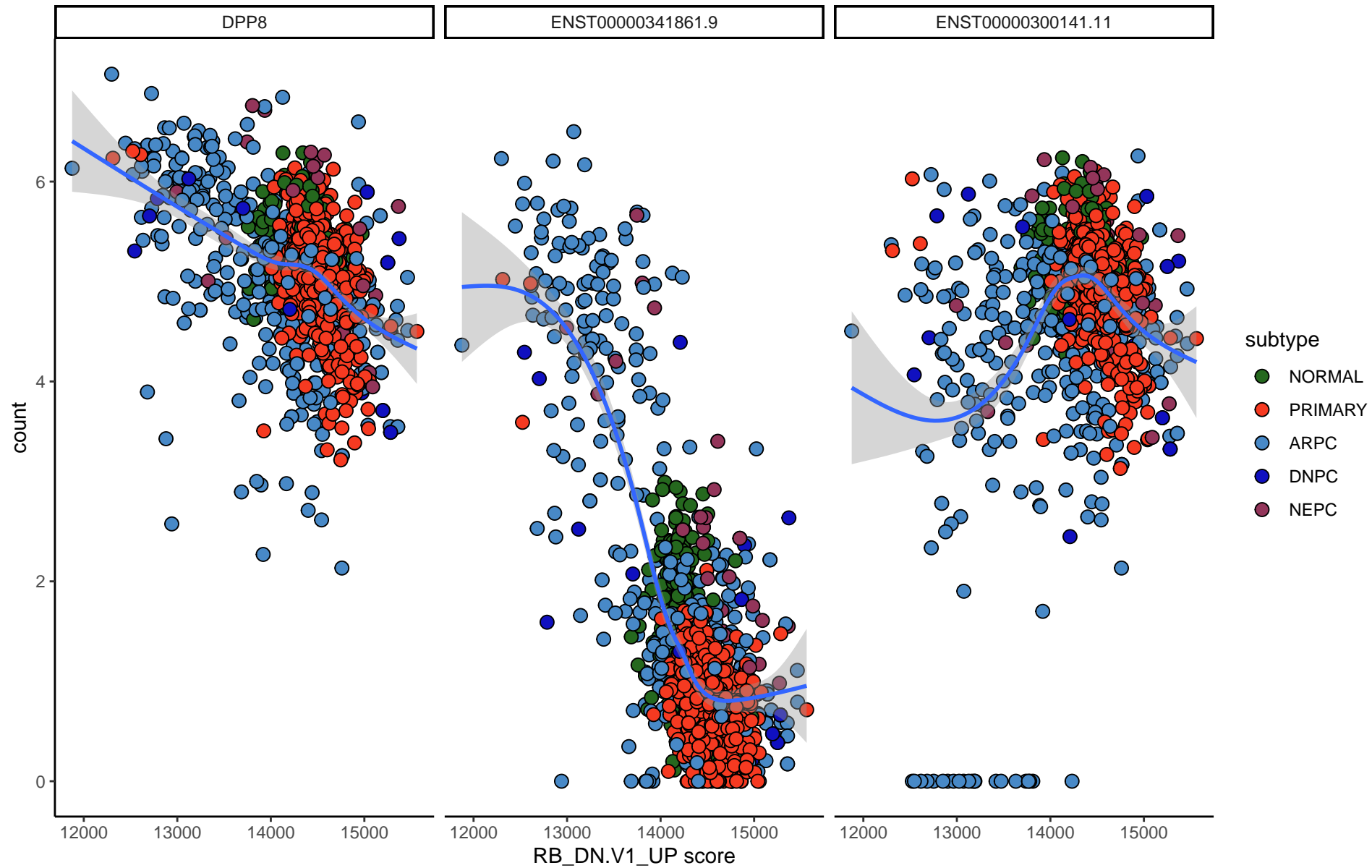
DPP8
Delta corr= 1.37
Gene corr= -0.42



HALLMARK_UV_RESPONSE_UP
ENST00000341861.9 – DPP8–203 PCC = -0.68
ENST00000300141.11 – DPP8–201 PCC = 0.69



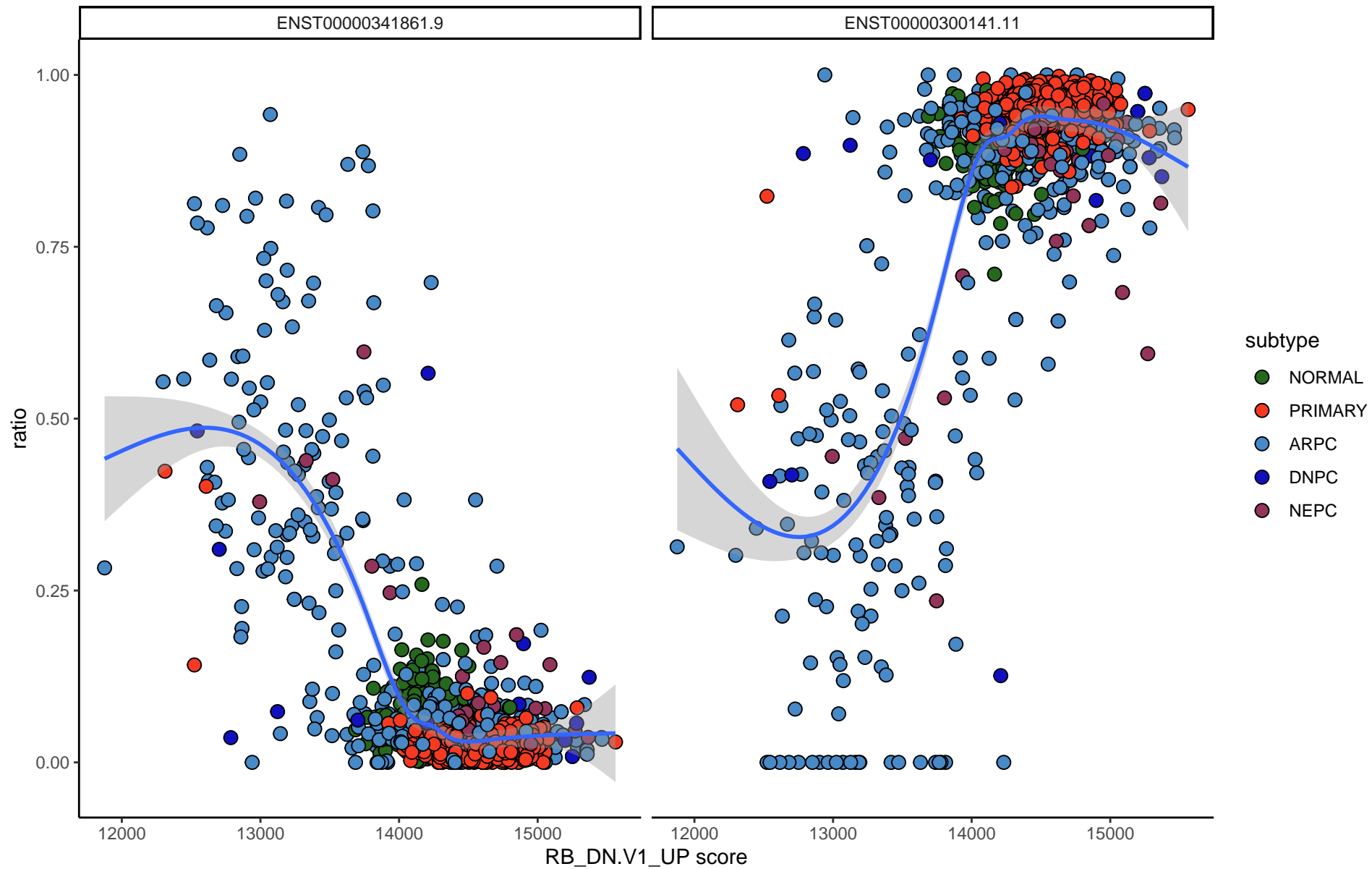
DPP8
Delta corr= 1.37
Gene corr= -0.41



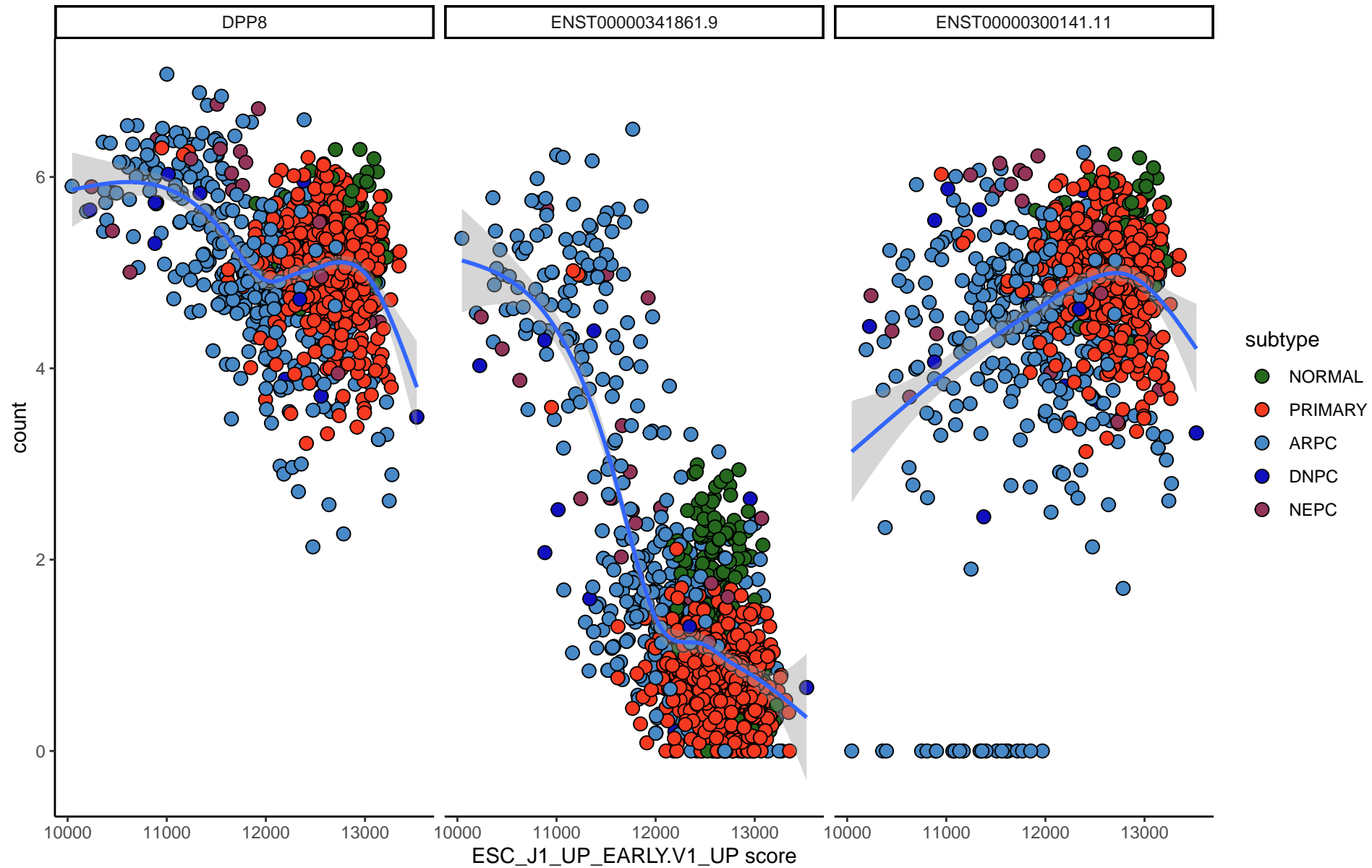
RB_DN.V1_UP

ENST00000341861.9 – DPP8–203 PCC = -0.68

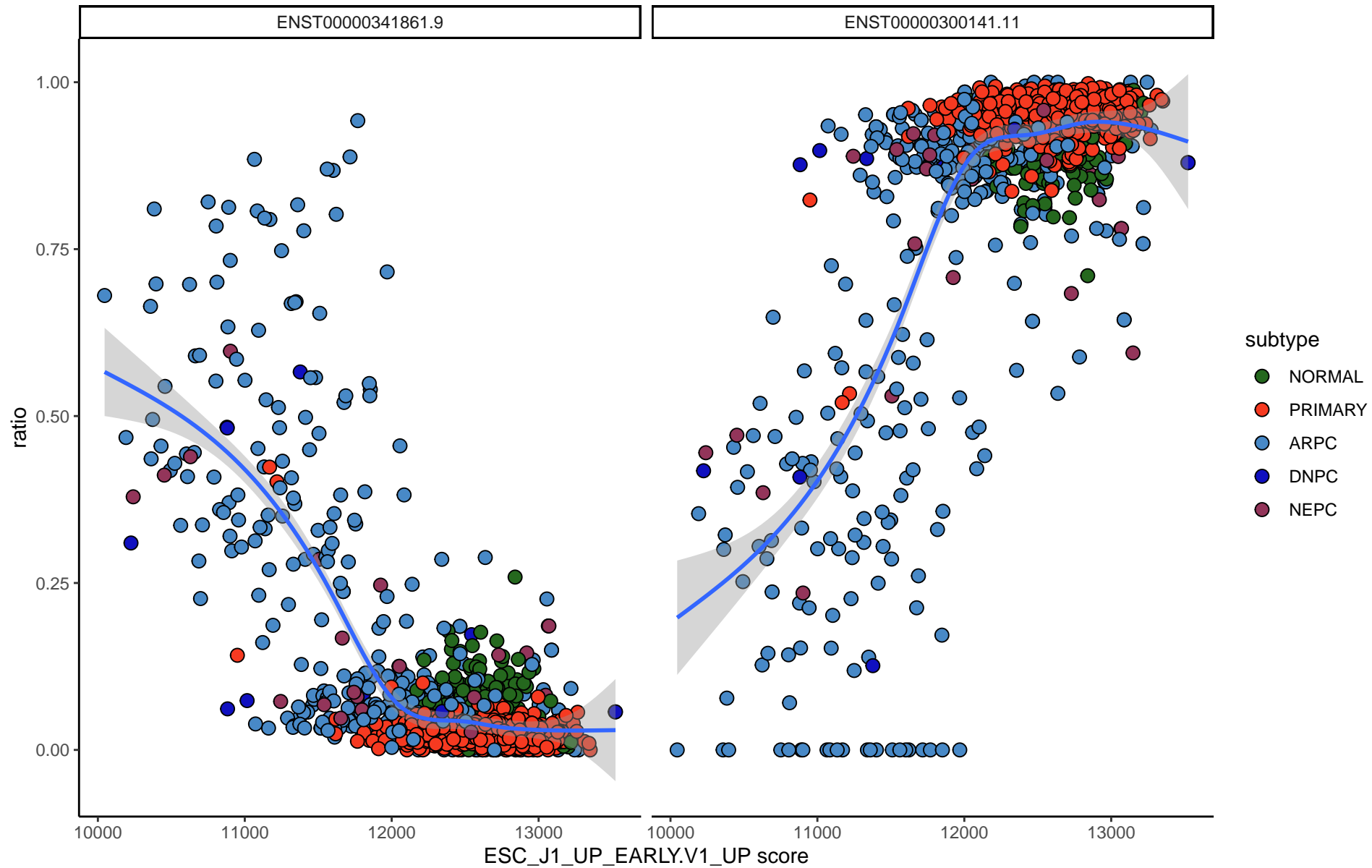
ENST00000300141.11 – DPP8–201 PCC = 0.69



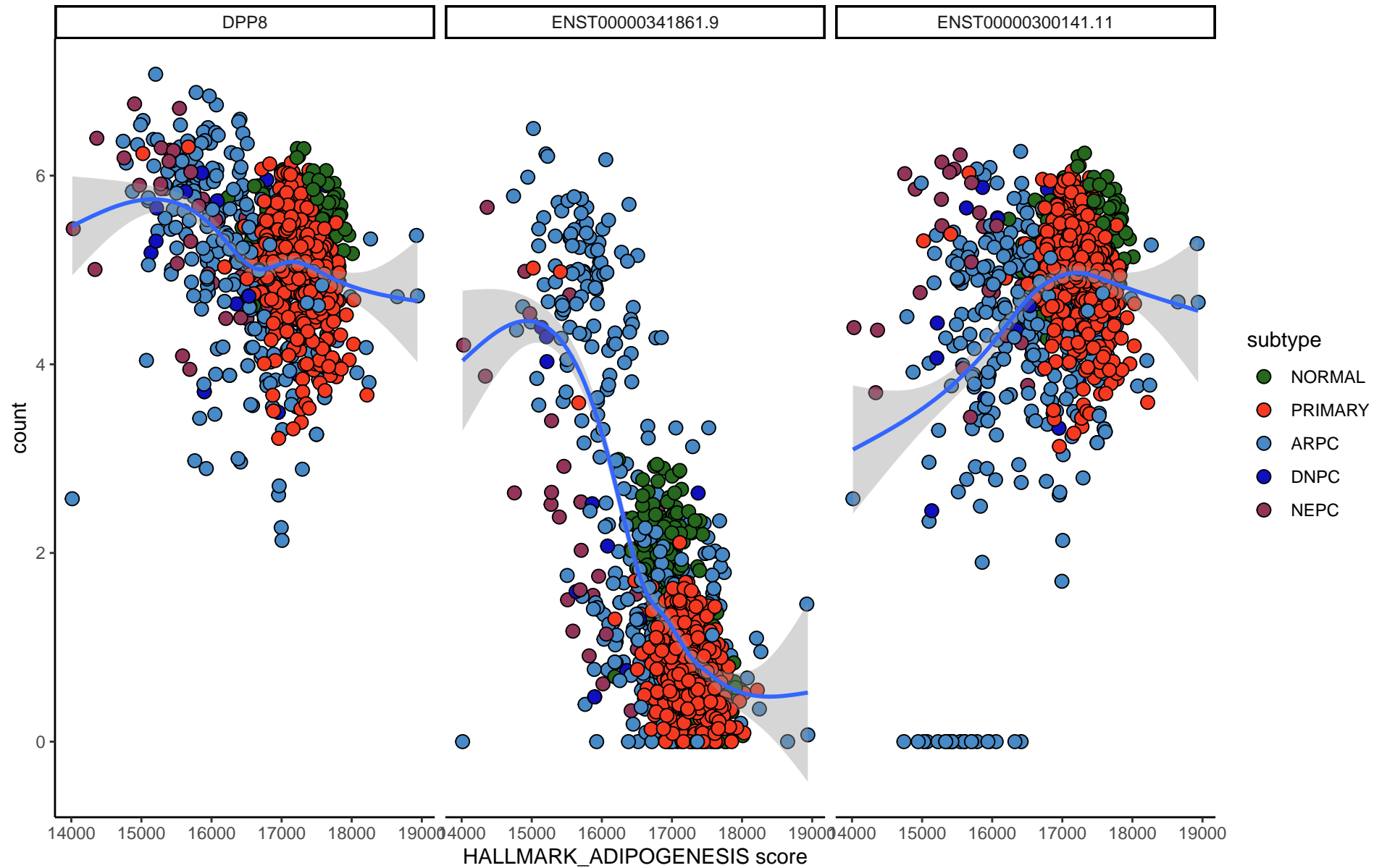
DPP8
Delta corr= 1.36
Gene corr= -0.31



ESC_J1_UP_EARLY.V1_UP
ENST00000341861.9 – DPP8–203 PCC = -0.67
ENST00000300141.11 – DPP8–201 PCC = 0.69



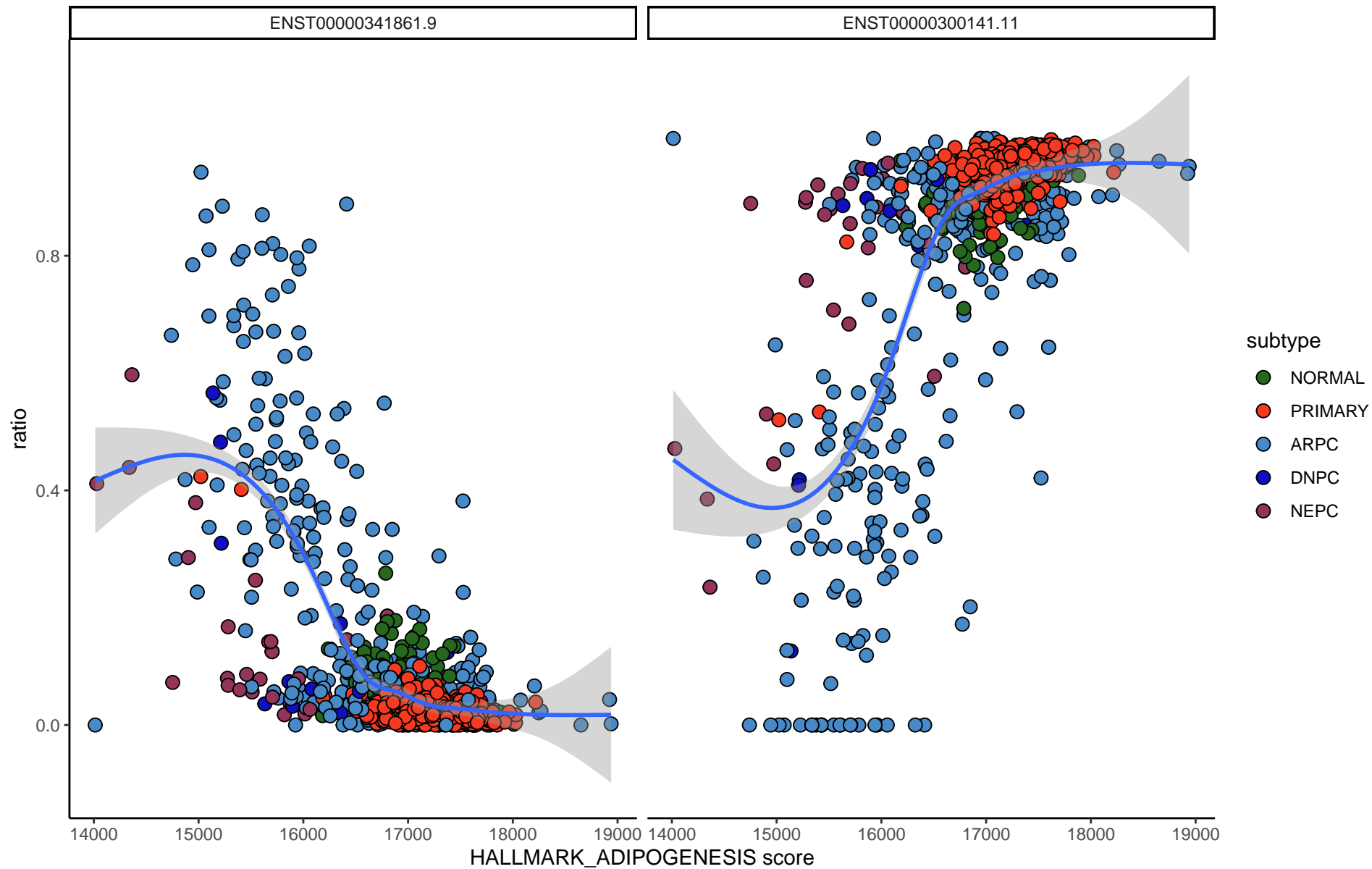
DPP8
Delta corr= 1.35
Gene corr= -0.28



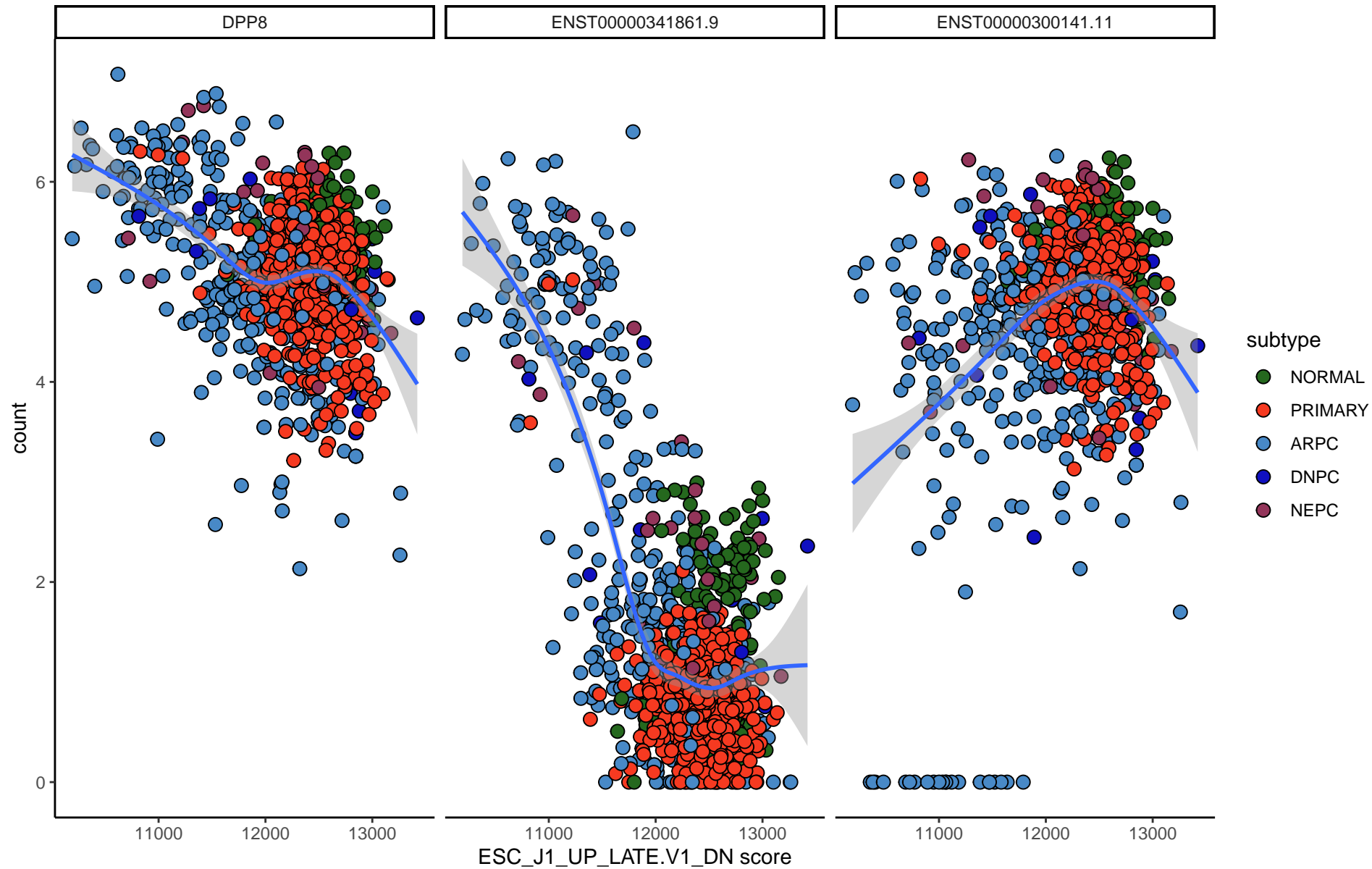
HALLMARK_ADIPOGENESIS

ENST00000341861.9 – DPP8–203 PCC = -0.67

ENST00000300141.11 – DPP8–201 PCC = 0.68



DPP8
Delta corr= 1.34
Gene corr= -0.33



ESC_J1_UP_LATE.V1_DN

ENST00000341861.9 – DPP8–203 PCC = -0.66

ENST00000300141.11 – DPP8–201 PCC = 0.68

