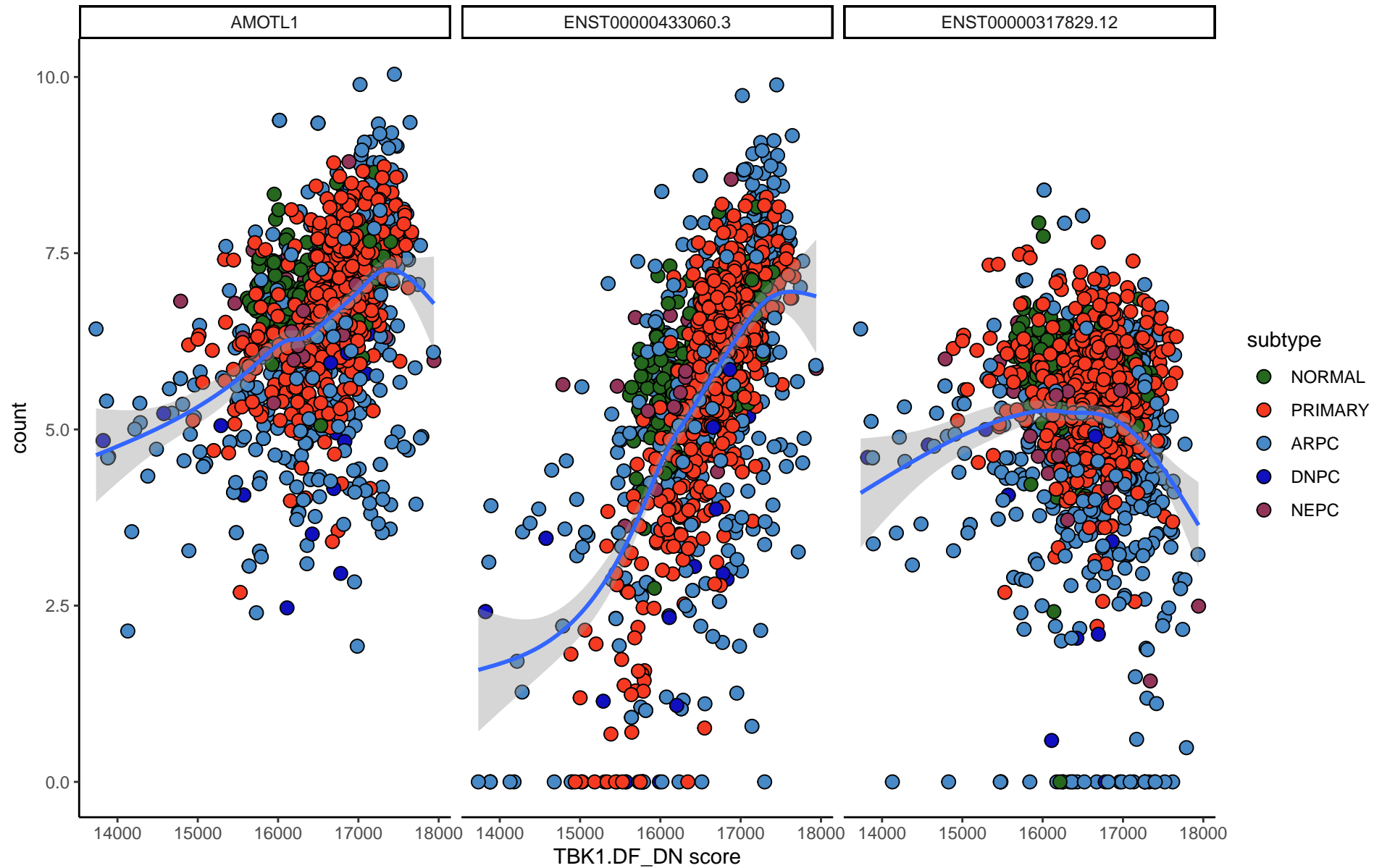


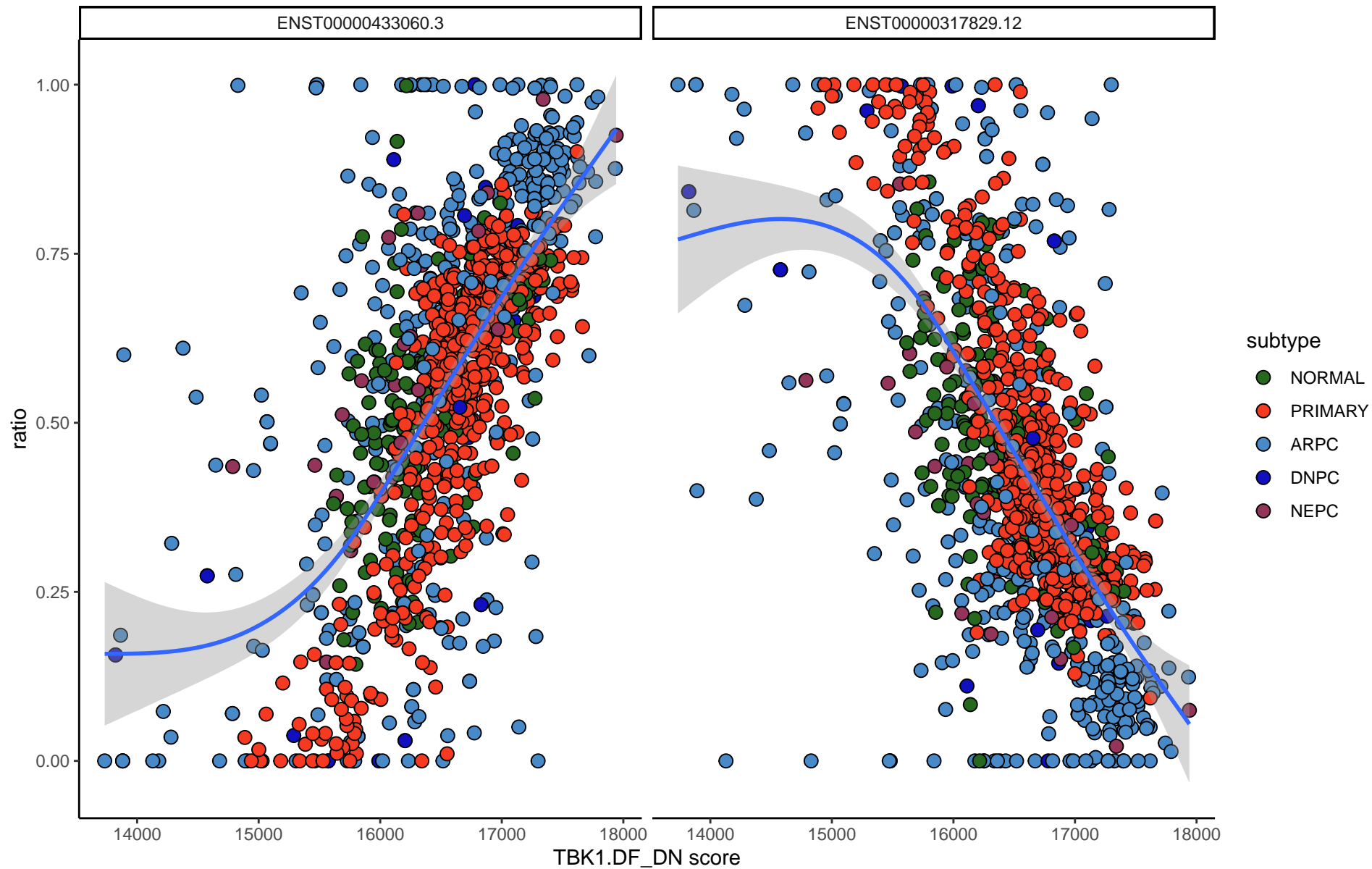
AMOTL1
Delta corr= 1.28
Gene corr= 0.42



TBK1.DF_DN

ENST00000433060.3 – AMOTL1–203 PCC = 0.65

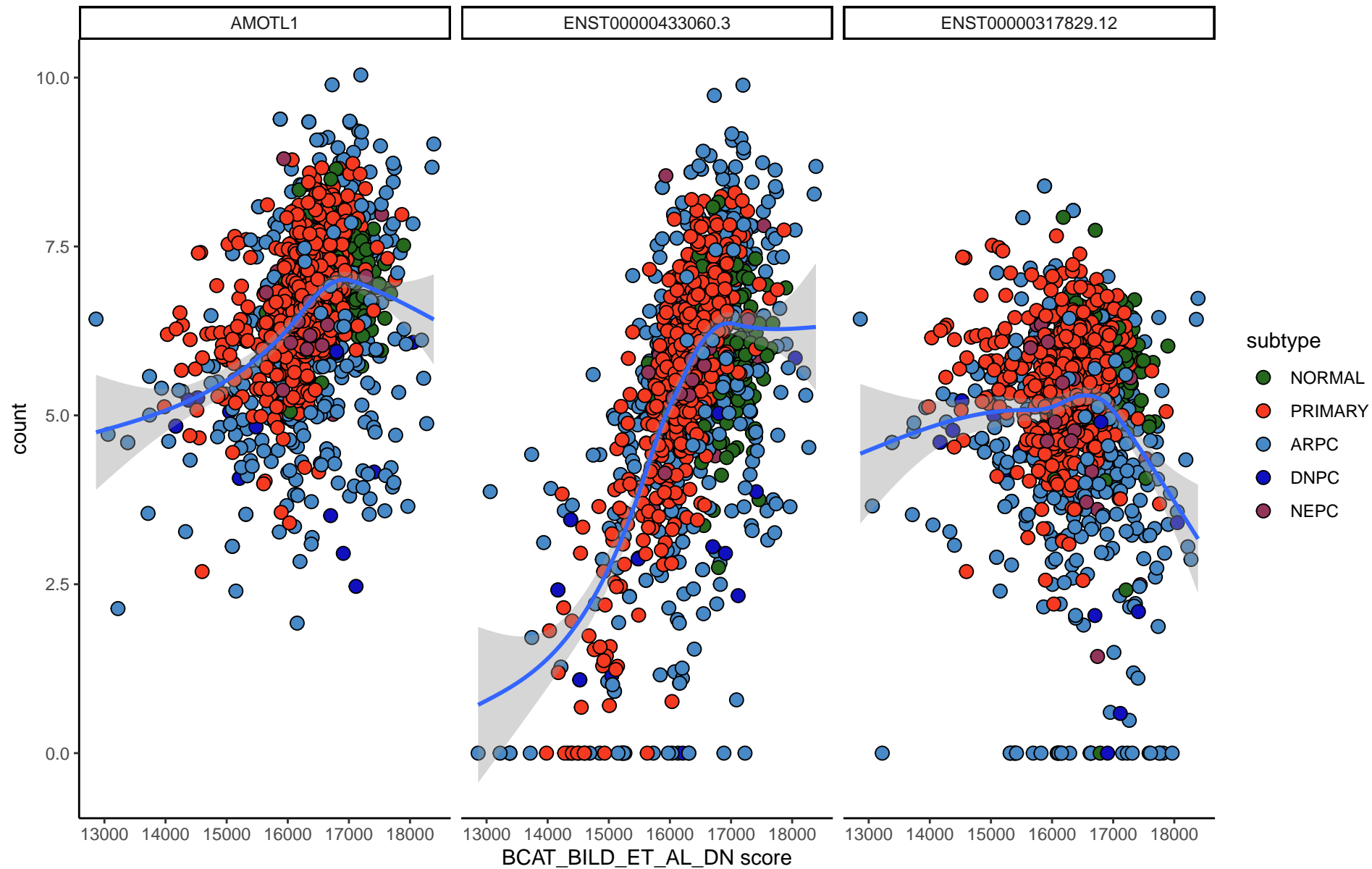
ENST00000317829.12 – AMOTL1–202 PCC = -0.64



AMOTL1

Delta corr= 1.19

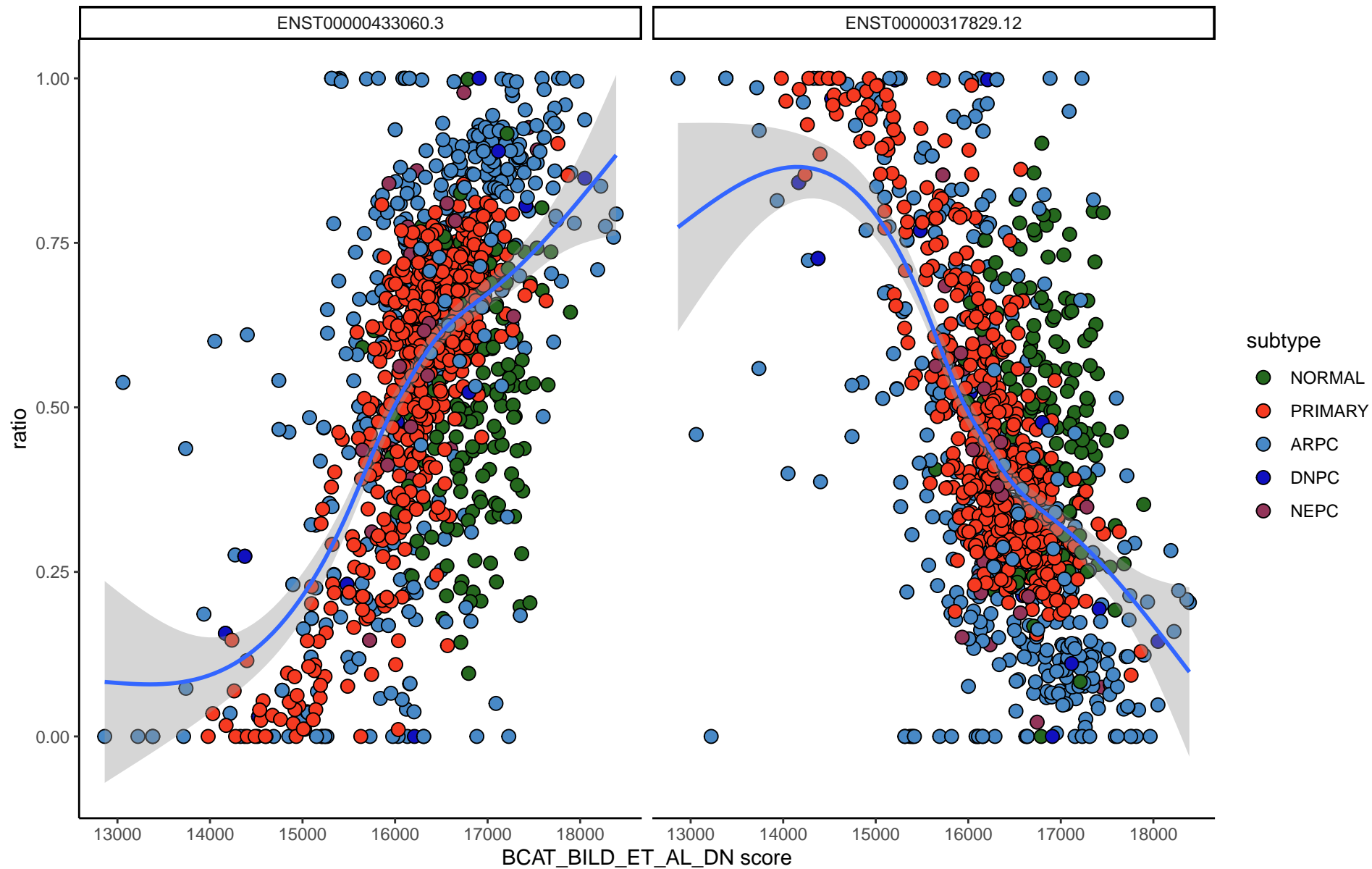
Gene corr= 0.37



BCAT_BILD_ET_AL_DN

ENST00000433060.3 – AMOTL1–203 PCC = 0.6

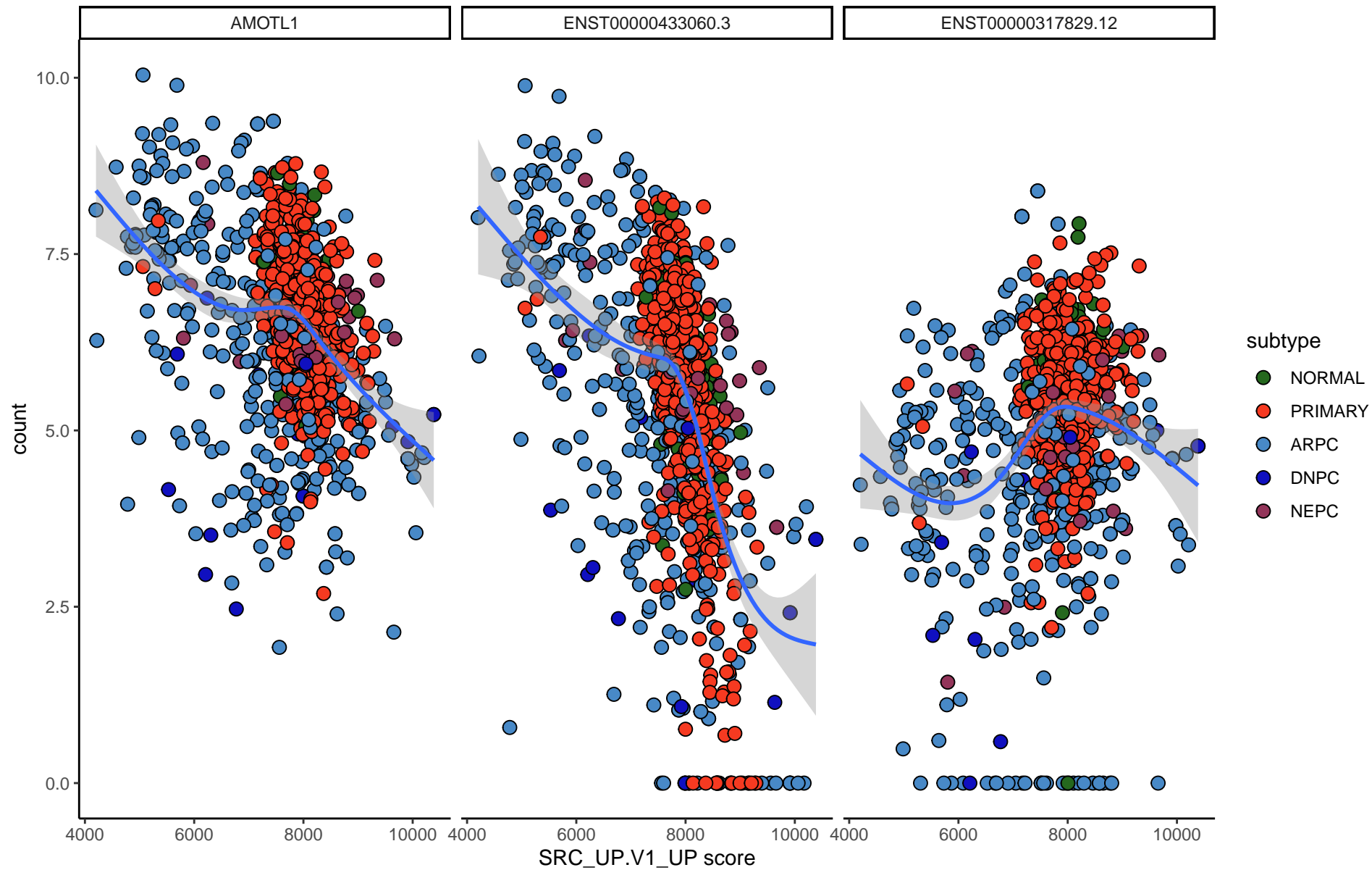
ENST00000317829.12 – AMOTL1–202 PCC = -0.59



AMOTL1

Delta corr= 1.18

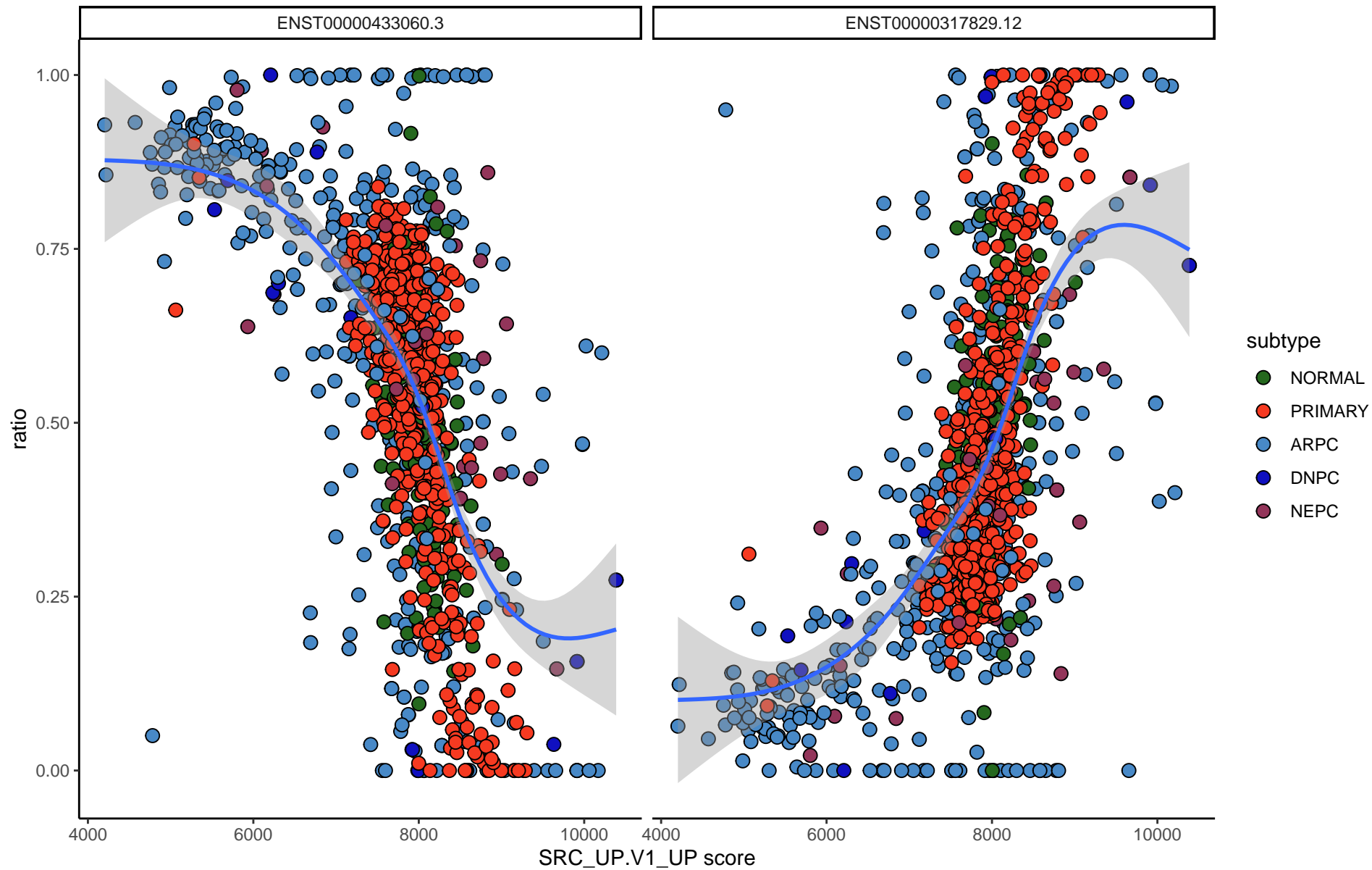
Gene corr= -0.33



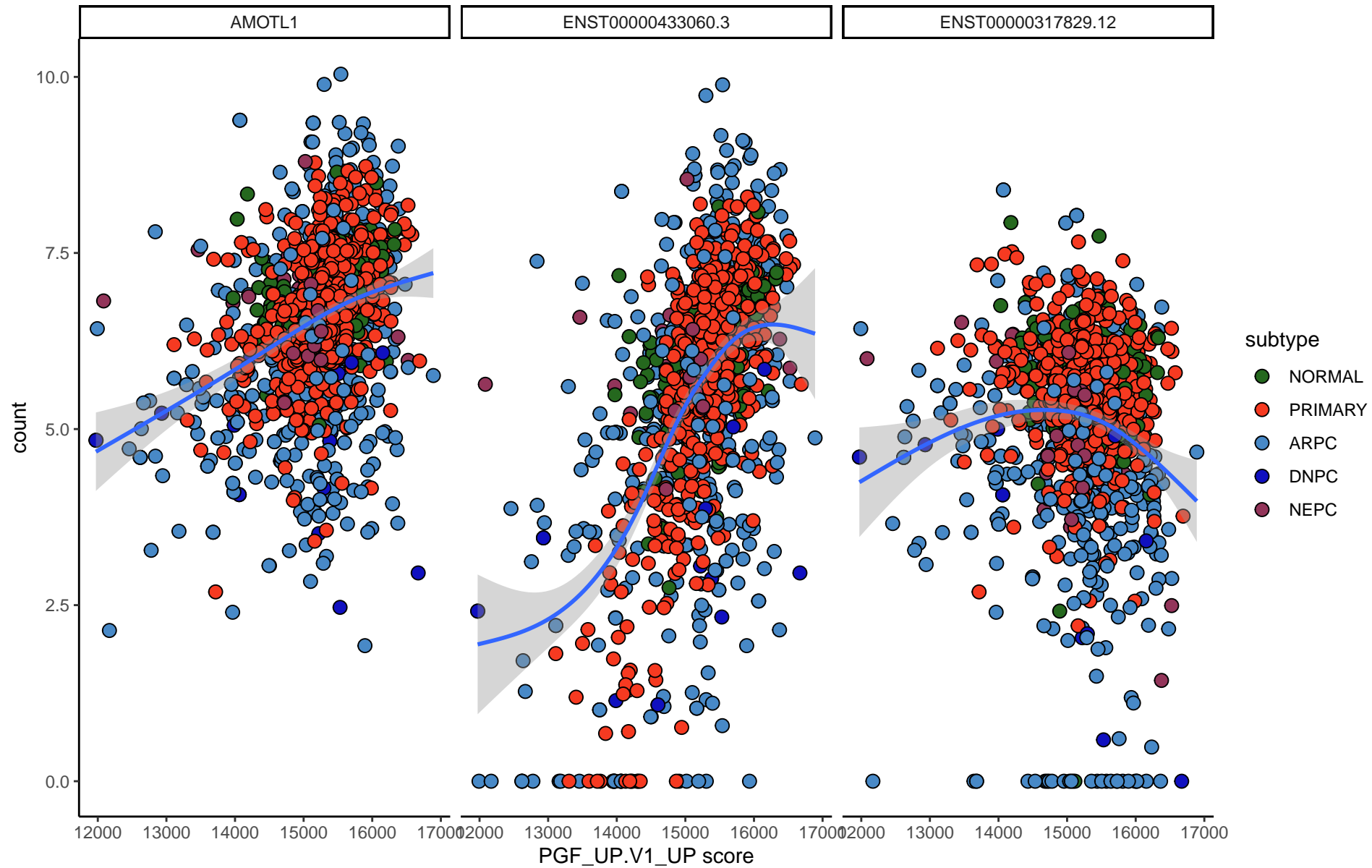
SRC_UP.V1_UP

ENST00000433060.3 – AMOTL1–203 PCC = -0.59

ENST00000317829.12 – AMOTL1–202 PCC = 0.59



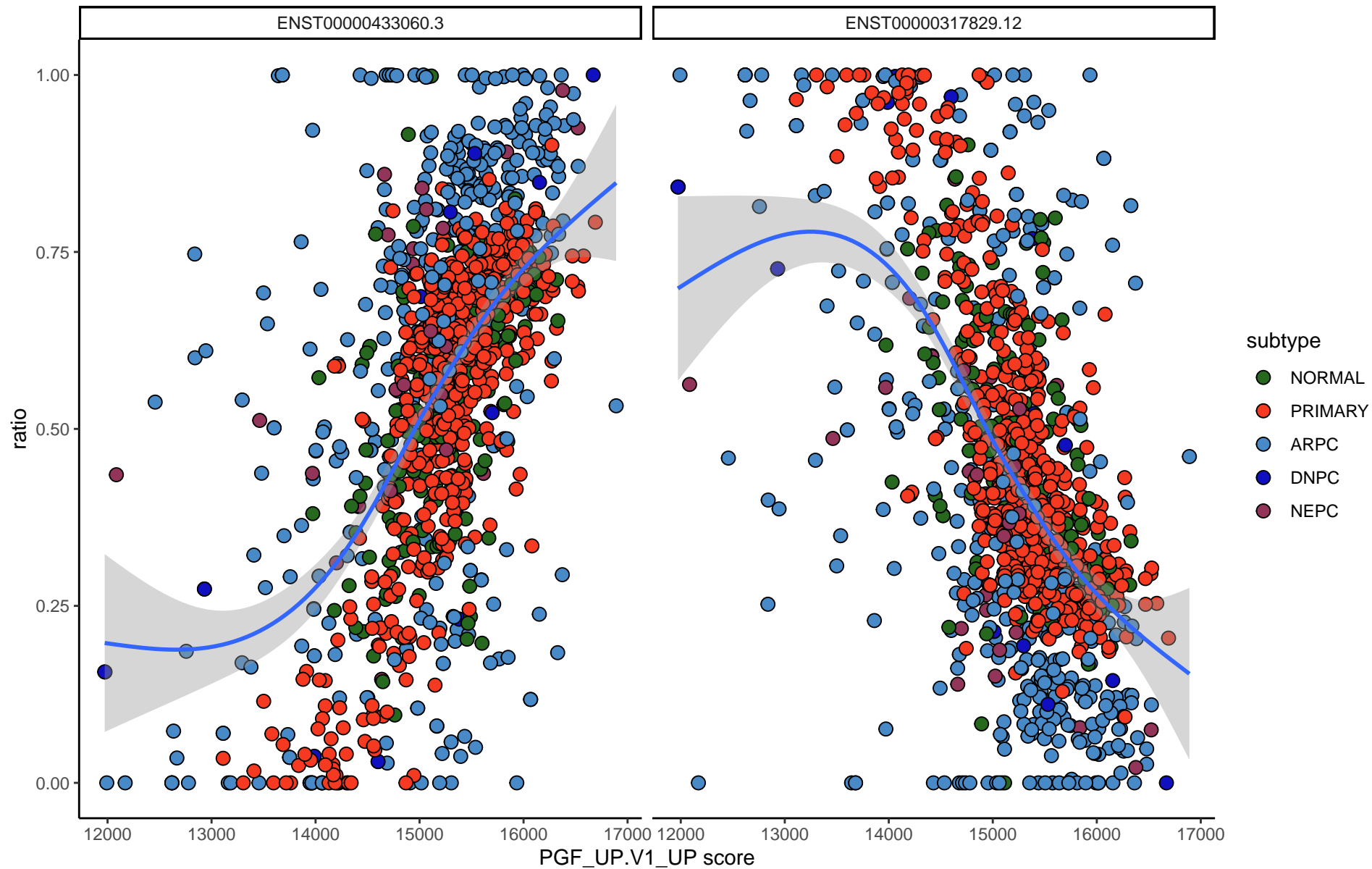
AMOTL1
Delta corr= 1.13
Gene corr= 0.33



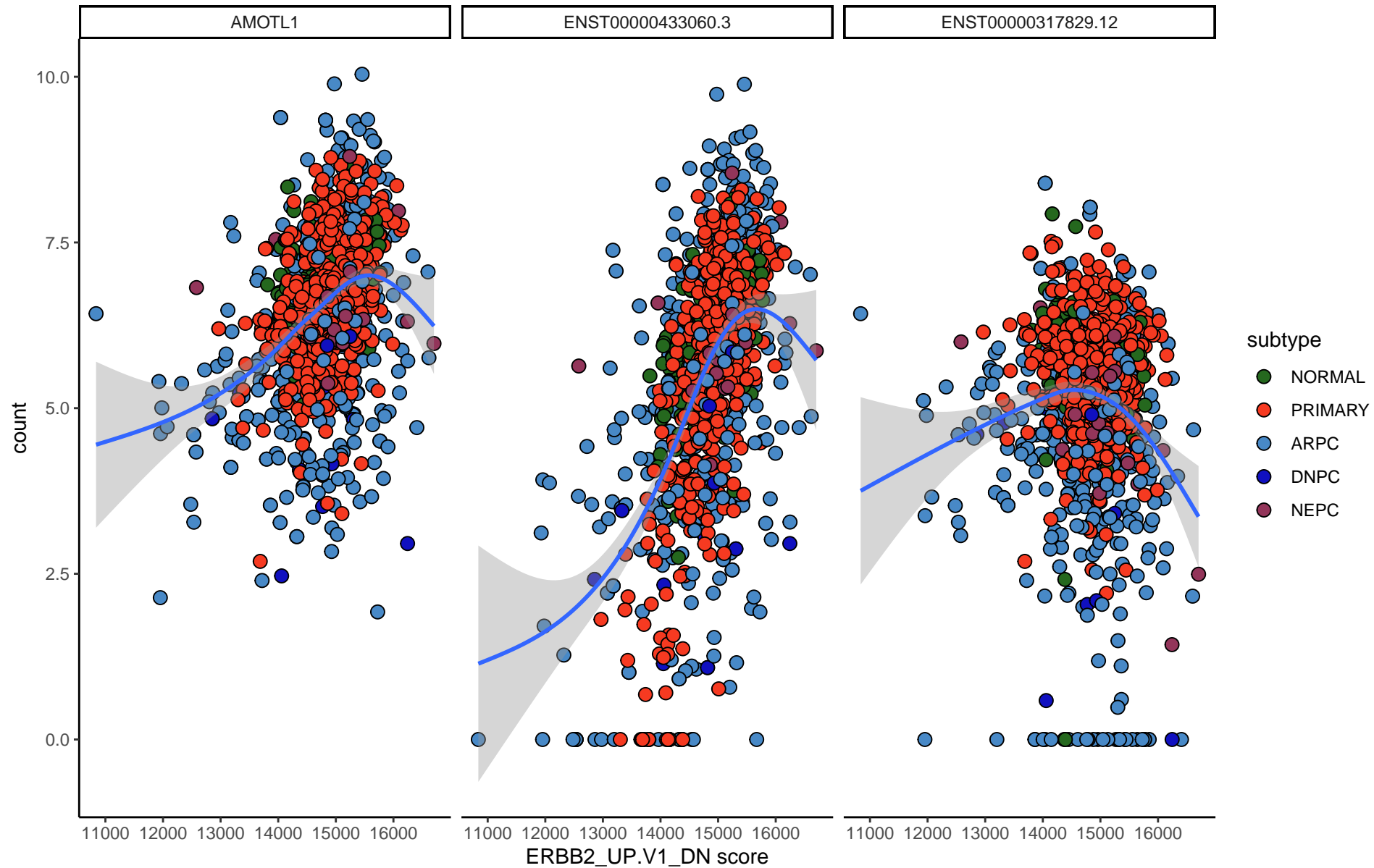
PGF_UP.V1_UP

ENST00000433060.3 – AMOTL1–203 PCC = 0.57

ENST00000317829.12 – AMOTL1–202 PCC = -0.56



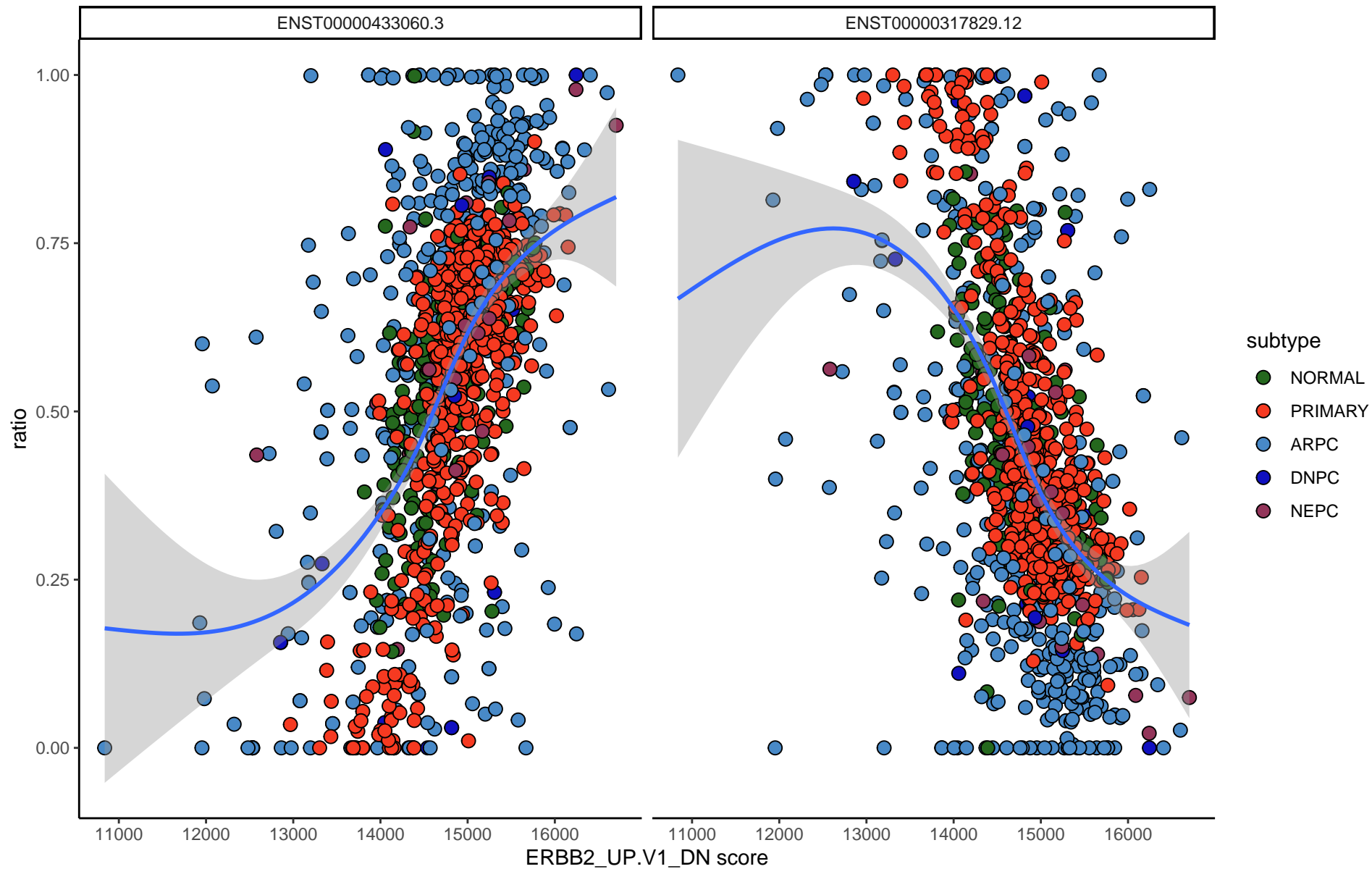
AMOTL1
Delta corr= 1.09
Gene corr= 0.35



ERBB2_UP.V1_DN

ENST00000433060.3 – AMOTL1–203 PCC = 0.55

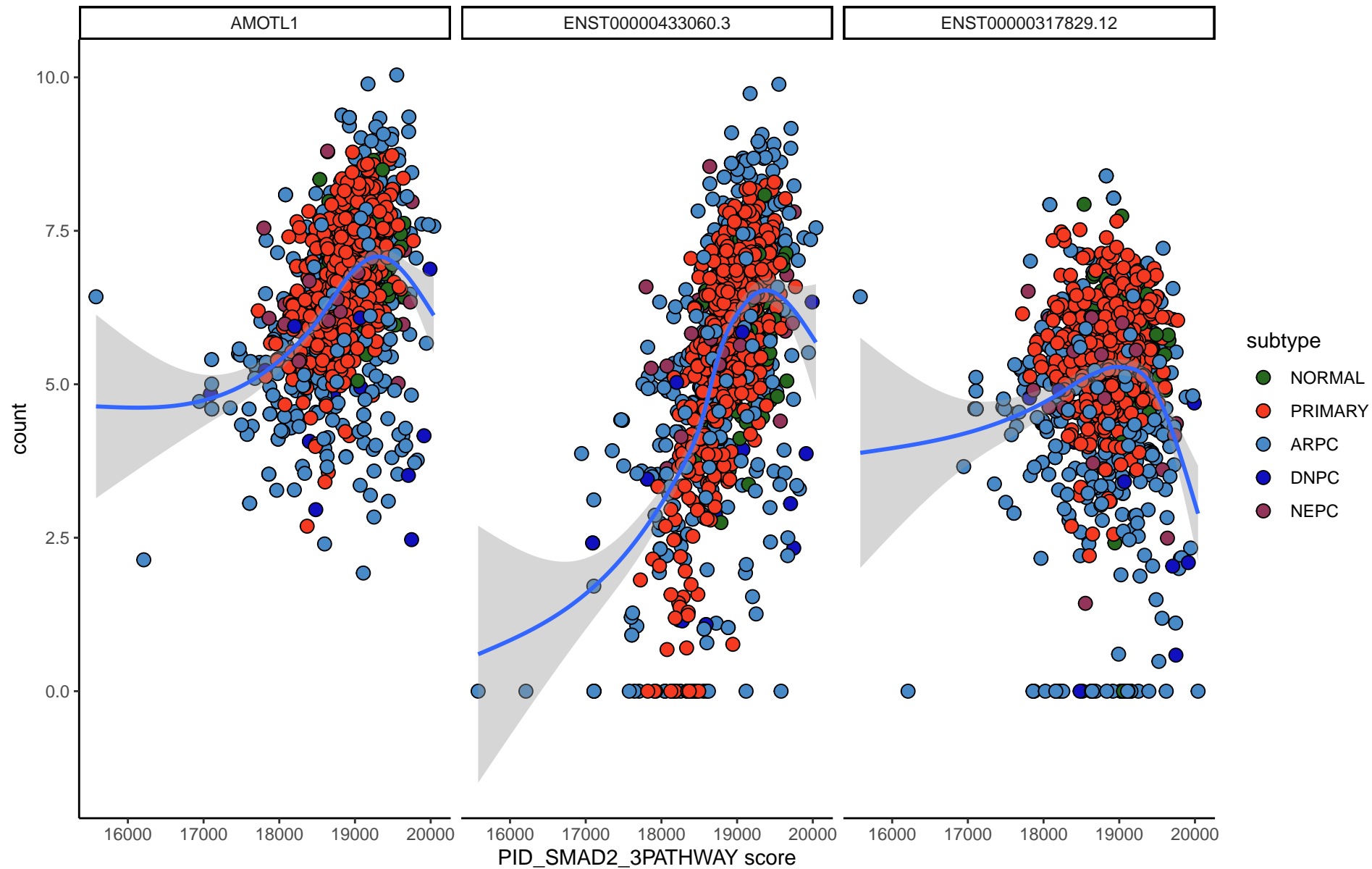
ENST00000317829.12 – AMOTL1–202 PCC = -0.54



AMOTL1

Delta corr= 1.09

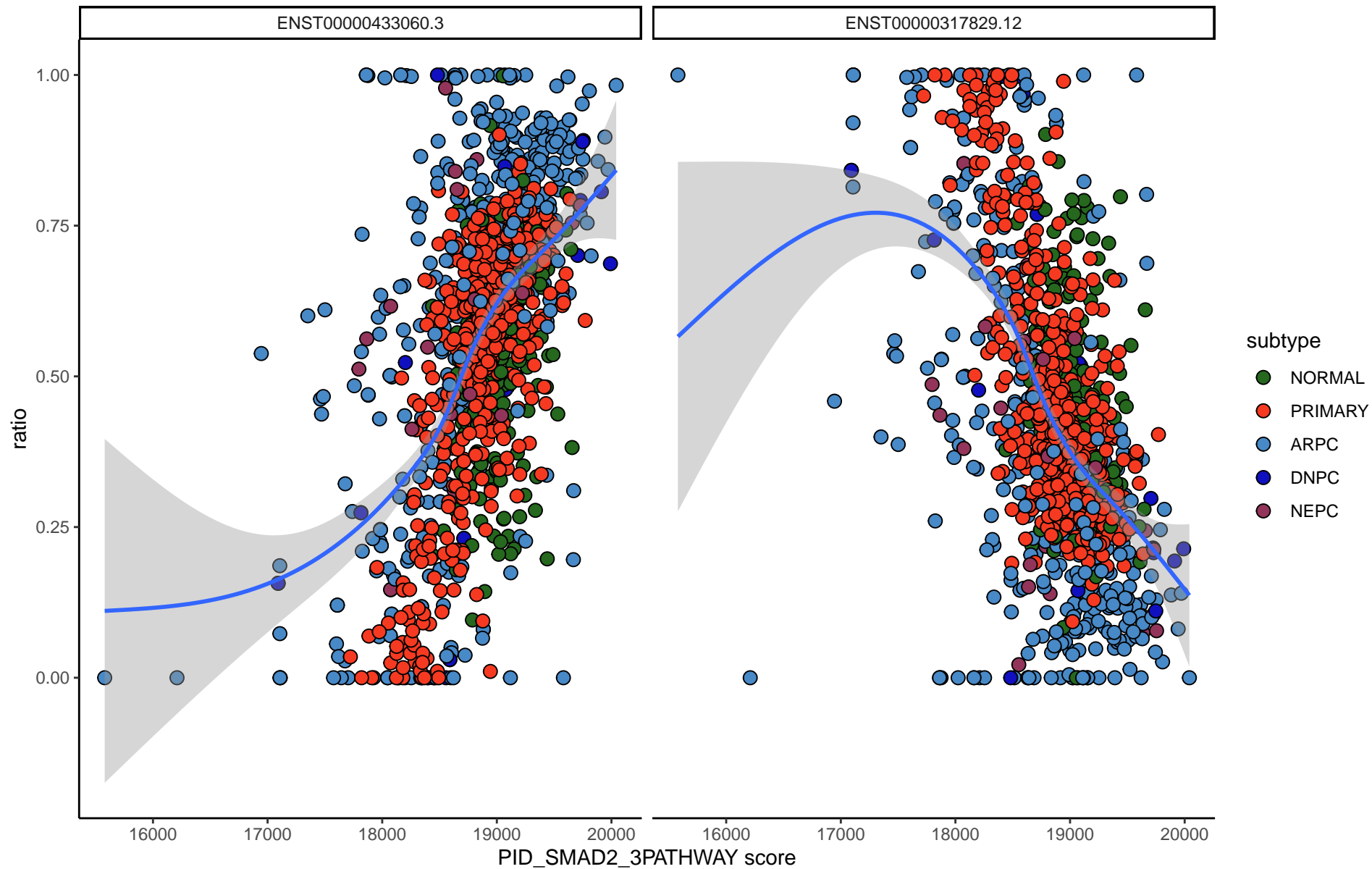
Gene corr= 0.42



PID_SMAD2_3PATHWAY

ENST00000433060.3 – AMOTL1–203 PCC = 0.55

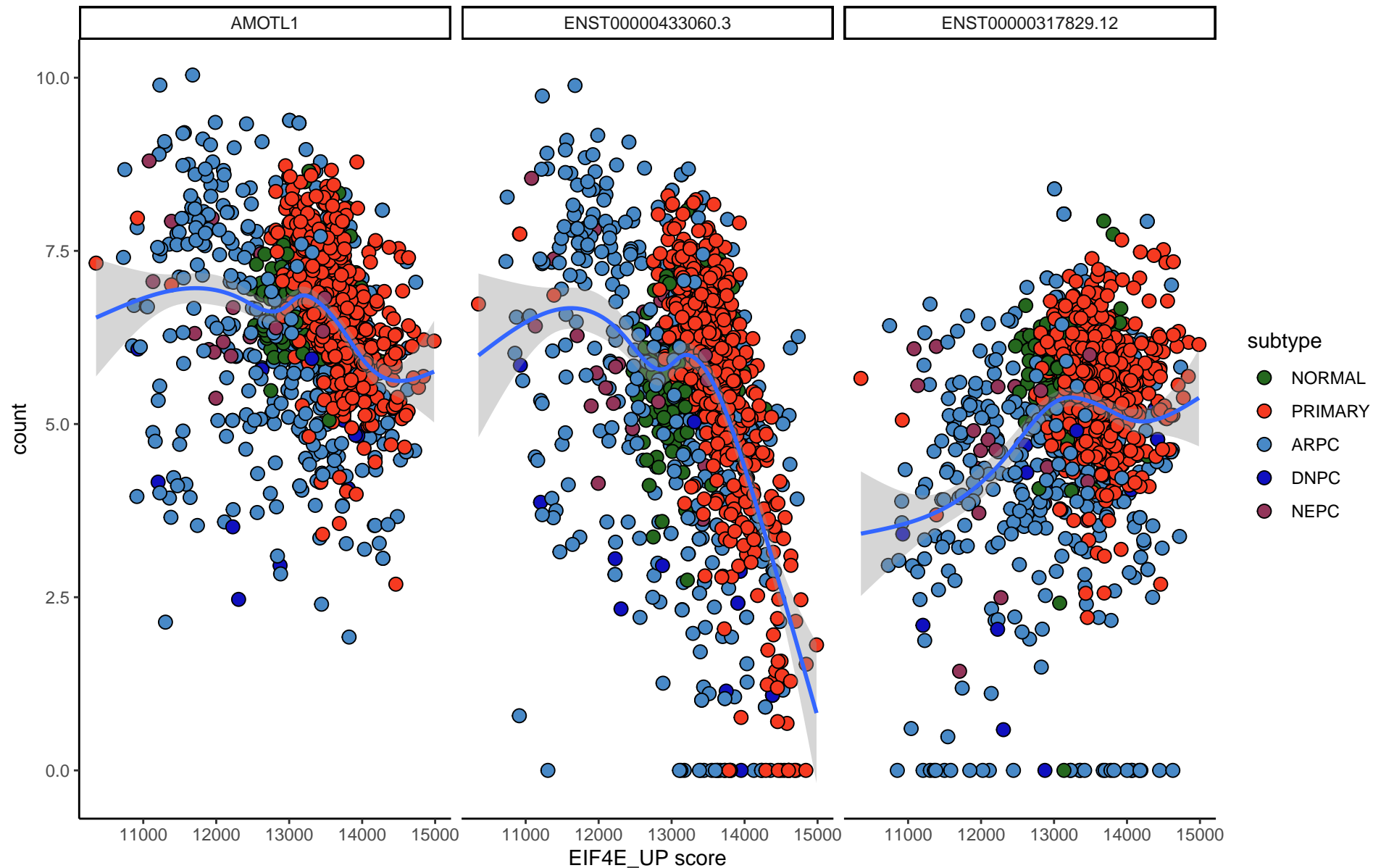
ENST00000317829.12 – AMOTL1–202 PCC = -0.54



AMOTL1

Delta corr= 1.08

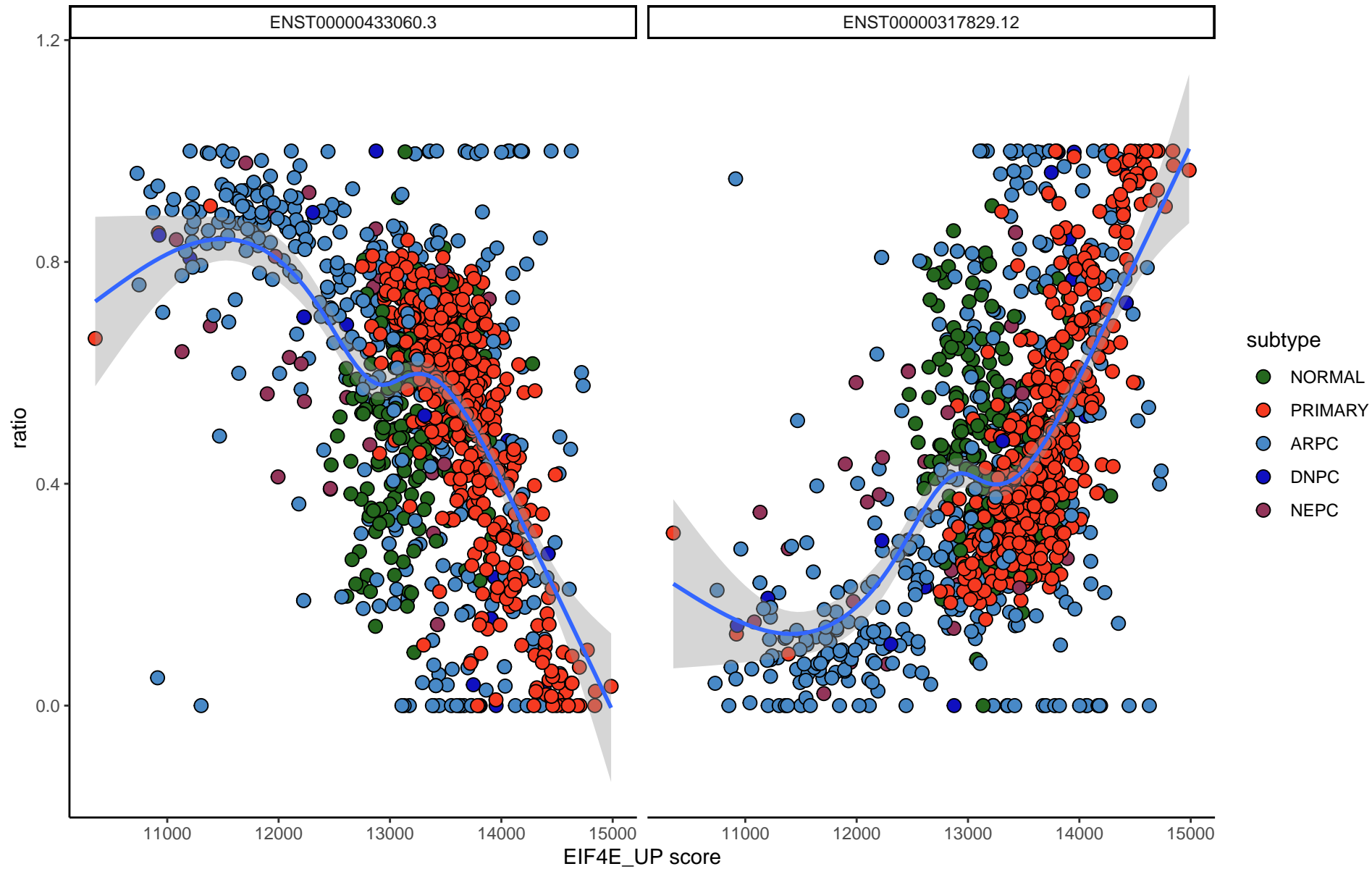
Gene corr= -0.24



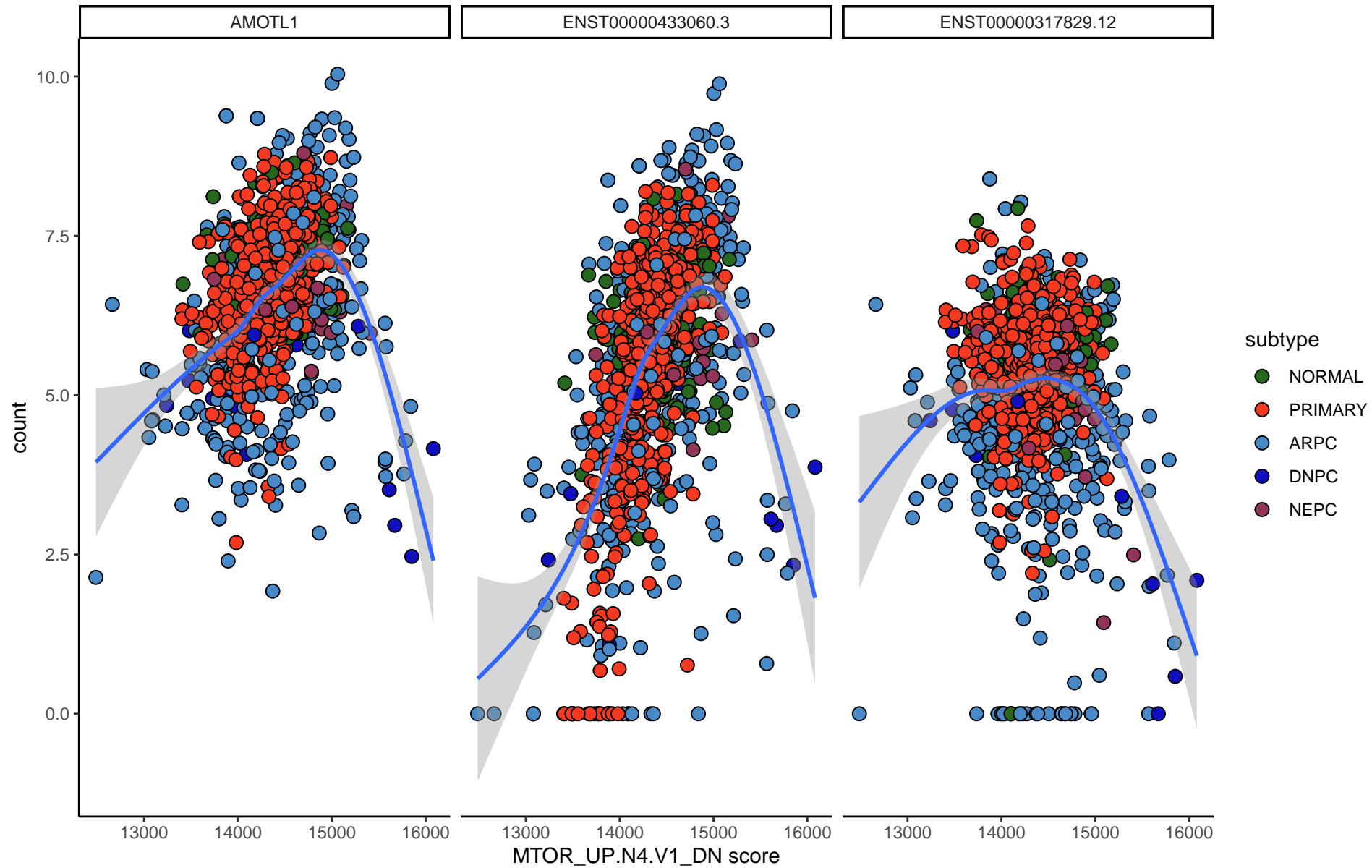
EIF4E_UP

ENST00000433060.3 – AMOTL1–203 PCC = -0.53

ENST00000317829.12 – AMOTL1–202 PCC = 0.55



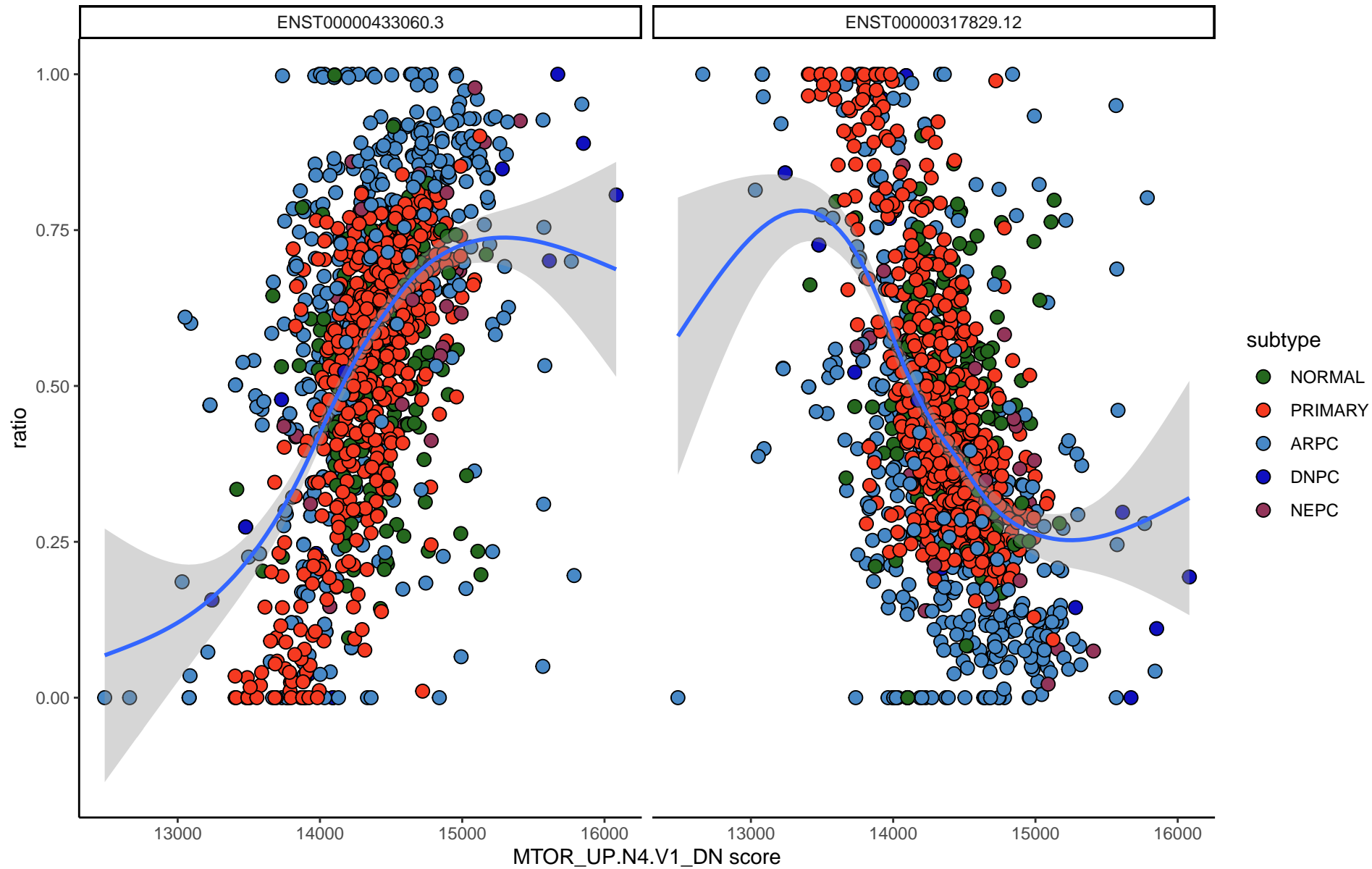
AMOTL1
Delta corr= 1.04
Gene corr= 0.33



MTOR_UP.N4.V1_DN

ENST00000433060.3 – AMOTL1–203 PCC = 0.53

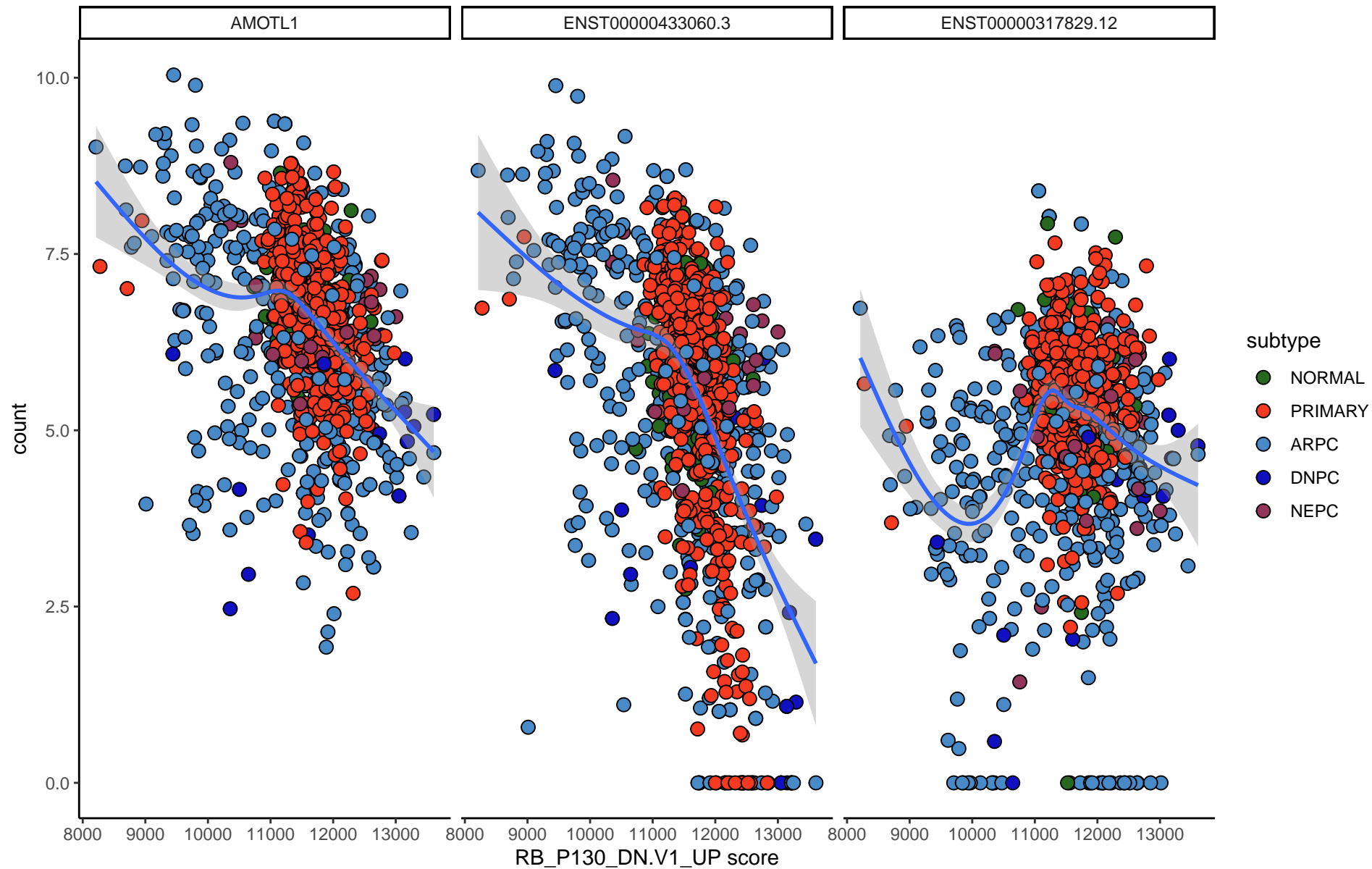
ENST00000317829.12 – AMOTL1–202 PCC = -0.52



AMOTL1

Delta corr= 1.04

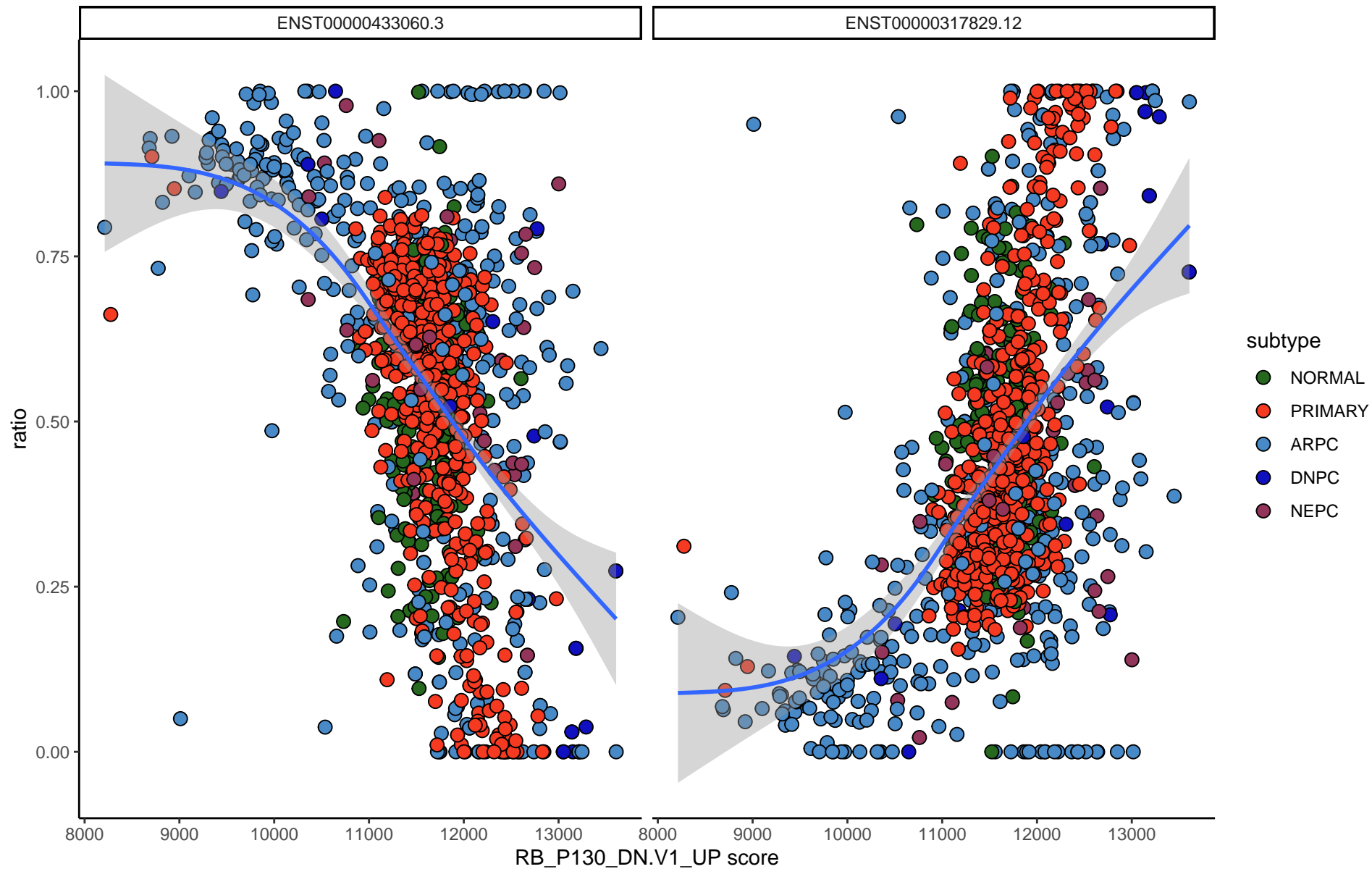
Gene corr= -0.37



RB_P130_DN.V1_UP

ENST00000433060.3 – AMOTL1–203 PCC = -0.52

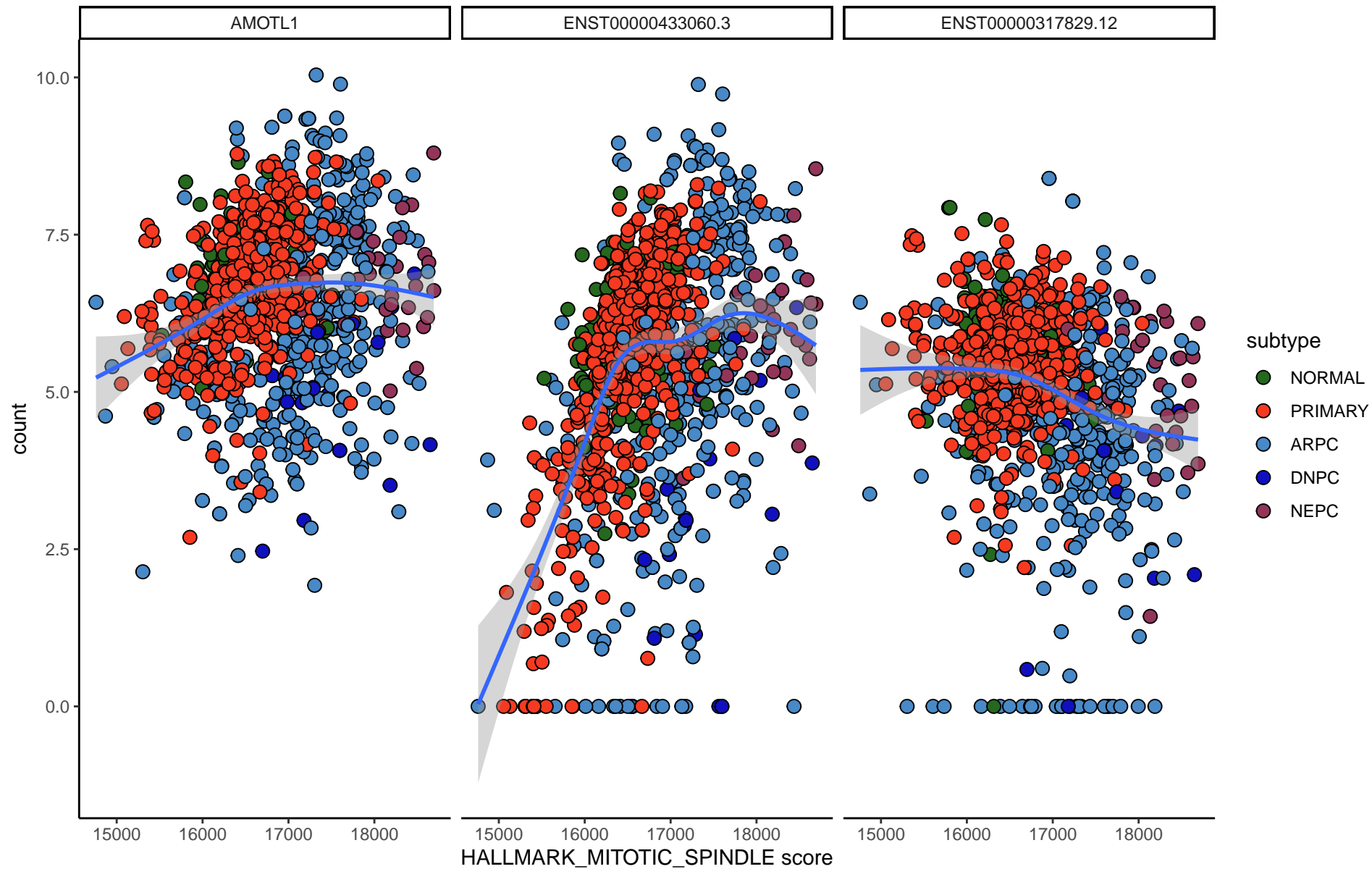
ENST00000317829.12 – AMOTL1–202 PCC = 0.53



AMOTL1

Delta corr= 1.01

Gene corr= 0.16



HALLMARK_MITOTIC_SPINDLE

ENST00000433060.3 – AMOTL1–203 PCC = 0.51

ENST00000317829.12 – AMOTL1–202 PCC = -0.5

