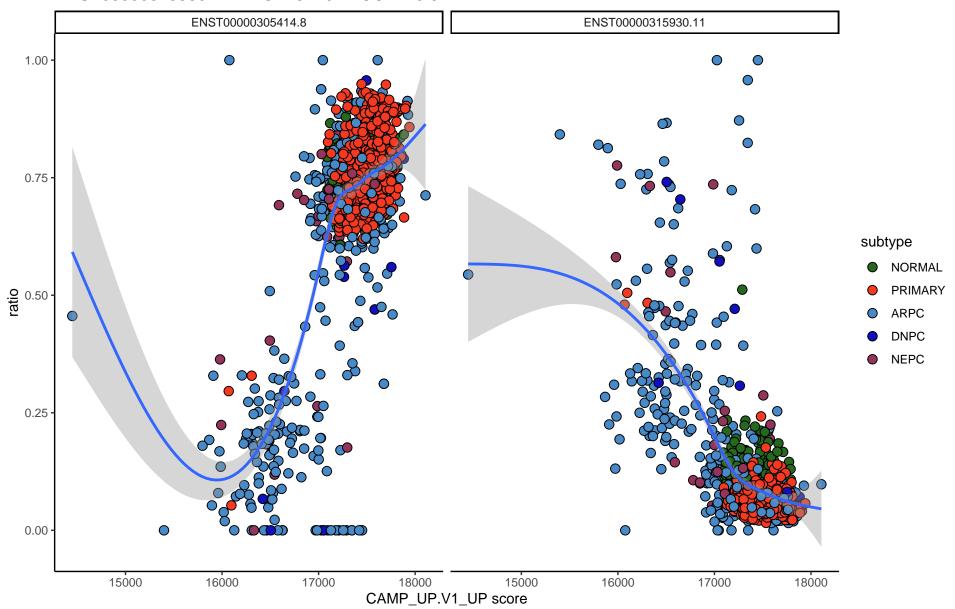
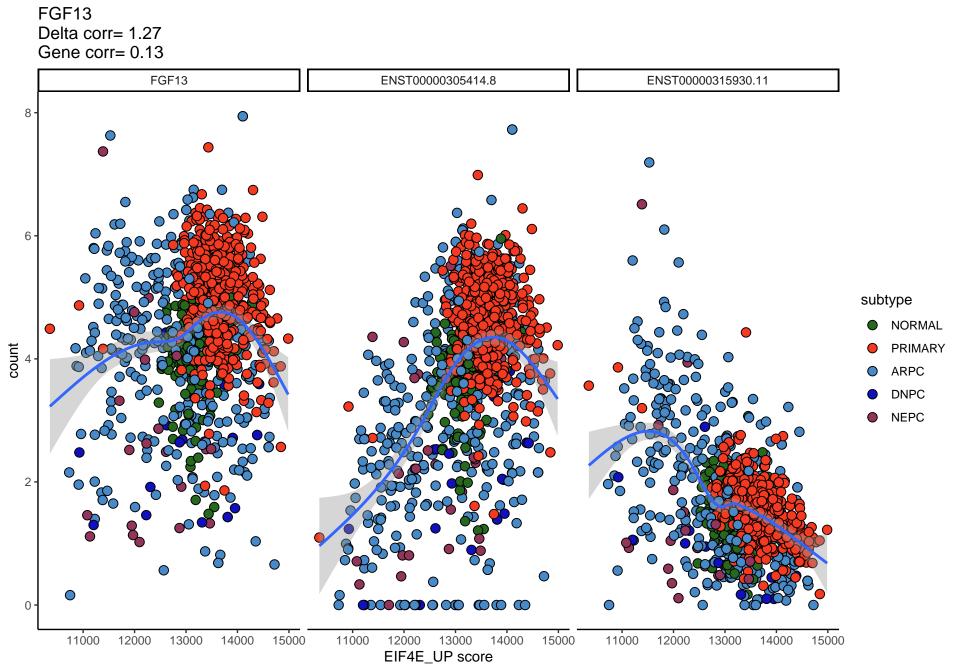
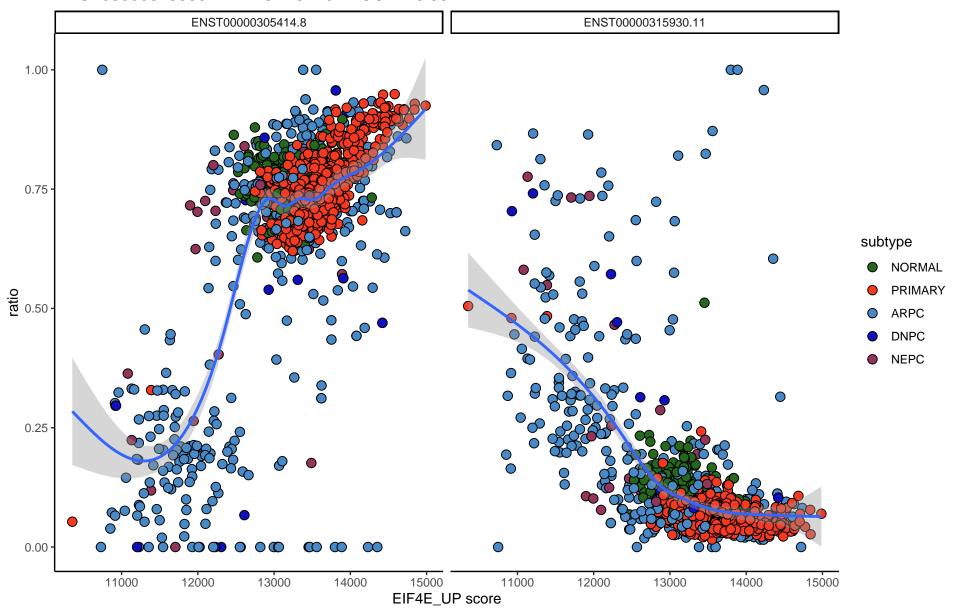
FGF13 Delta corr= 1.35 Gene corr= 0.12 FGF13 ENST00000305414.8 ENST00000315930.11 8 subtype NORMAL count **PRIMARY** ARPC DNPC NEPC 2 -0. 16000 17000 18000 15000 16000 18000 15000 16000 17000 18000 15000 17000 CAMP_UP.V1_UP score

CAMP_UP.V1_UP ENST00000305414.8 - FGF13-201 PCC = 0.71 ENST00000315930.11 - FGF13-202 PCC = -0.64



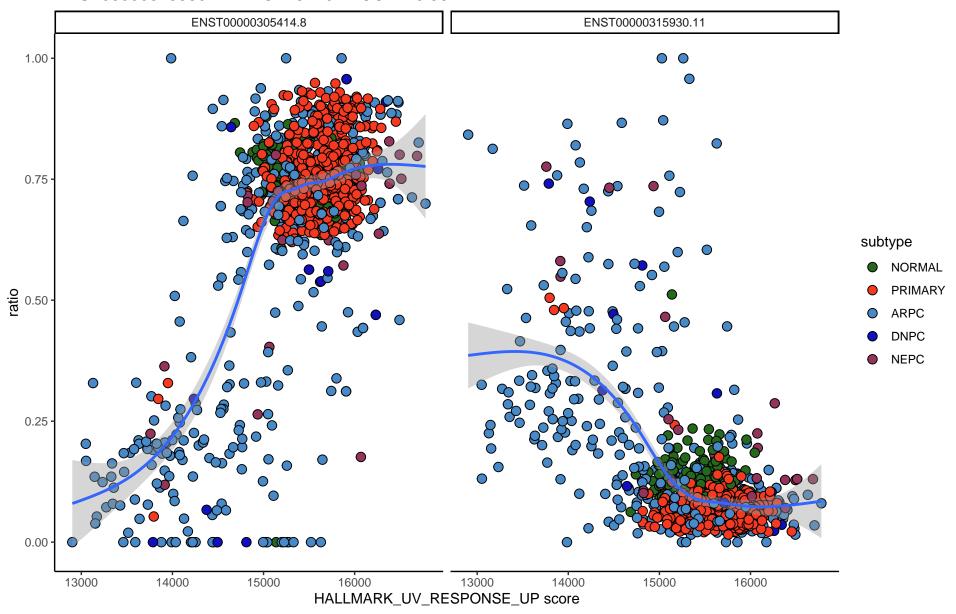


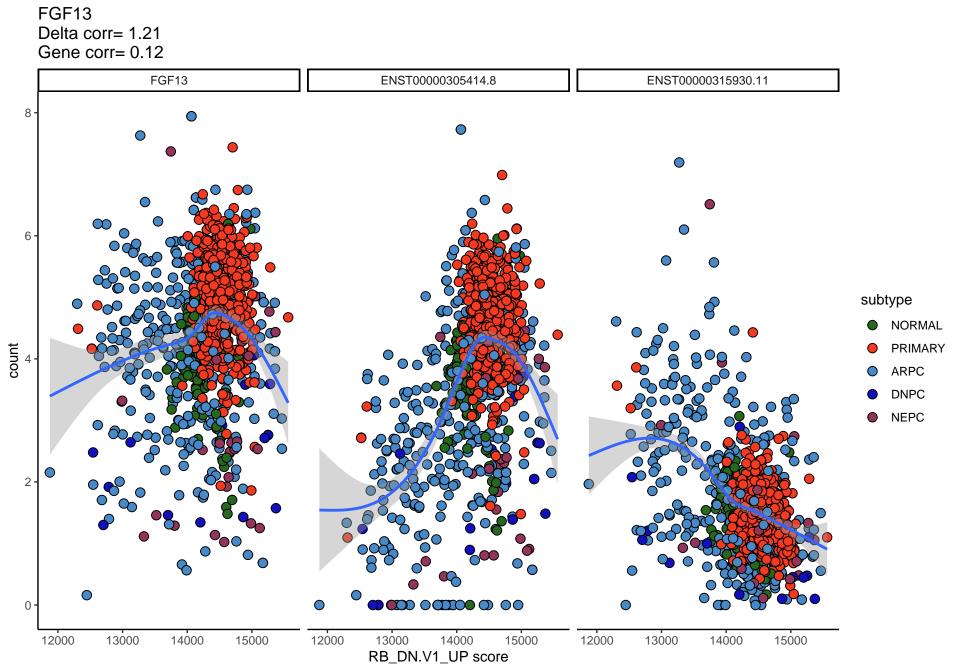
EIF4E_UP ENST00000305414.8 - FGF13-201 PCC = 0.68 ENST00000315930.11 - FGF13-202 PCC = -0.59



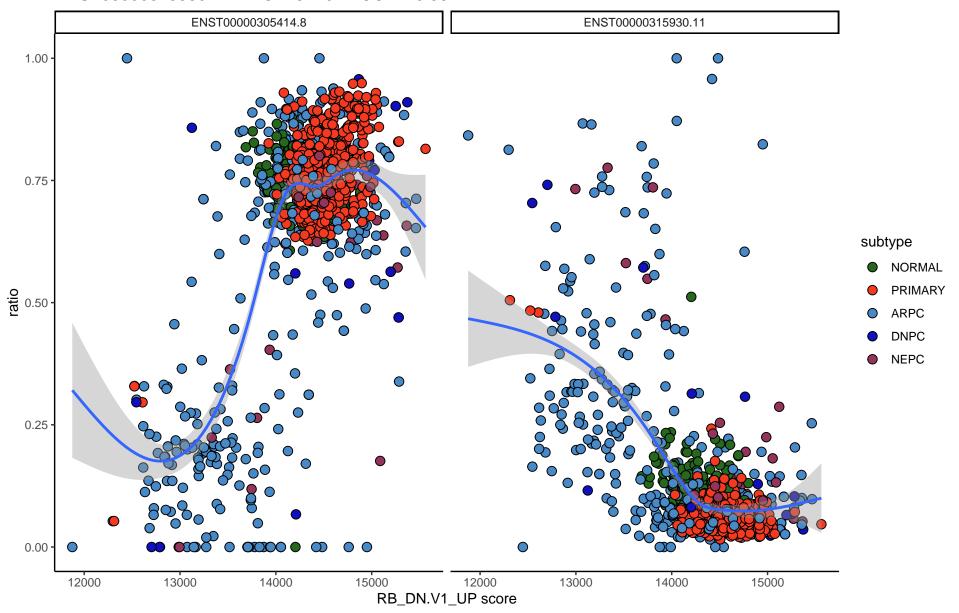
FGF13 Delta corr= 1.23 Gene corr= 0.04 FGF13 ENST00000305414.8 ENST00000315930.11 8 subtype NORMAL connt **PRIMARY** ARPC DNPC **NEPC** 2 -0 . 15000 16000 13000 13000 14000 15000 16000 13000 14000 14000 15000 16000 HALLMARK_UV_RESPONSE_UP score

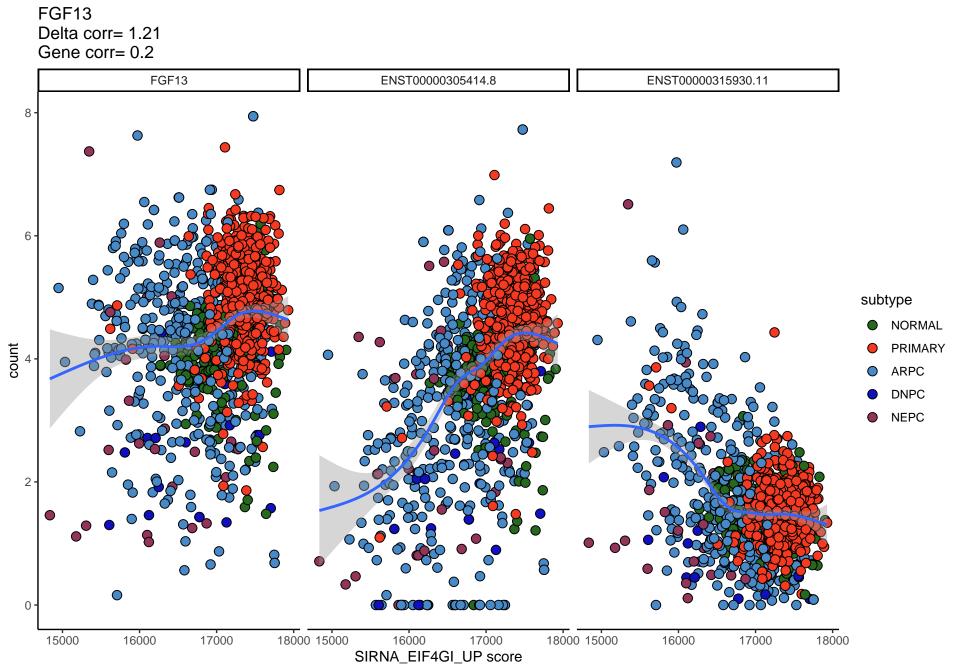
HALLMARK_UV_RESPONSE_UP ENST00000305414.8 - FGF13-201 PCC = 0.68 ENST00000315930.11 - FGF13-202 PCC = -0.56

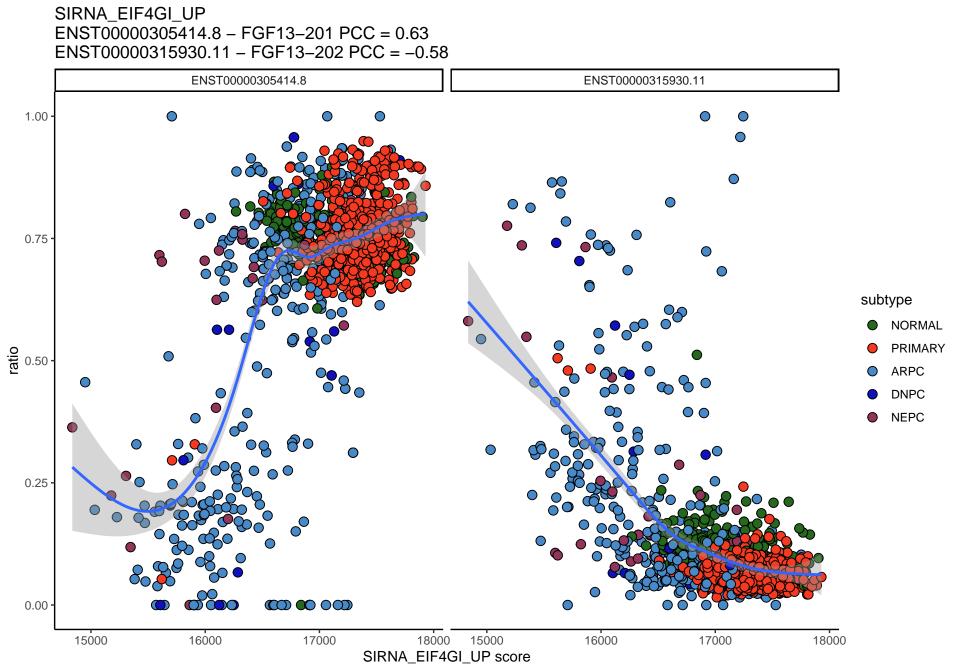


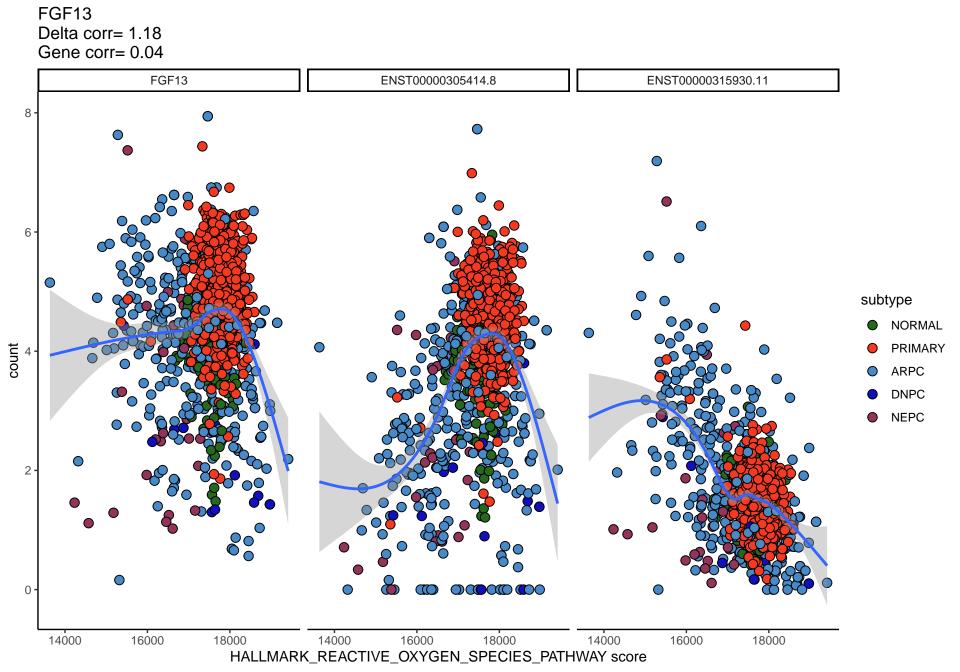


RB_DN.V1_UP ENST00000305414.8 - FGF13-201 PCC = 0.64 ENST00000315930.11 - FGF13-202 PCC = -0.58

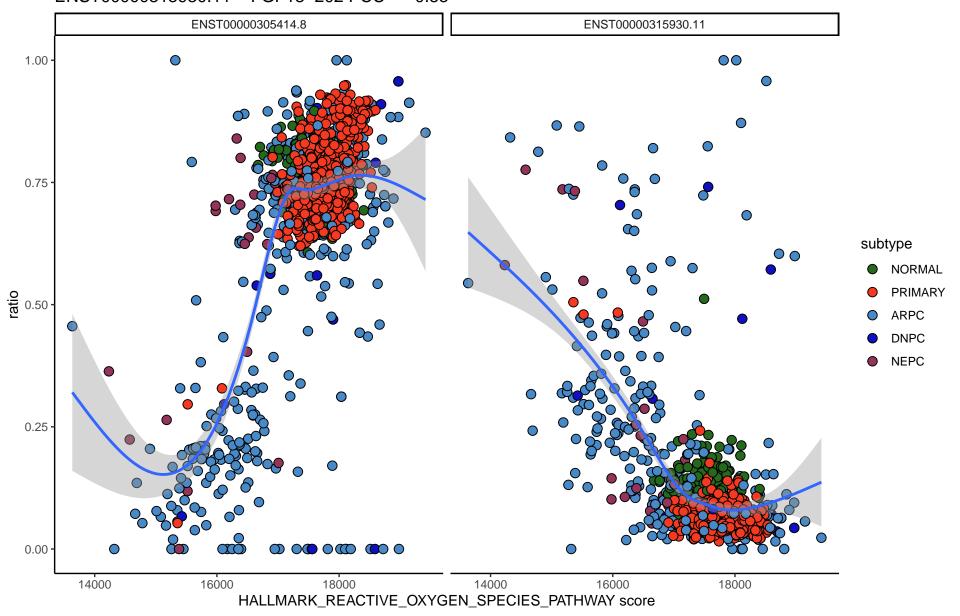


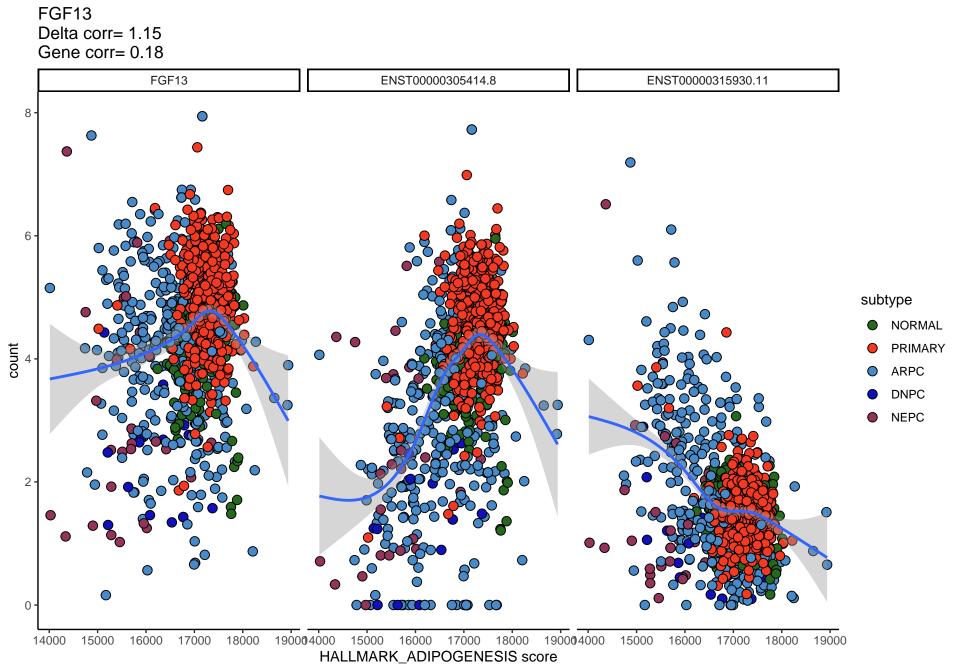




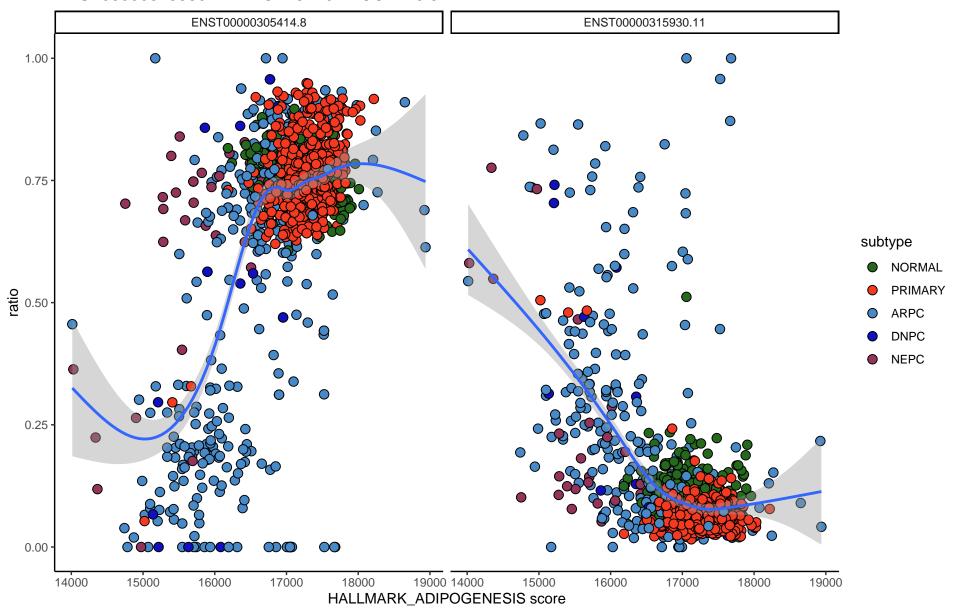


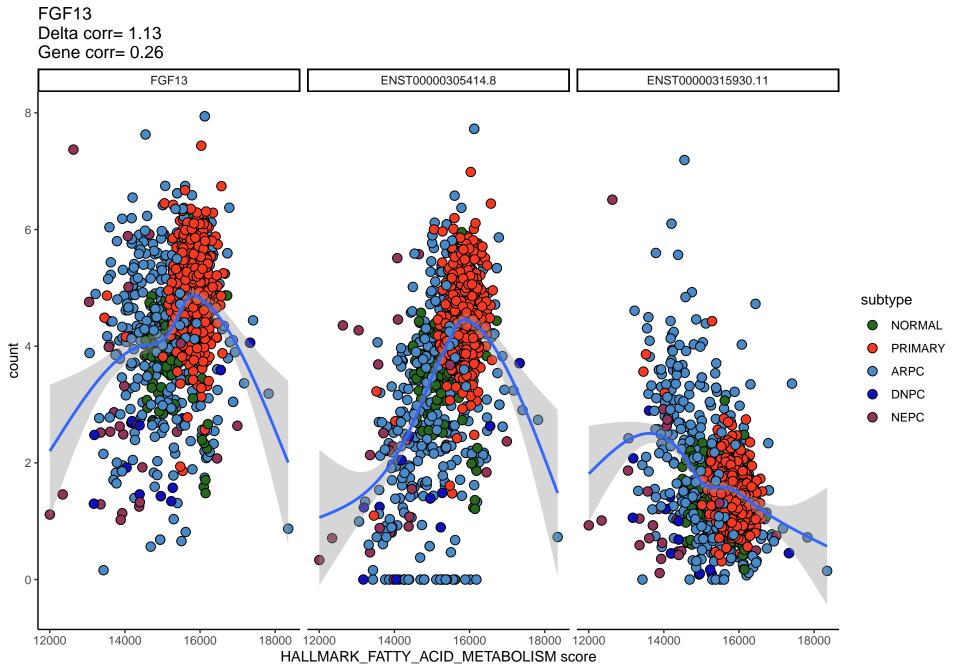
HALLMARK_REACTIVE_OXYGEN_SPECIES_PATHWAY ENST00000305414.8 - FGF13-201 PCC = 0.63 ENST00000315930.11 - FGF13-202 PCC = -0.55



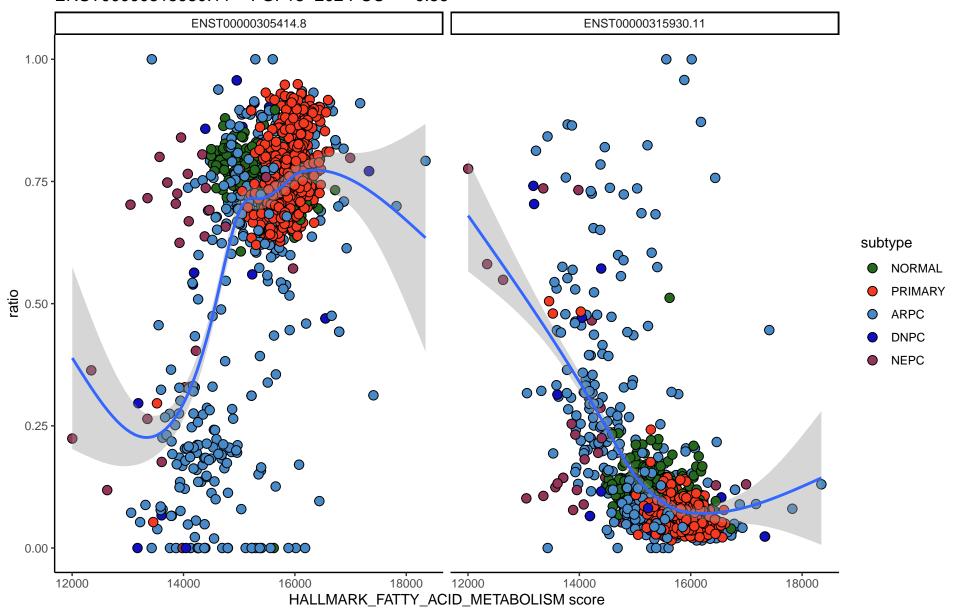


HALLMARK_ADIPOGENESIS ENST00000305414.8 - FGF13-201 PCC = 0.61 ENST00000315930.11 - FGF13-202 PCC = -0.54



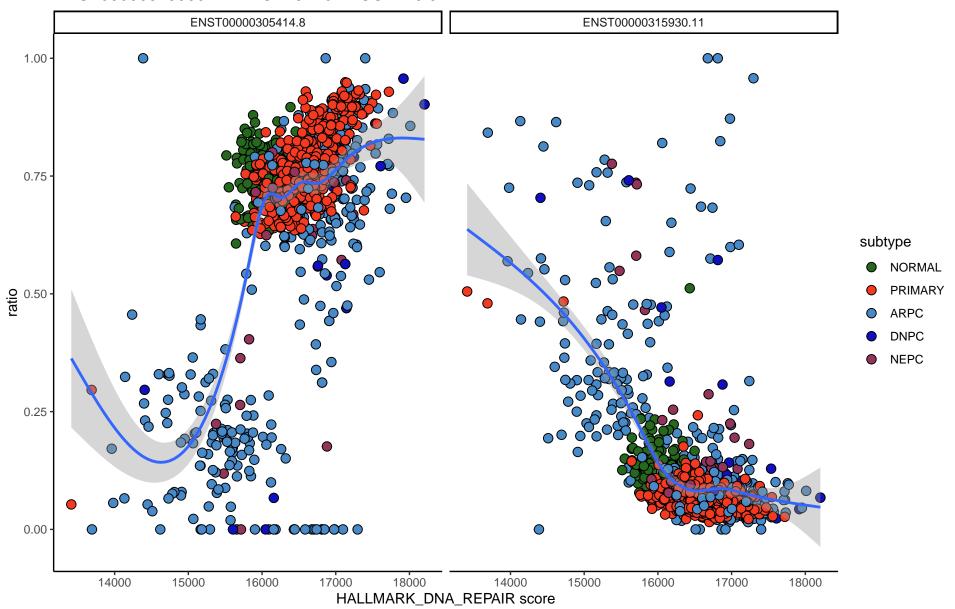


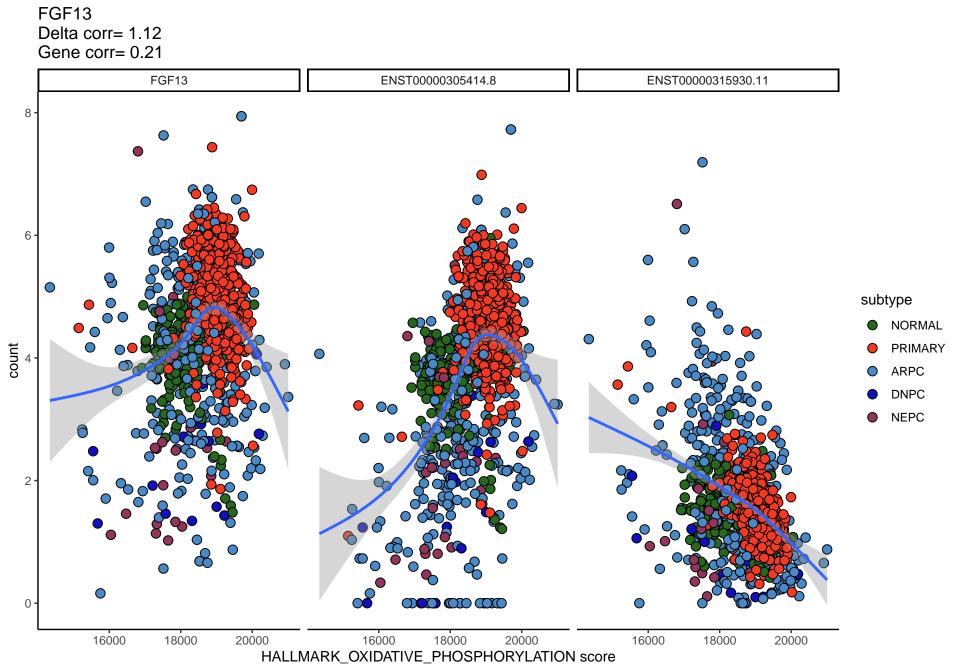
HALLMARK_FATTY_ACID_METABOLISM ENST00000305414.8 - FGF13-201 PCC = 0.57 ENST00000315930.11 - FGF13-202 PCC = -0.56



FGF13 Delta corr= 1.12 Gene corr= 0.04 FGF13 ENST00000305414.8 ENST00000315930.11 8 -6 subtype NORMAL count 4 **PRIMARY** ARPC DNPC NEPC 0 -14000 15000 16000 17000 18000 14000 15000 16000 17000 18000 14000 15000 16000 17000 18000 HALLMARK_DNA_REPAIR score

HALLMARK_DNA_REPAIR ENST00000305414.8 - FGF13-201 PCC = 0.58 ENST00000315930.11 - FGF13-202 PCC = -0.54





HALLMARK_OXIDATIVE_PHOSPHORYLATION ENST00000305414.8 - FGF13-201 PCC = 0.55 ENST00000315930.11 - FGF13-202 PCC = -0.56

