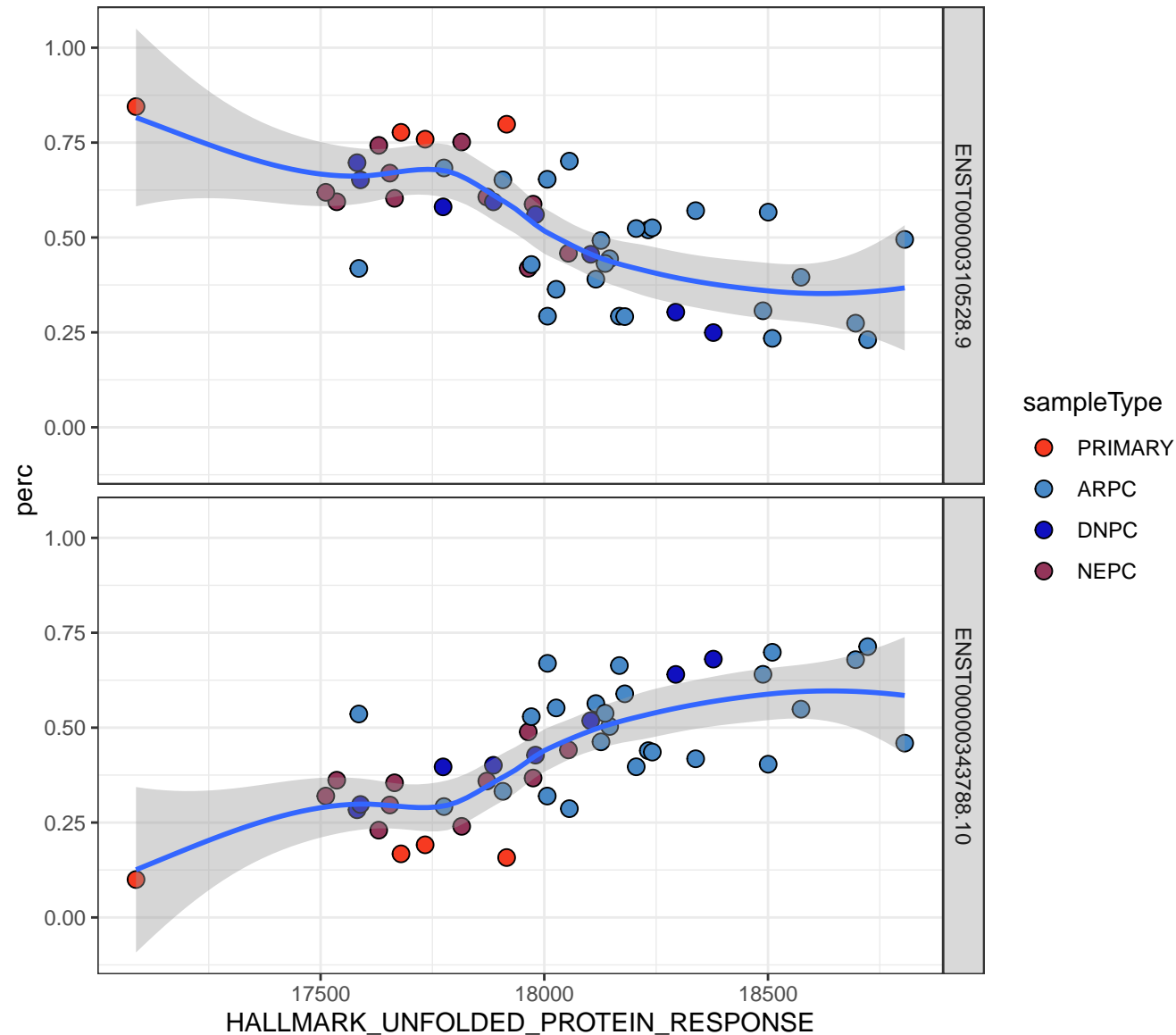


Gene = SUGT1

Target = HALLMARK\_UNFOLDED\_PROTEIN\_RESPONSE

ENST00000343788.10 (up) = 0.705698162598252

ENST00000310528.9 (down) = -0.688930902990535

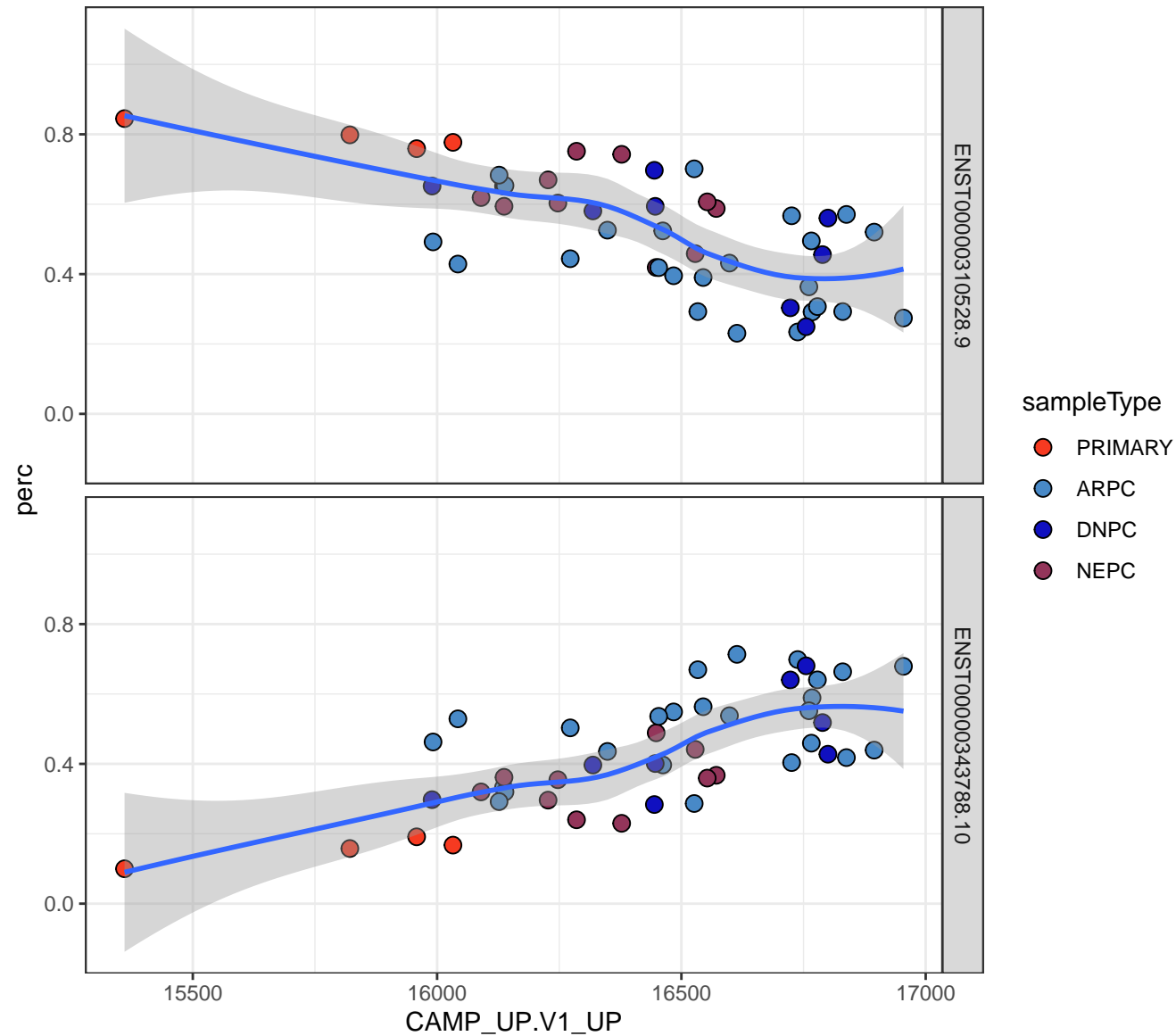


Gene = SUGT1

Target = CAMP\_UP.V1\_UP

ENST00000343788.10 (up) = 0.708865550785823

ENST00000310528.9 (down) = -0.679935323367582

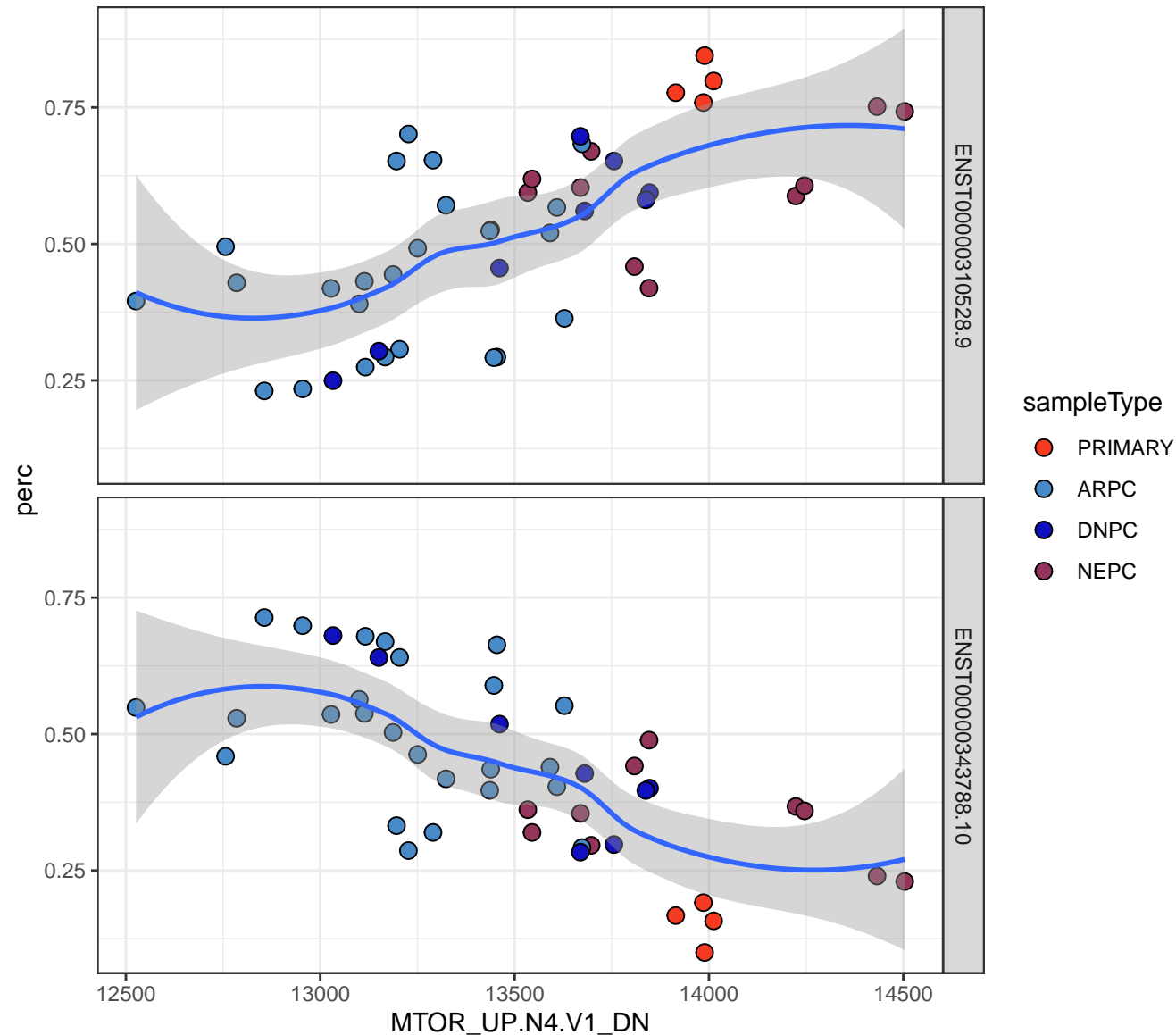


Gene = SUGT1

Target = MTOR\_UP.N4.V1\_DN

ENST00000310528.9 (up) = 0.665164336877199

ENST00000343788.10 (down) = -0.683838610278662

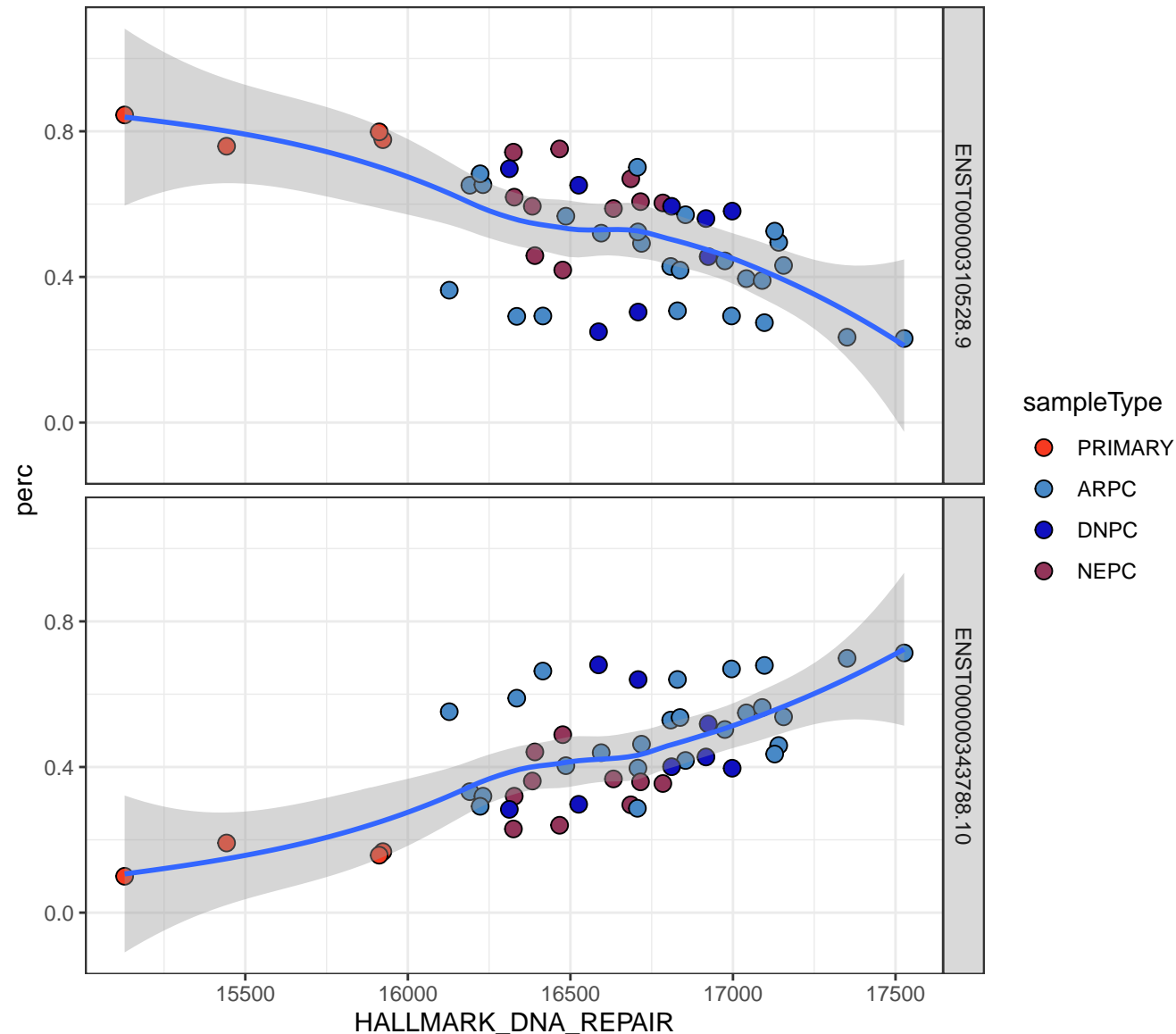


Gene = SUGT1

Target = HALLMARK\_DNA\_REPAIR

ENST00000343788.10 (up) = 0.683804509706719

ENST00000310528.9 (down) = -0.627270497321838

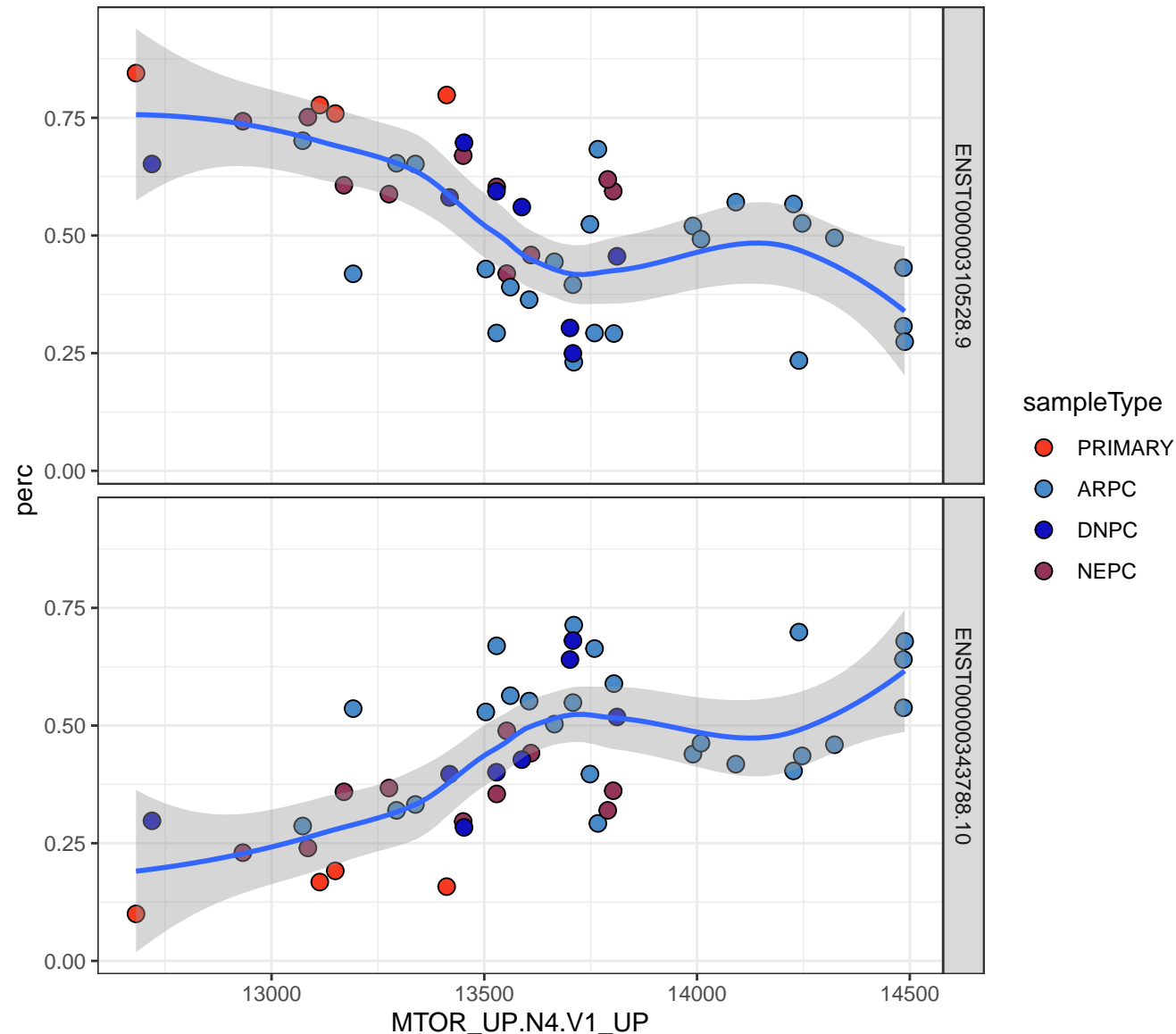


Gene = SUGT1

Target = MTOR\_UP.N4.V1\_UP

ENST00000343788.10 (up) = 0.607027359023741

ENST00000310528.9 (down) = -0.589677990128392

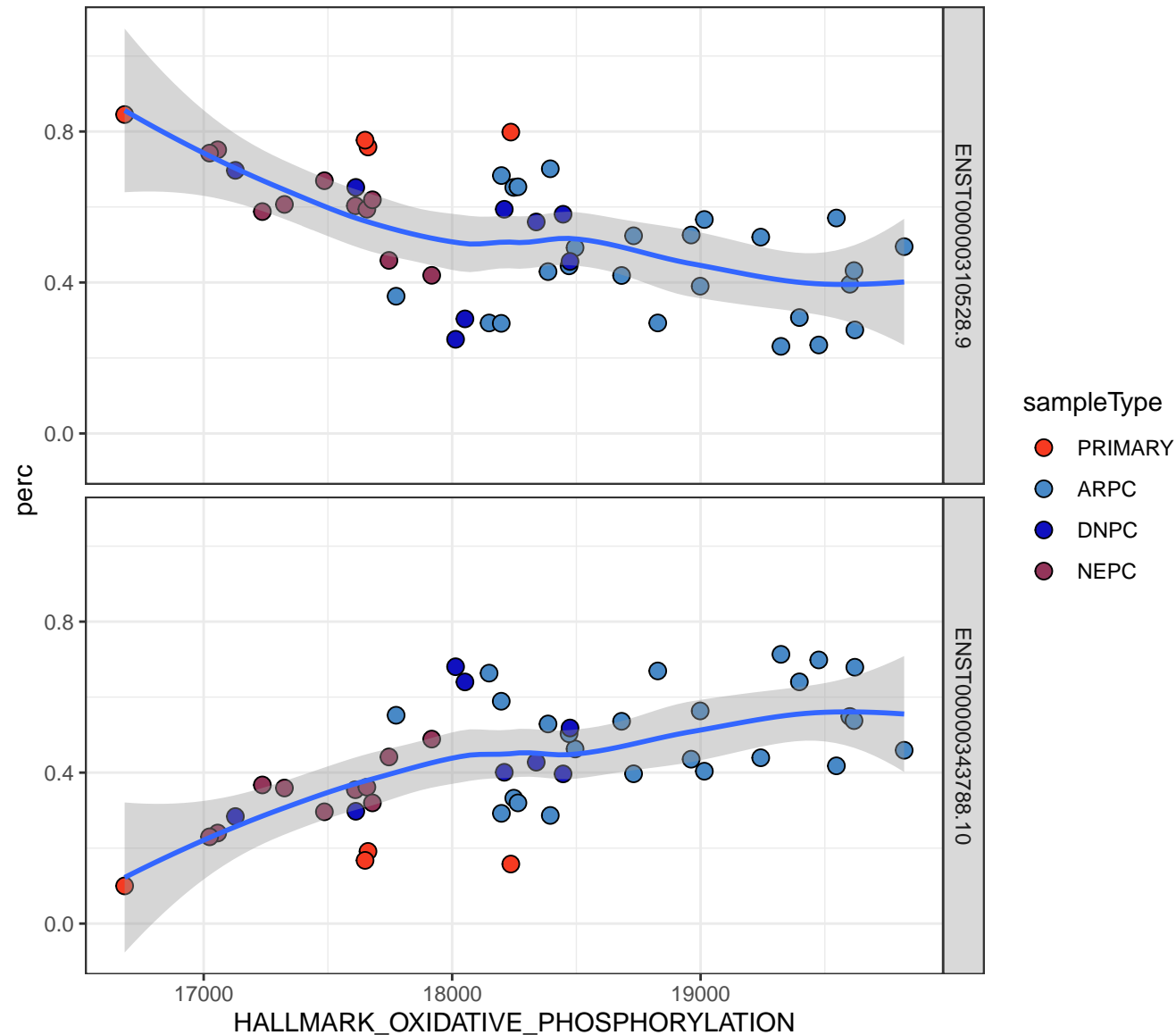


Gene = SUGT1

Target = HALLMARK\_OXIDATIVE\_PHOSPHORYLATION

ENST00000343788.10 (up) = 0.615695902370469

ENST00000310528.9 (down) = -0.578485062213859

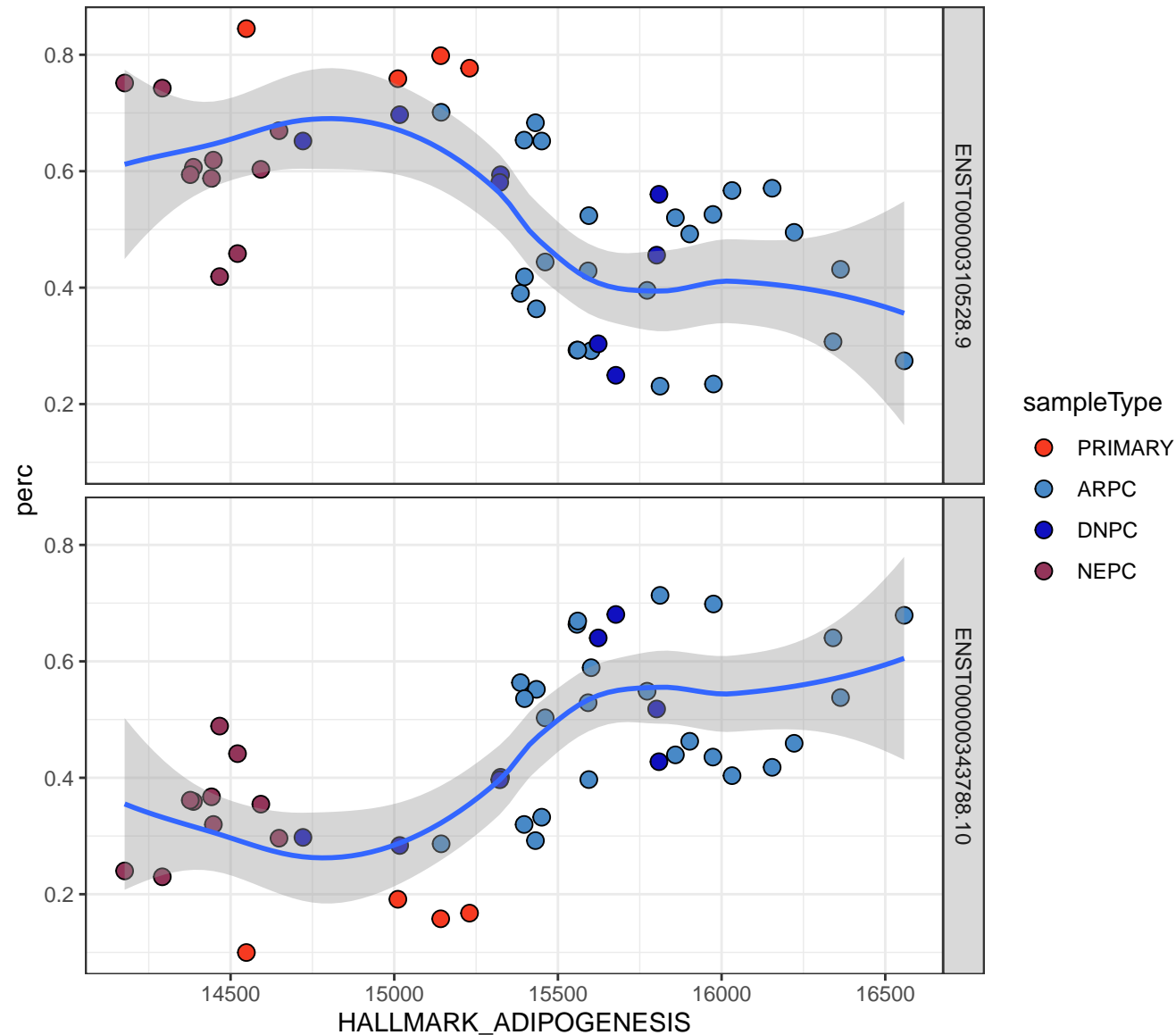


Gene = SUGT1

Target = HALLMARK\_ADIPOGENESIS

ENST00000343788.10 (up) = 0.60669322248511

ENST00000310528.9 (down) = -0.565925740249386

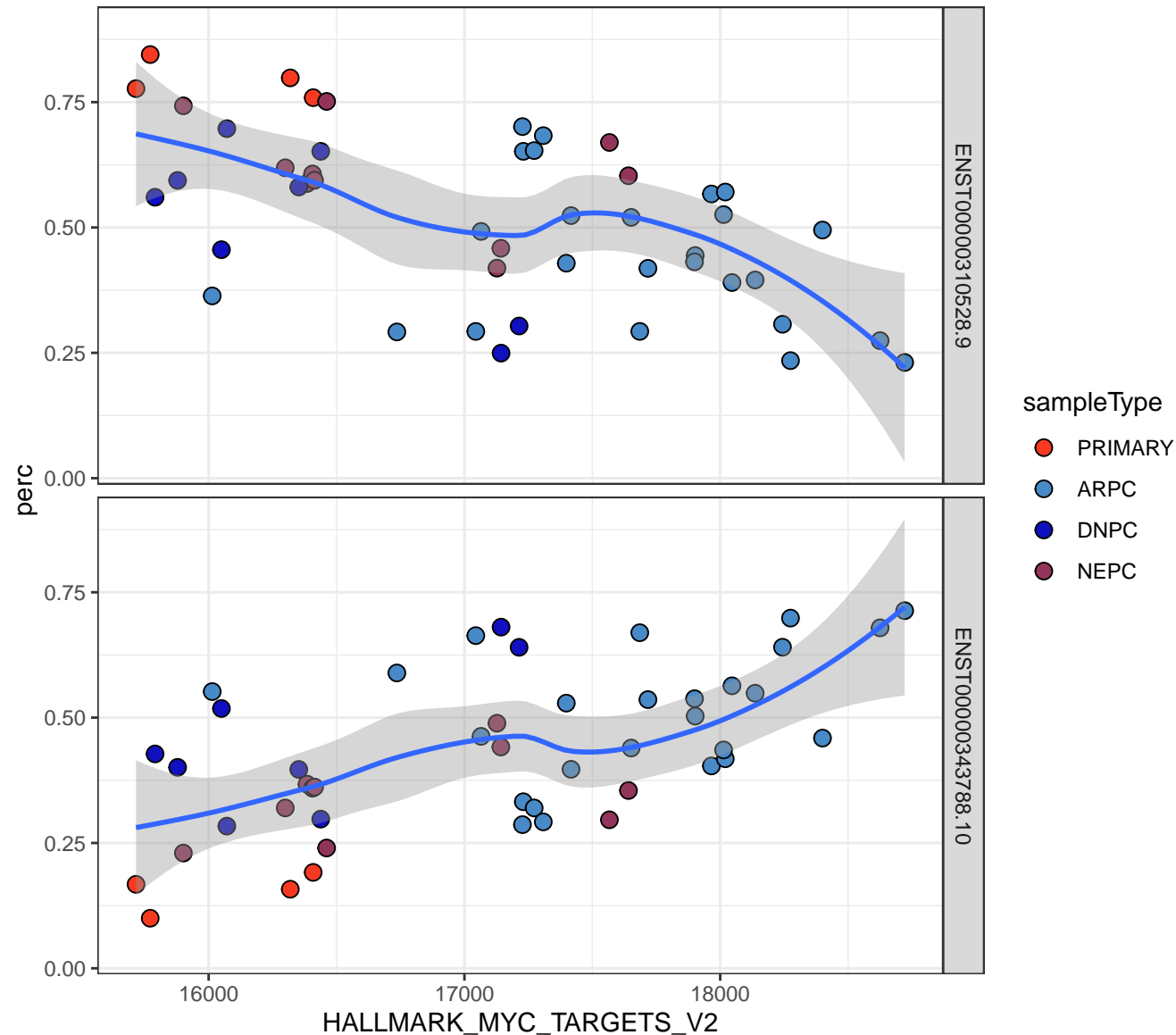


Gene = SUGT1

Target = HALLMARK\_MYC\_TARGETS\_V2

ENST00000343788.10 (up) = 0.593910914468131

ENST00000310528.9 (down) = -0.574253932469995



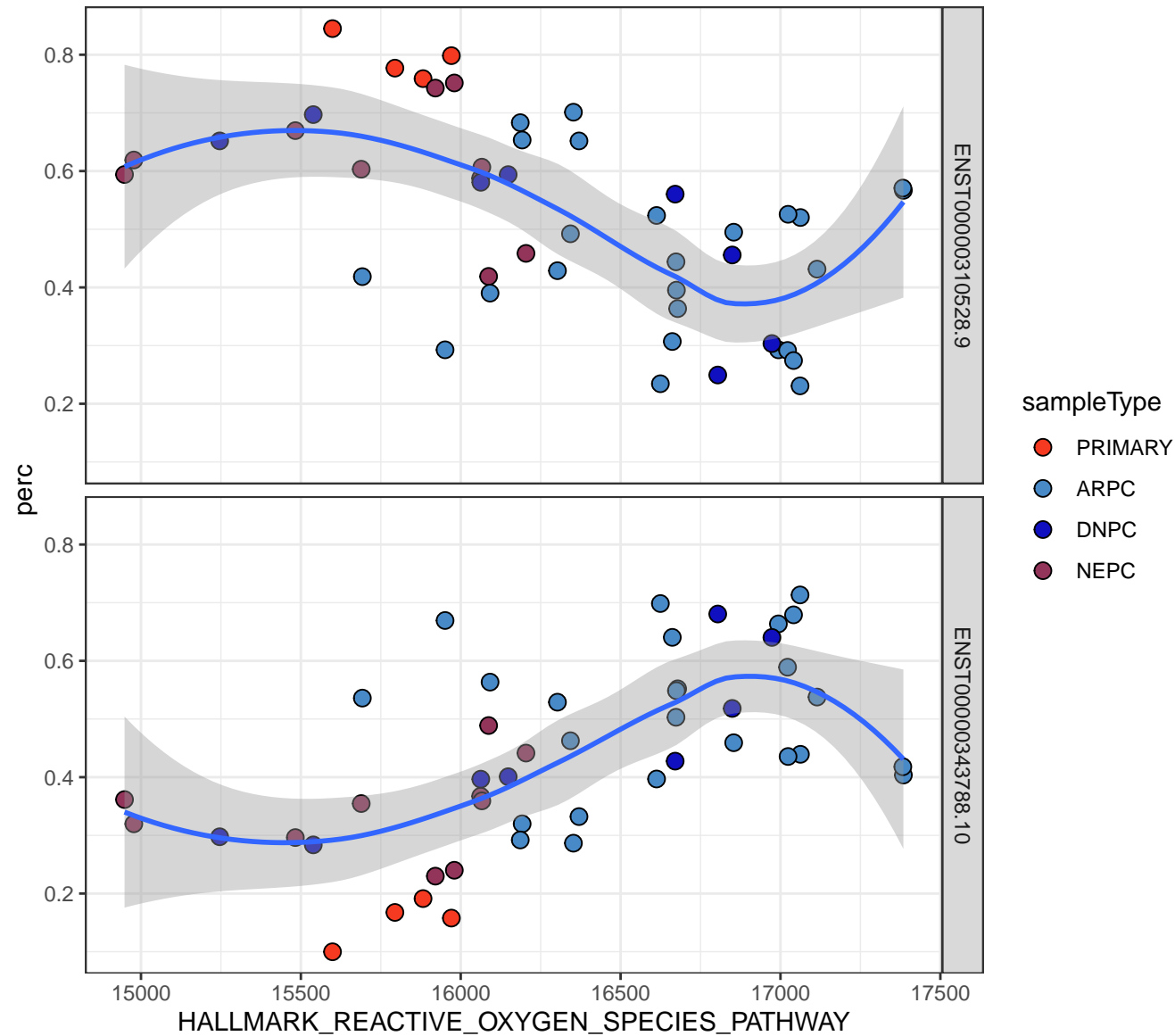


Gene = SUGT1

Target = HALLMARK\_REACTIVE\_OXYGEN\_SPECIES\_PATHWAY

ENST00000343788.10 (up) = 0.579442411062404

ENST00000310528.9 (down) = -0.555018790142142



Gene = SUGT1

Target = HALLMARK\_MYC\_TARGETS\_V1

ENST00000343788.10 (up) = 0.55956862320791

ENST00000310528.9 (down) = -0.518838095047805

