

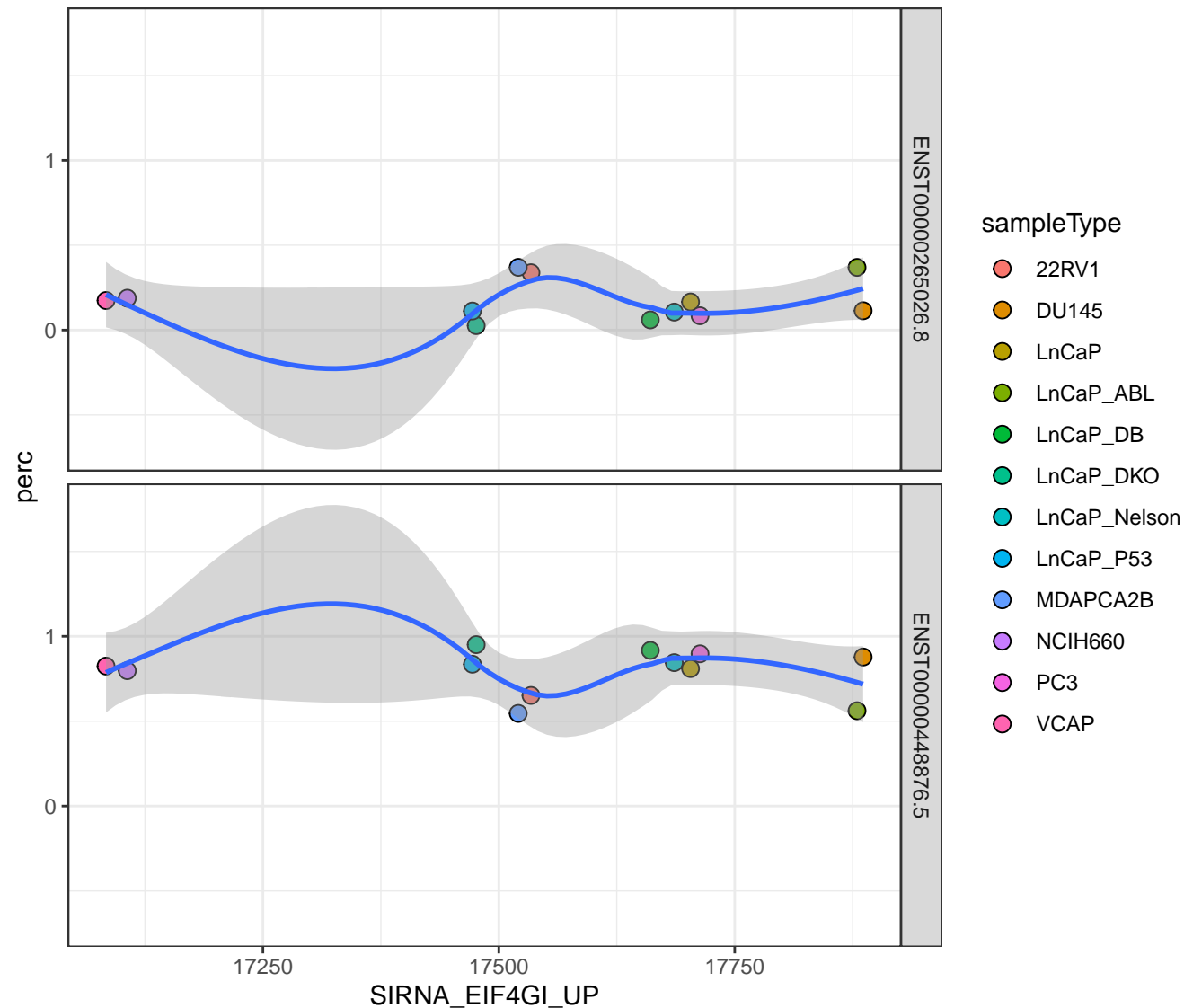
CellLine

Gene = MAP3K13

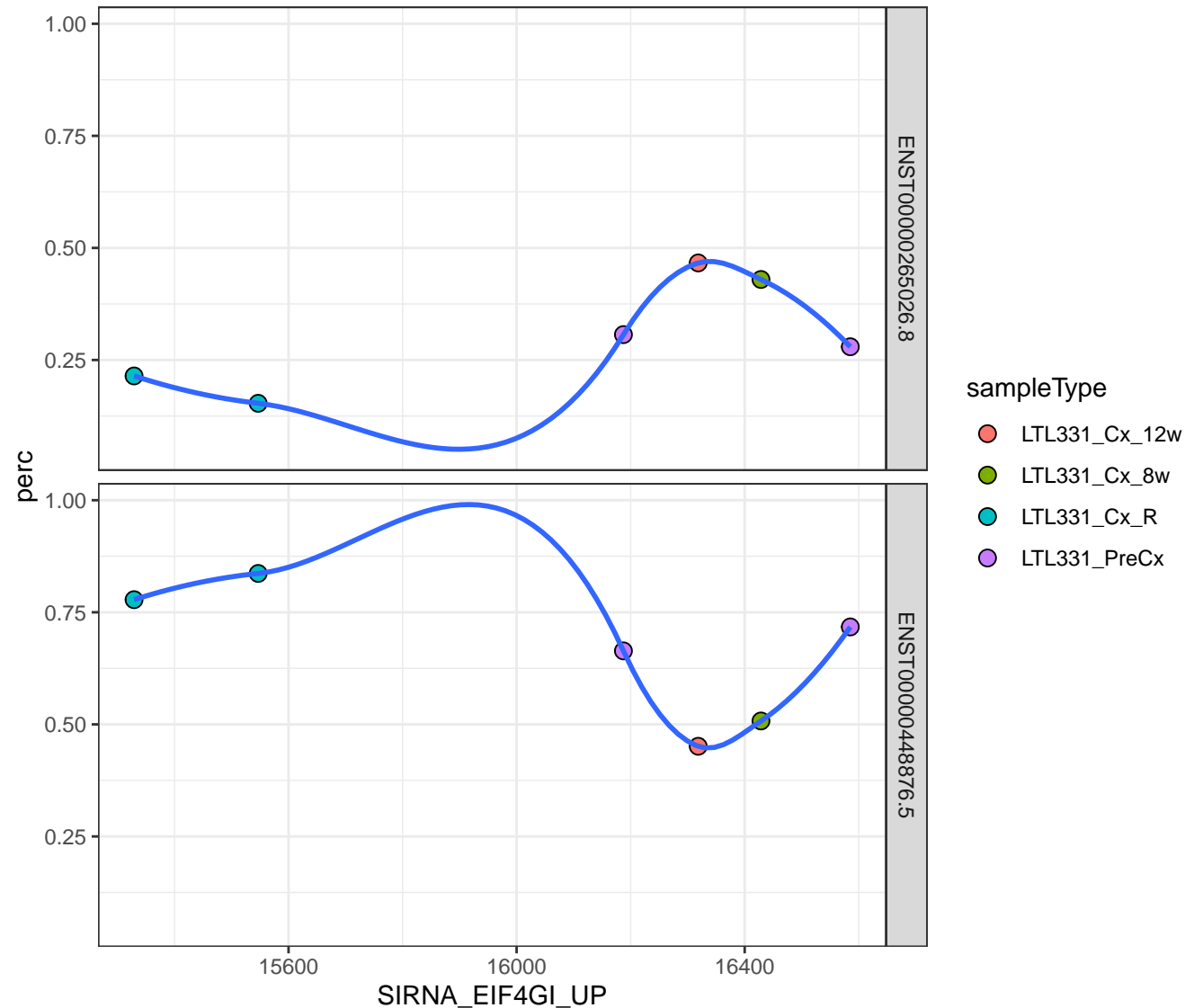
Target = SIRNA_EIF4GI_UP

ENST00000448876.5 (up) = -0.0776760244592958

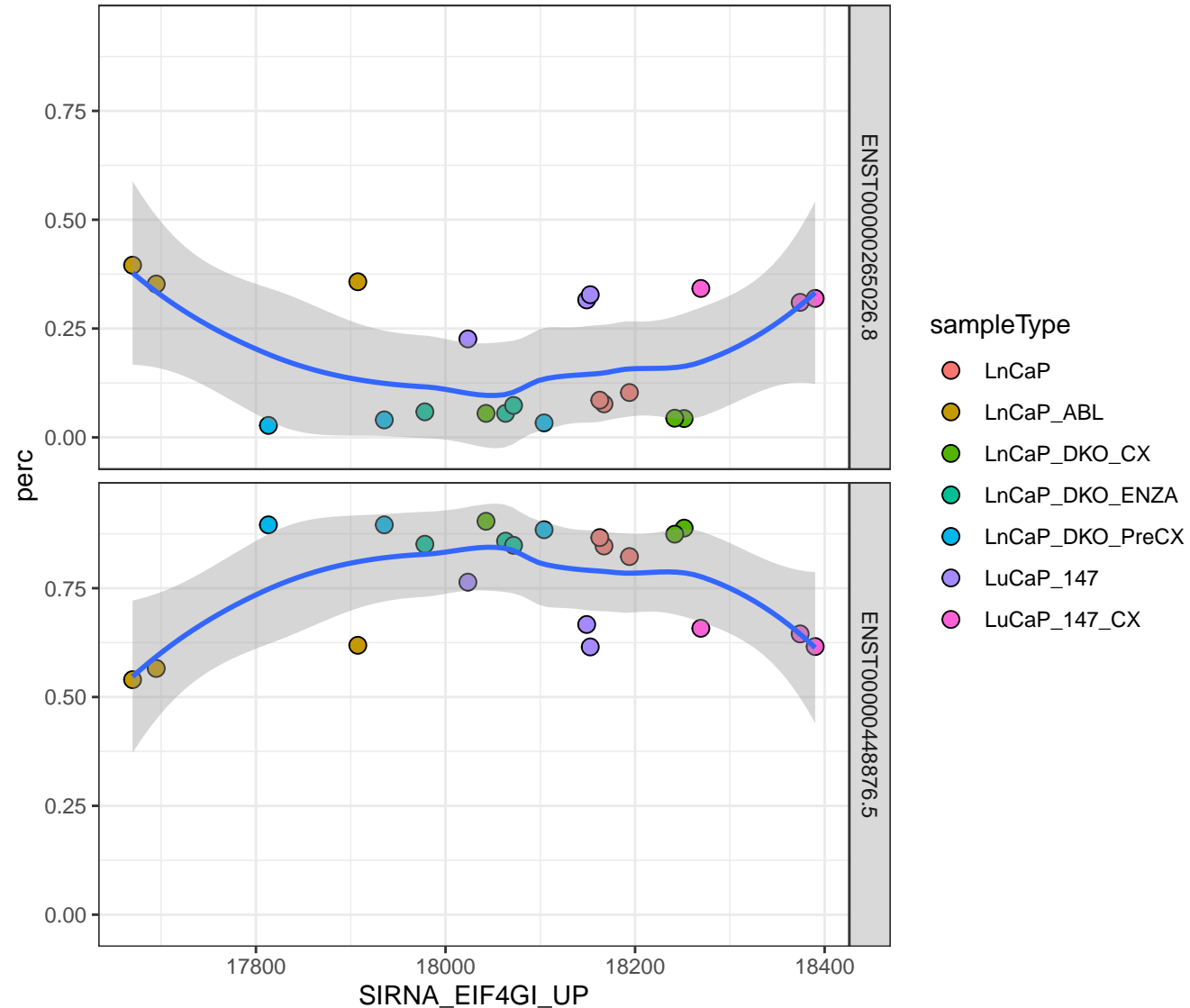
ENST00000265026.8 (down) = 0.0214500348271675



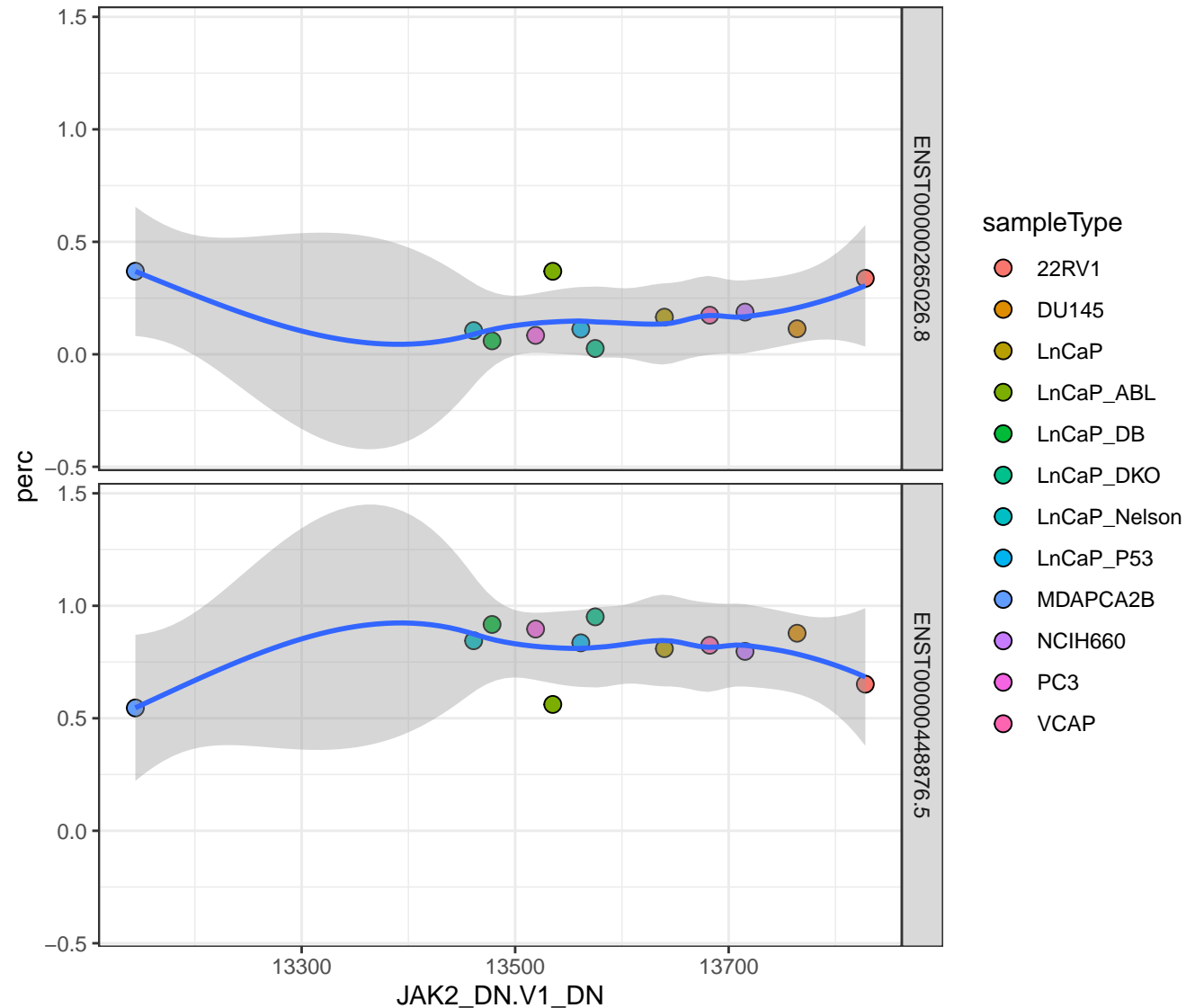
LTL_longitudinal
Gene = MAP3K13
Target = SIRNA_EIF4GI_UP
ENST00000448876.5 (up) = -0.664610150531121
ENST00000265026.8 (down) = 0.709150124858977



PDX_longitudinal
 Gene = MAP3K13
 Target = SIRNA_EIF4GI_UP
 ENST00000448876.5 (up) = 0.137008303422035
 ENST00000265026.8 (down) = -0.0900732399819509



CellLine
 Gene = MAP3K13
 Target = JAK2_DN.V1_DN
 ENST00000265026.8 (up) = -0.152708672154312
 ENST00000448876.5 (down) = 0.291962078609515



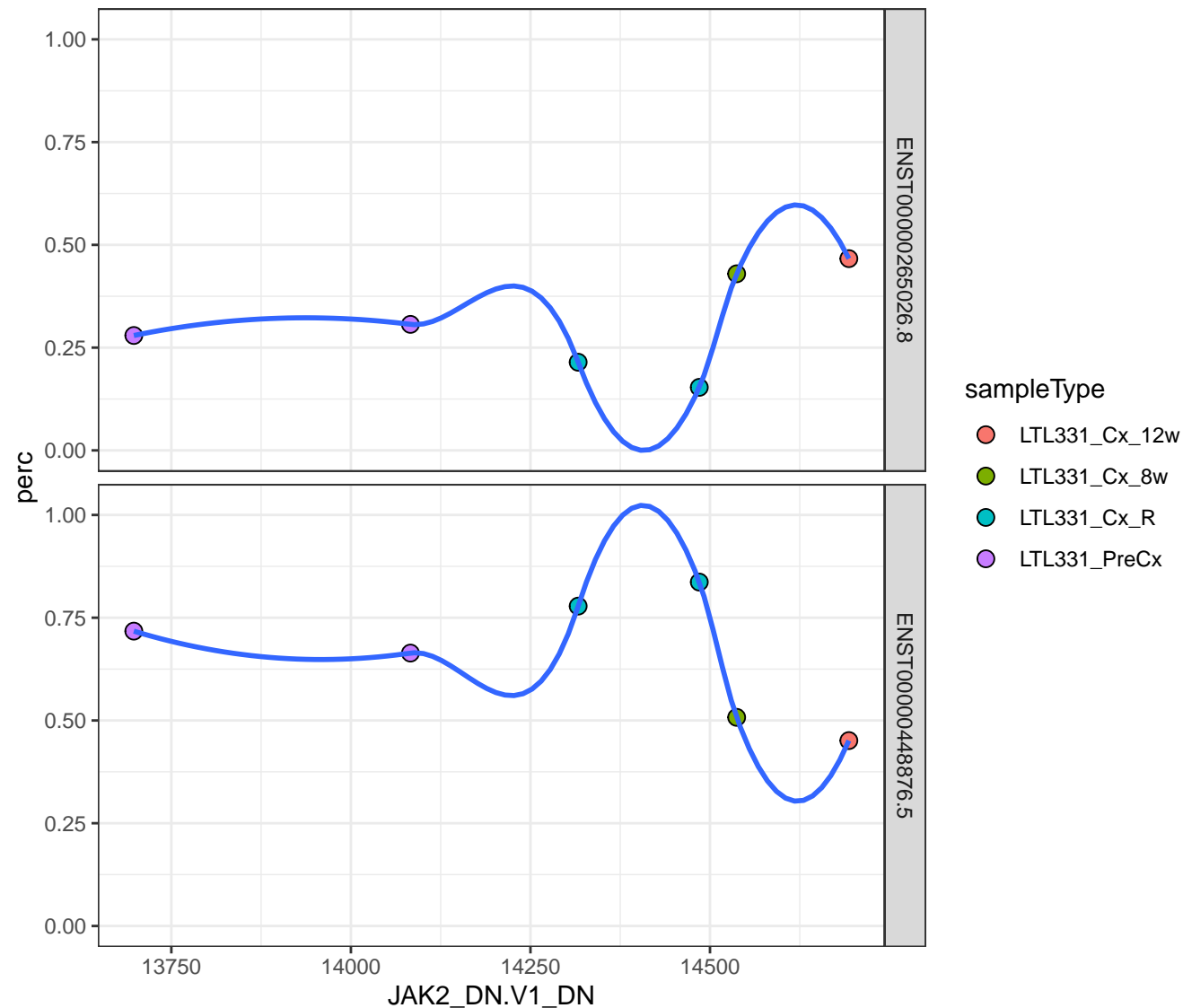
LTL_longitudinal

Gene = MAP3K13

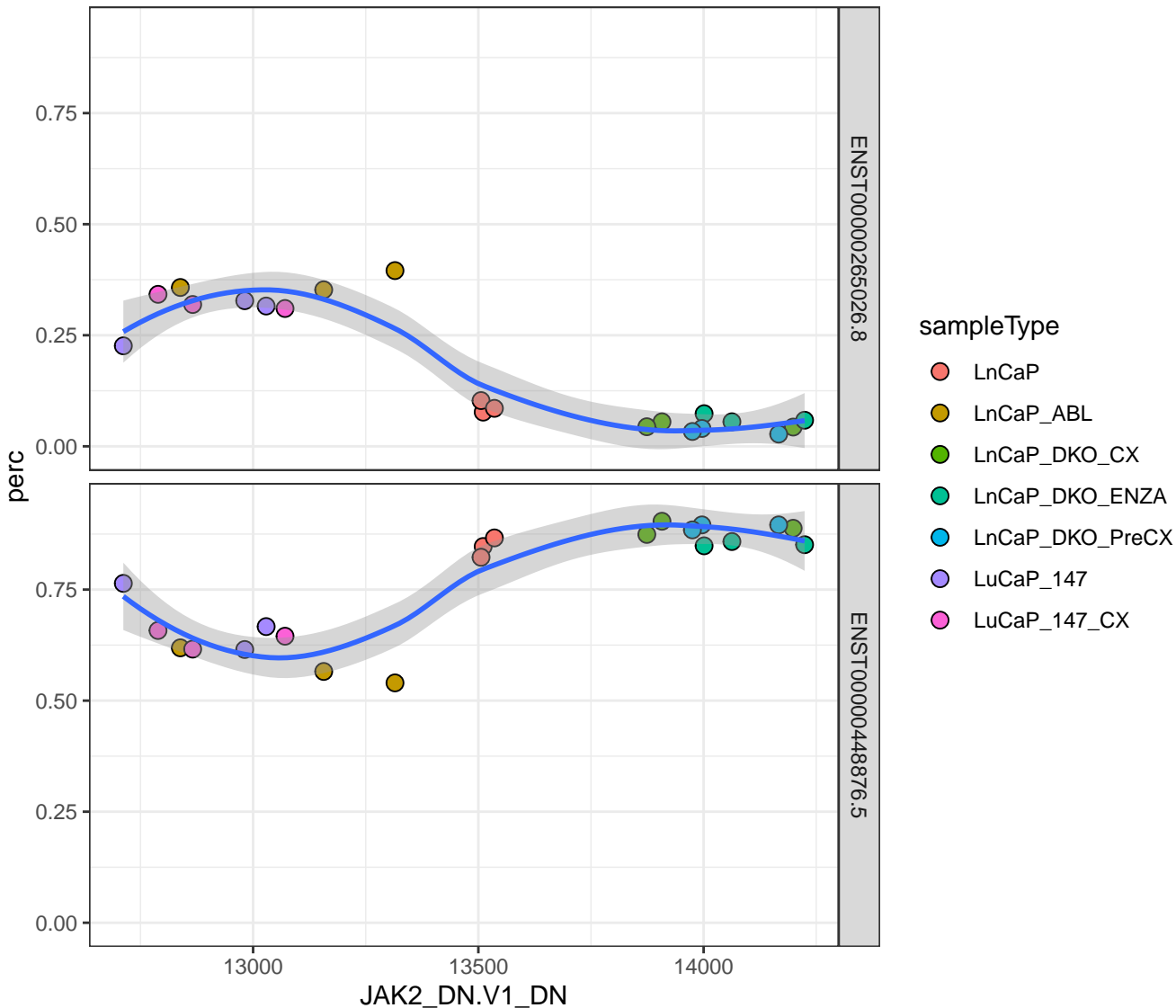
Target = JAK2_DN.V1_DN

ENST00000265026.8 (up) = 0.357853218754399

ENST00000448876.5 (down) = -0.432985415844348



PDX_longitudinal
Gene = MAP3K13
Target = JAK2_DN.V1_DN
ENST00000265026.8 (up) = -0.877245099536979
ENST00000448876.5 (down) = 0.814499341115142



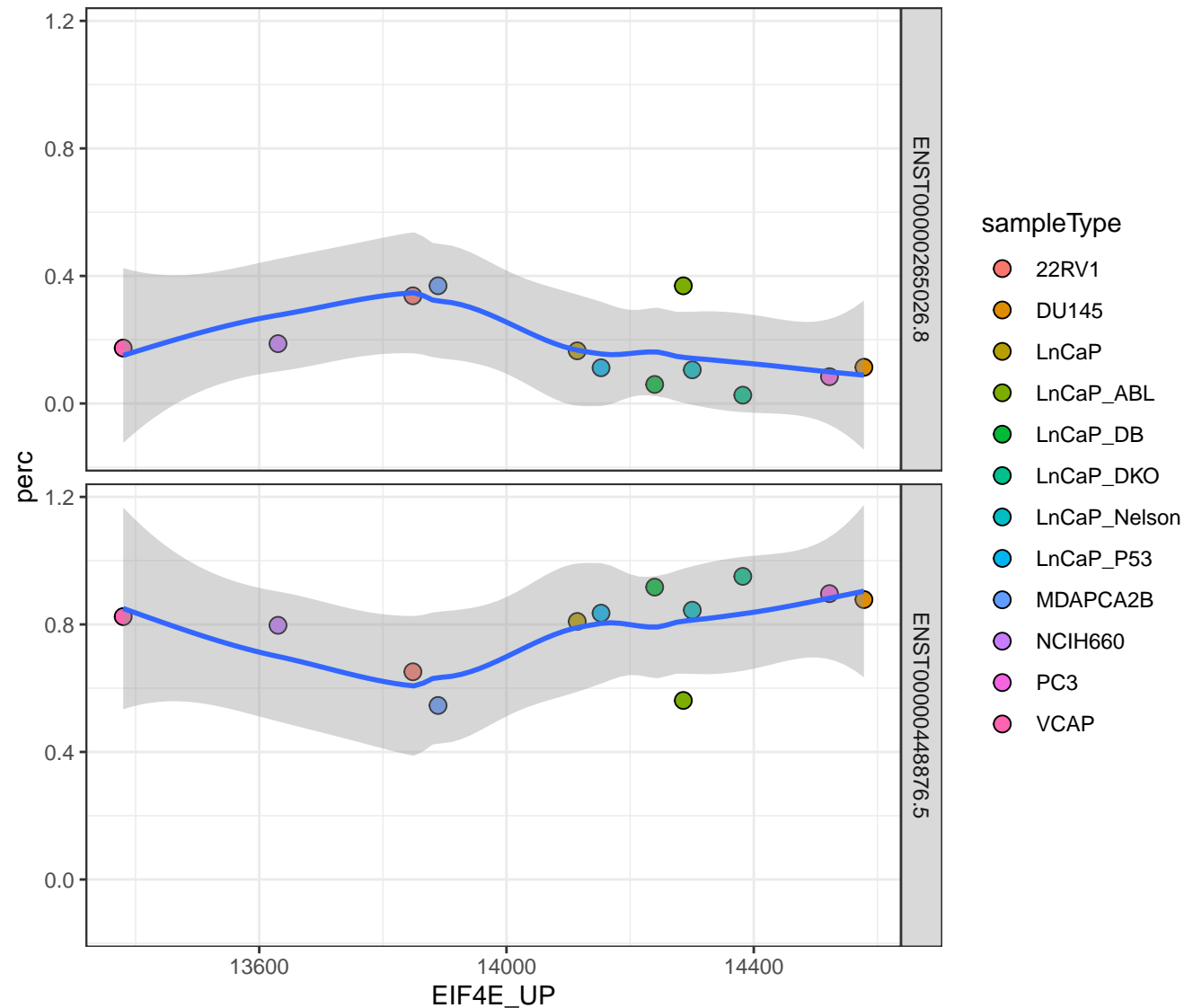
CellLine

Gene = MAP3K13

Target = EIF4E_UP

ENST00000448876.5 (up) = 0.336044109027338

ENST00000265026.8 (down) = -0.408160110406909



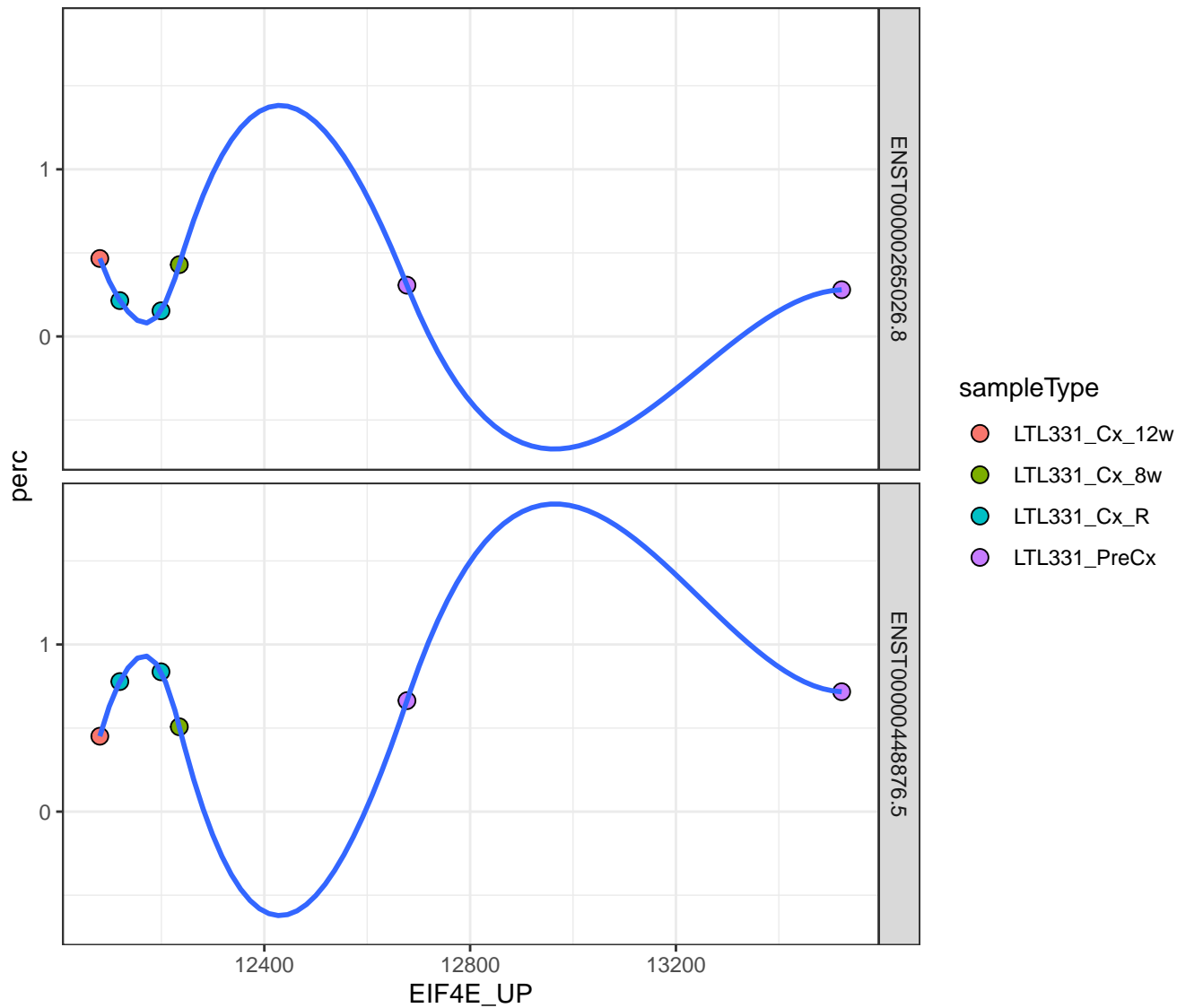
LTL_longitudinal

Gene = MAP3K13

Target = EIF4E_UP

ENST00000448876.5 (up) = 0.210480783069331

ENST00000265026.8 (down) = -0.1360579497821



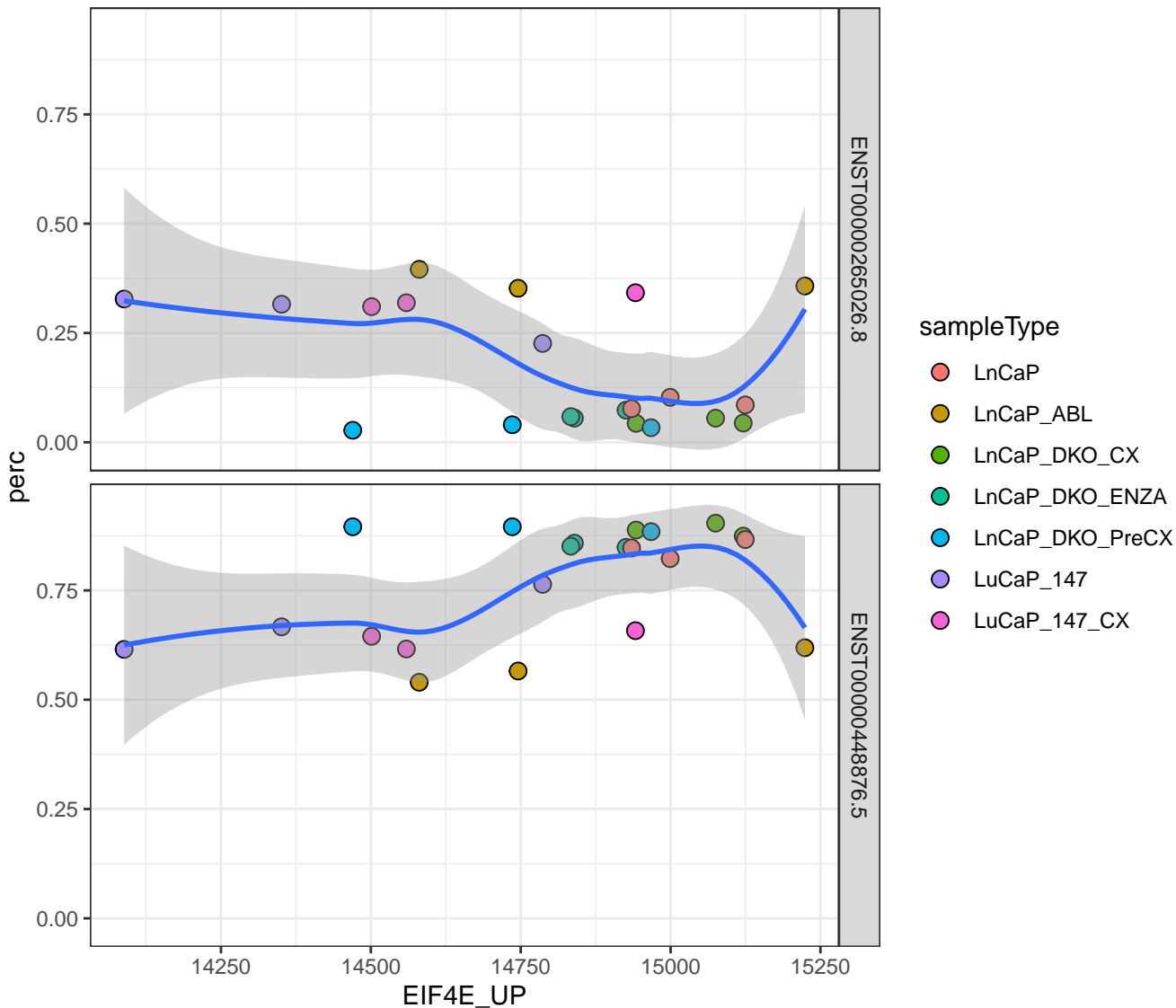
PDX_longitudinal

Gene = MAP3K13

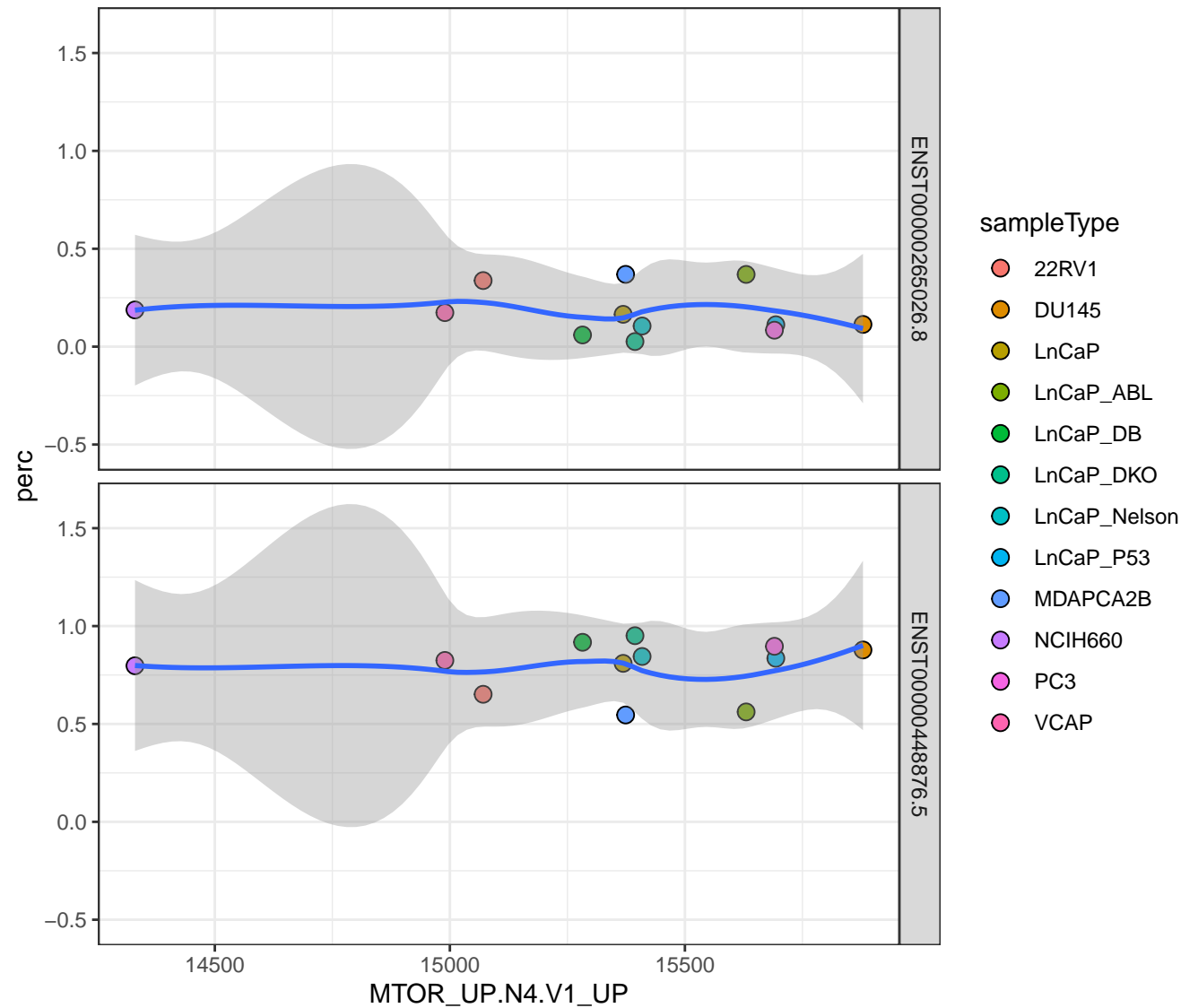
Target = EIF4E_UP

ENST00000448876.5 (up) = 0.460678553482726

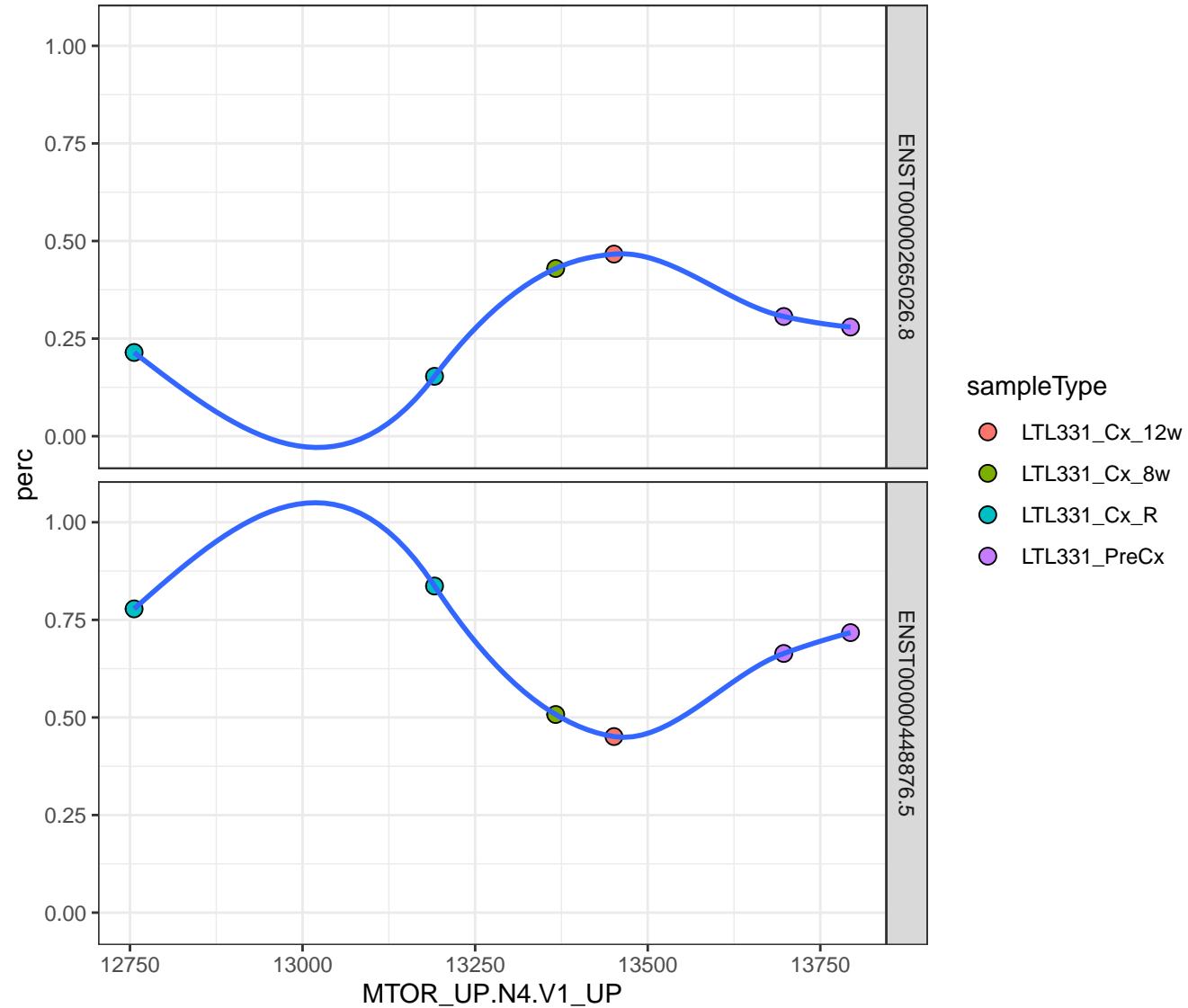
ENST00000265026.8 (down) = -0.427218016728387



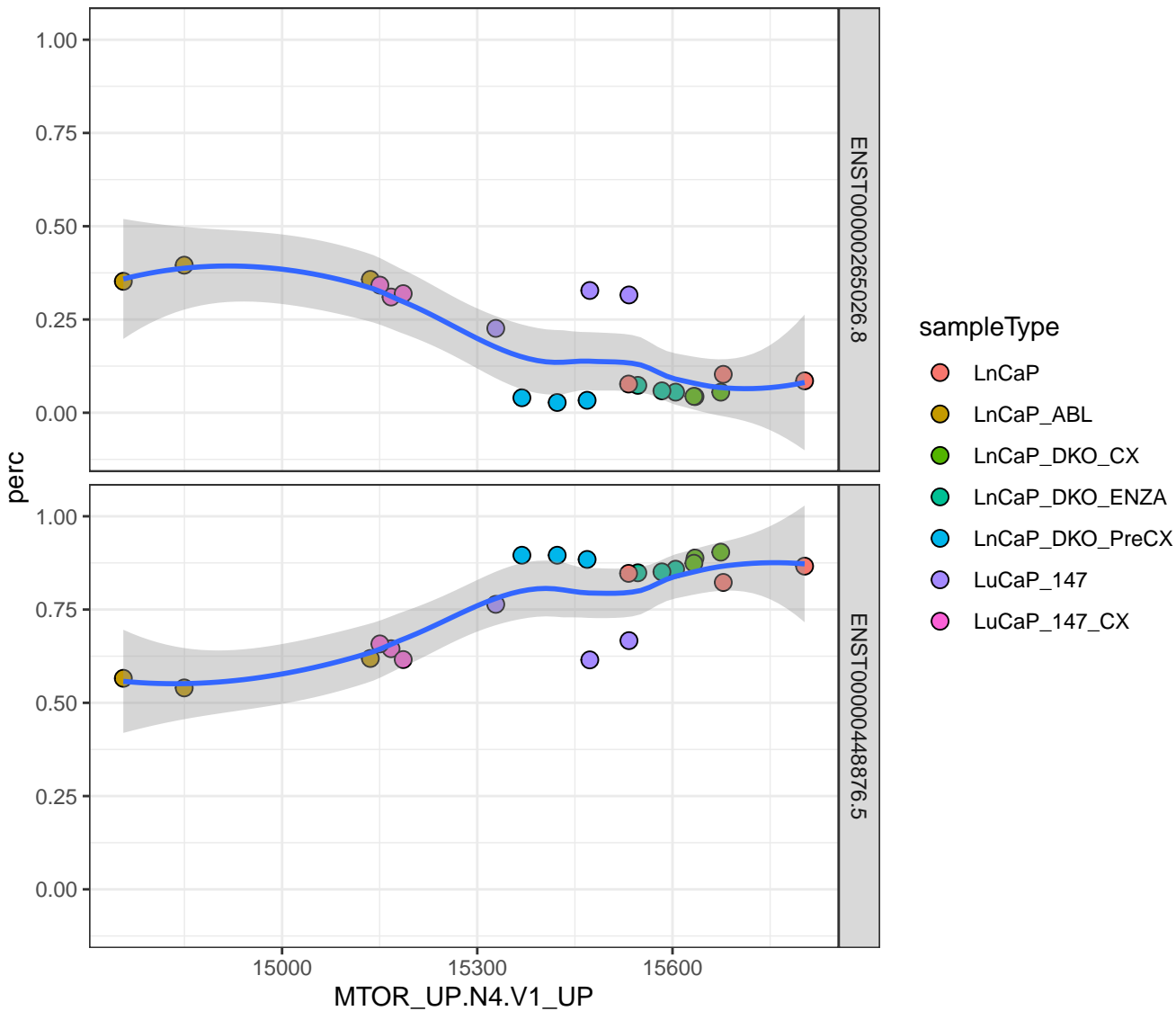
CellLine
 Gene = MAP3K13
 Target = MTOR_UP.N4.V1_UP
 ENST00000448876.5 (up) = 0.0818778895603529
 ENST00000265026.8 (down) = -0.160629839979234



LTL_longitudinal
Gene = MAP3K13
Target = MTOR_UP.N4.V1_UP
ENST00000448876.5 (up) = -0.3338127477249
ENST00000265026.8 (down) = 0.374490718254445



ENST00000265026.8 (down) = -0.765635625849923



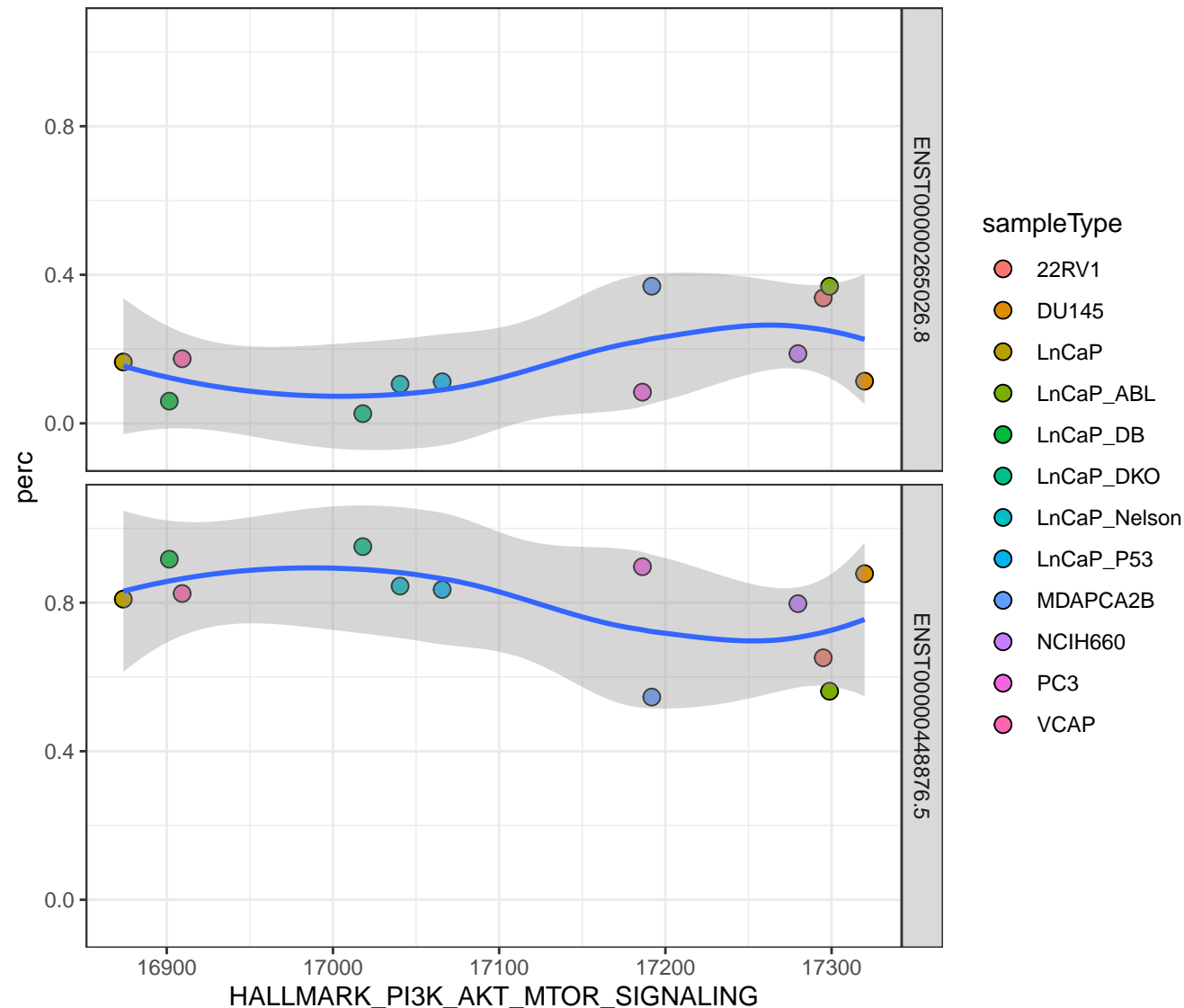
CellLine

Gene = MAP3K13

Target = HALLMARK_PI3K_AKT_MTOR_SIGNALING

ENST00000448876.5 (up) = -0.48245670107779

ENST00000265026.8 (down) = 0.509450658322378



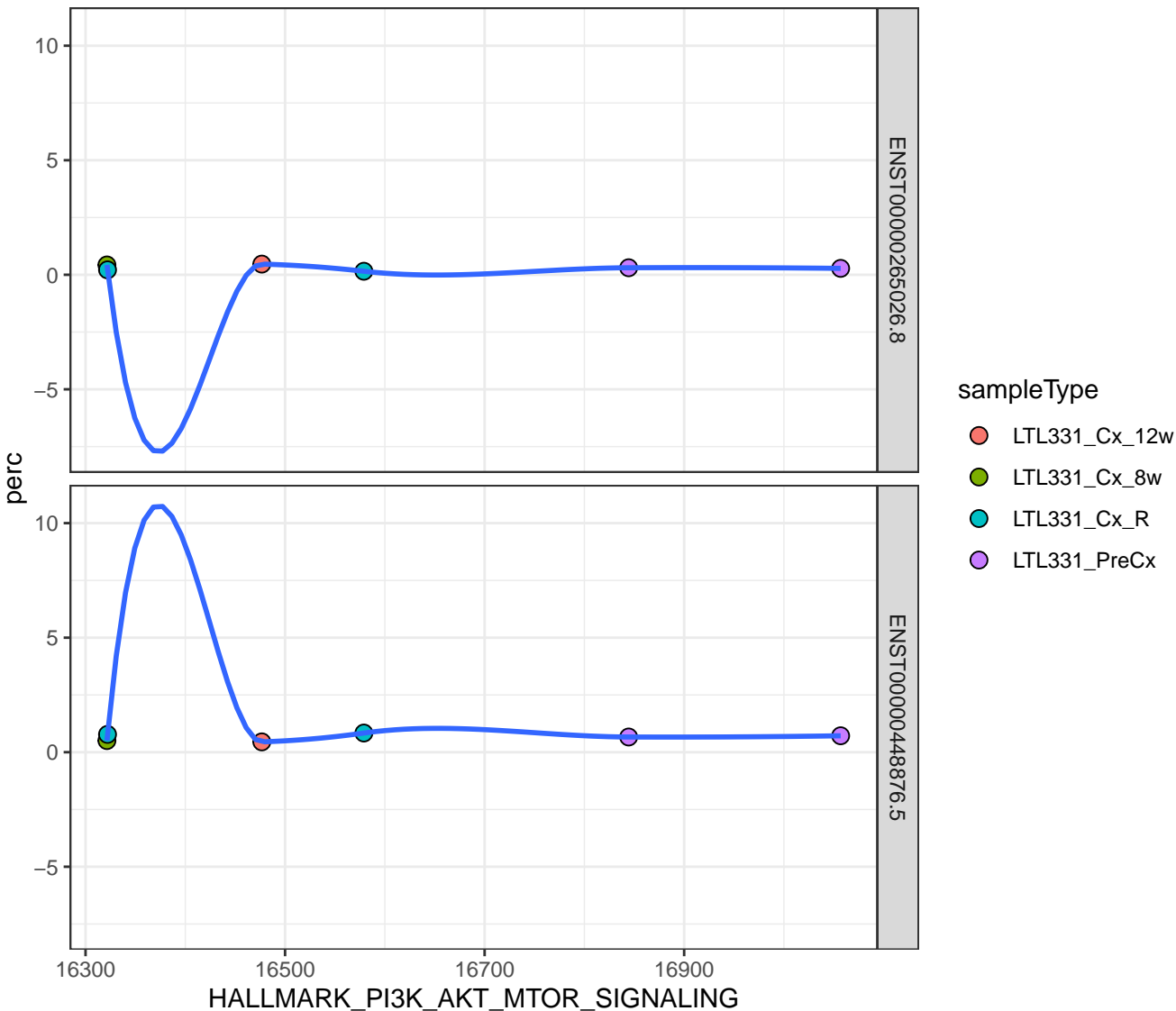
LTL_longitudinal

Gene = MAP3K13

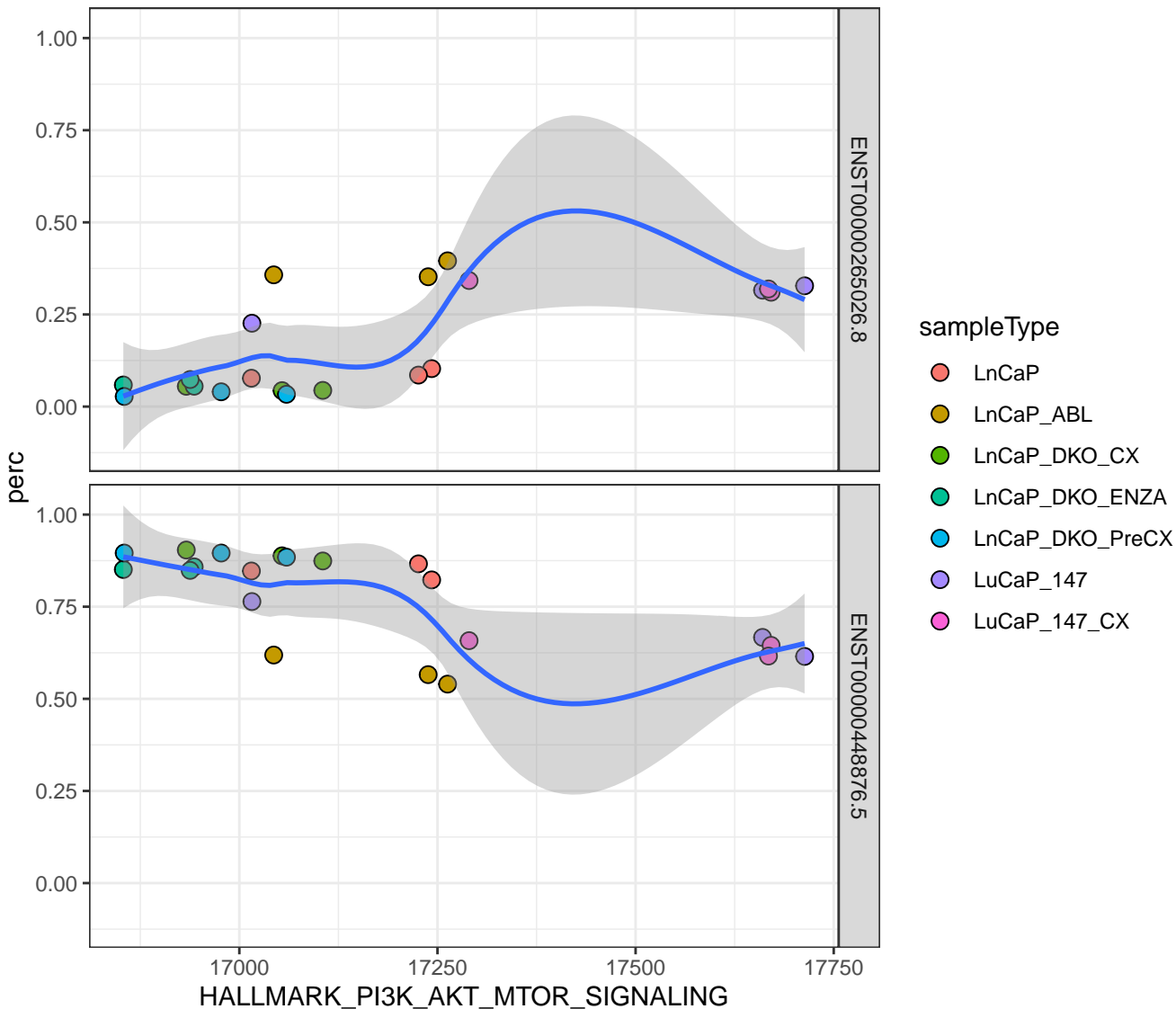
Target = HALLMARK_PI3K_AKT_MTOR_SIGNALING

ENST00000448876.5 (up) = 0.261339919912792

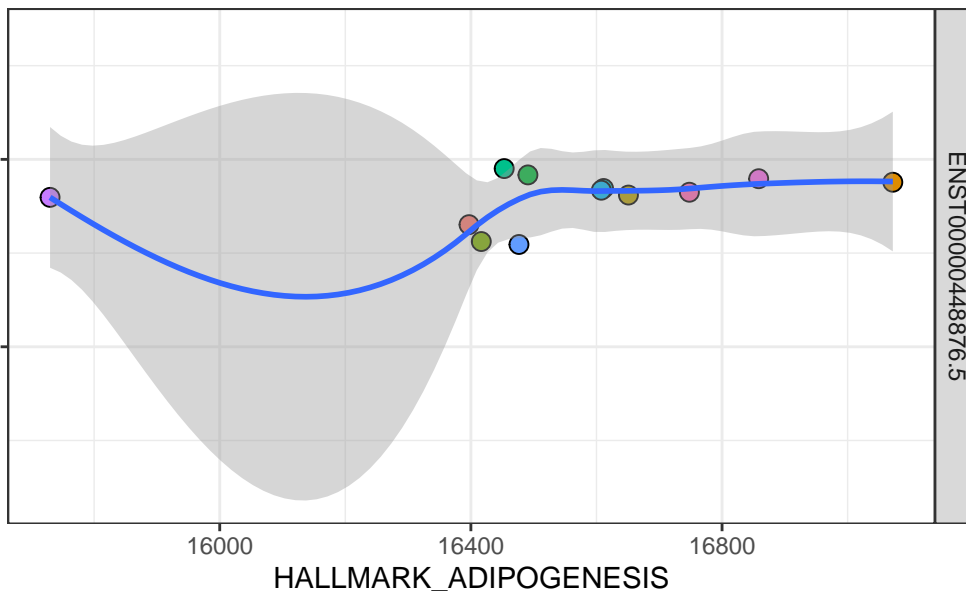
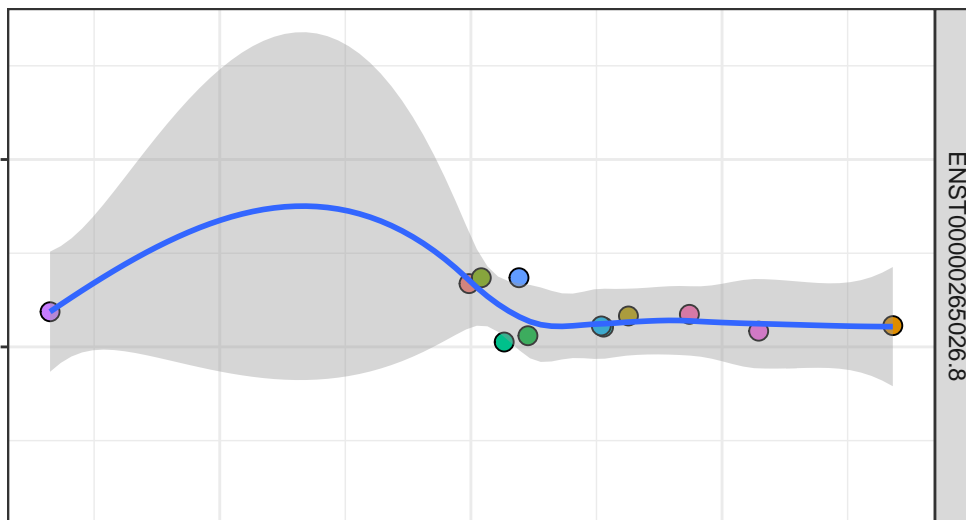
ENST00000265026.8 (down) = -0.209095897722384



PDX_longitudinal
Gene = MAP3K13
Target = HALLMARK_PI3K_AKT_MTOR_SIGNALING
ENST00000448876.5 (up) = -0.6934662984232
ENST00000265026.8 (down) = 0.694949321862588



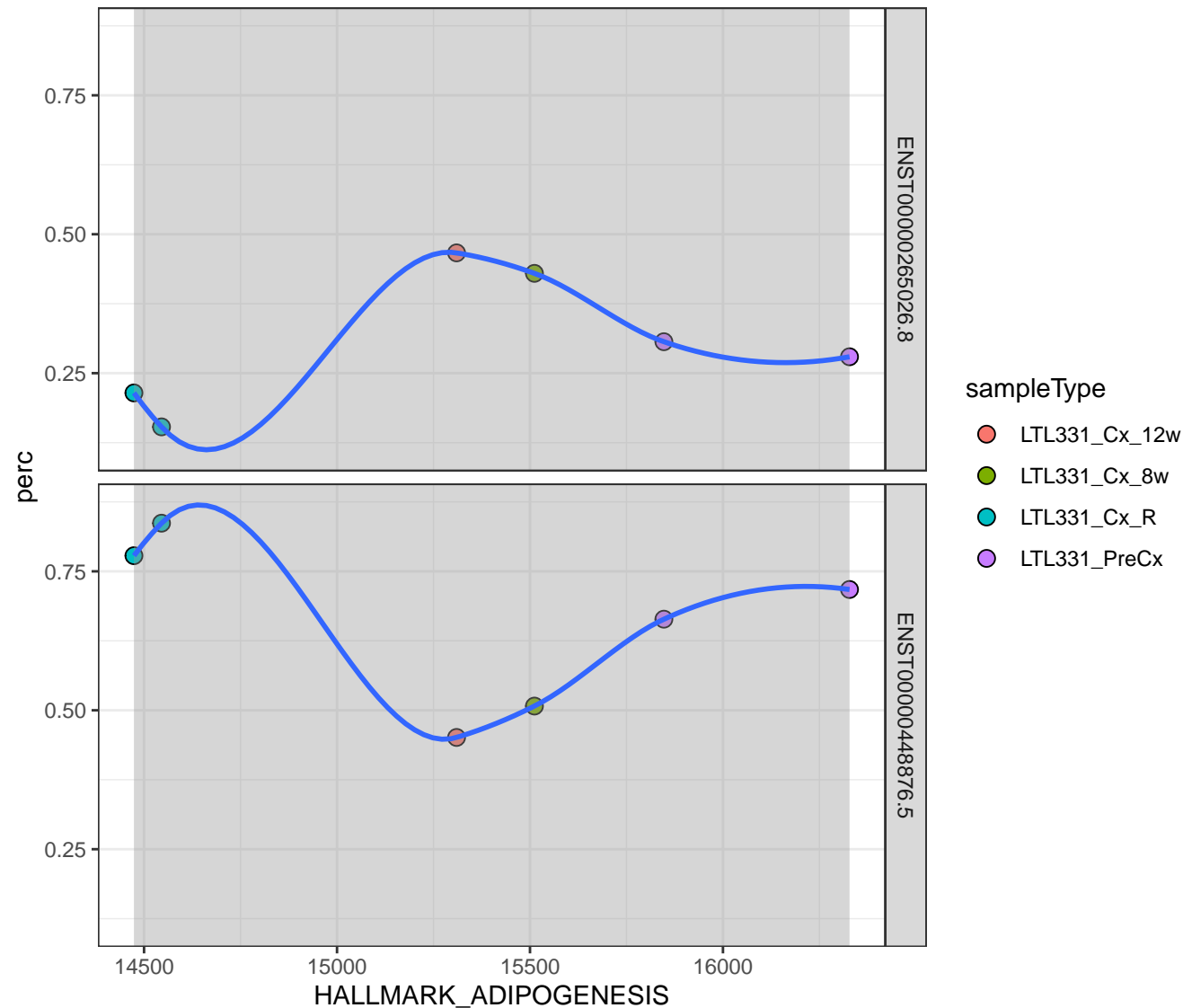
CellLine
Gene = MAP3K13
Target = HALLMARK_ADIPOGENESIS
ENST00000448876.5 (up) = 0.280417647204692
ENST00000265026.8 (down) = -0.28832942437635



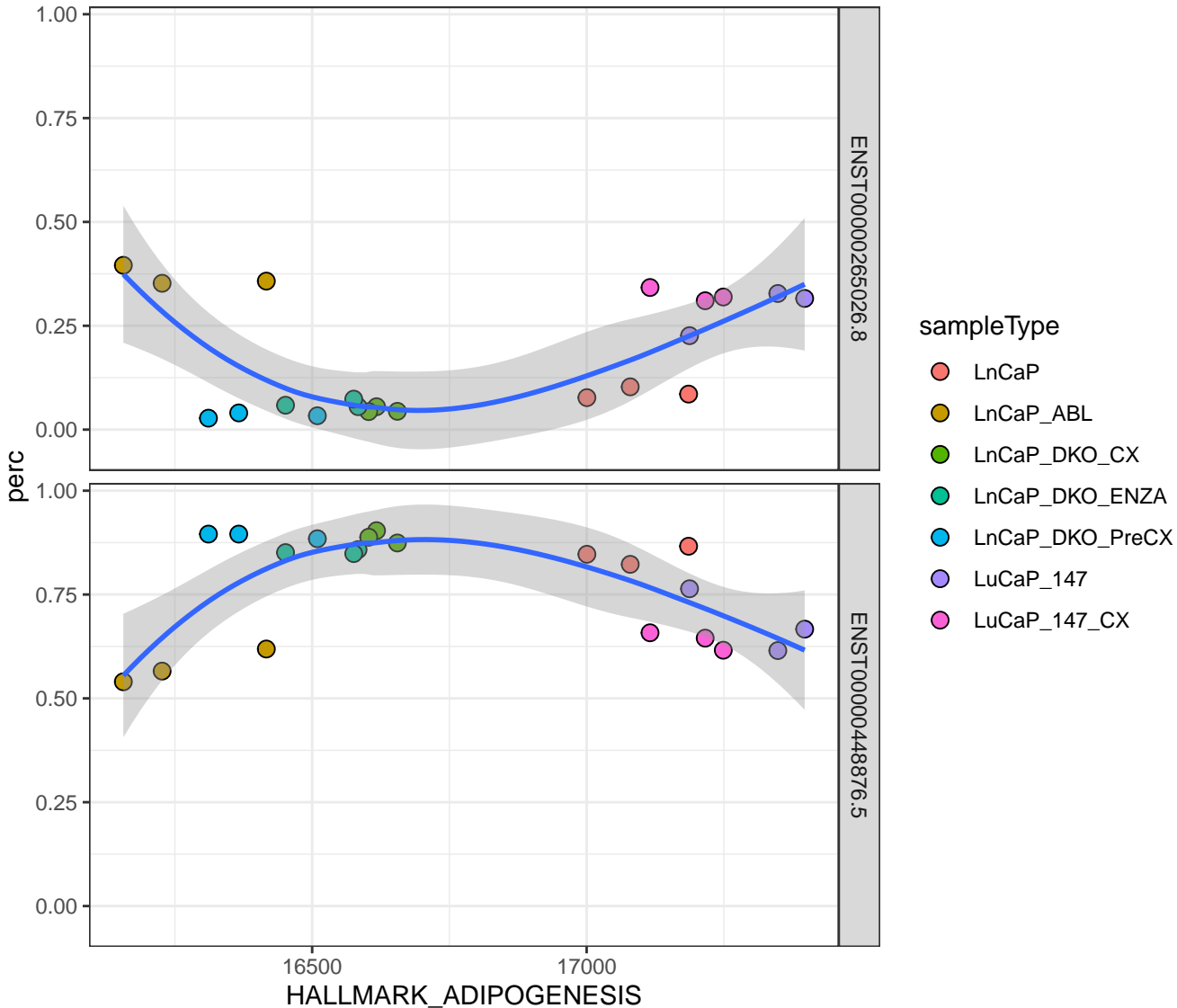
sampleType

- 22RV1
- DU145
- LnCaP
- LnCaP_ABL
- LnCaP_DB
- LnCaP_DKO
- LnCaP_Nelson
- LnCaP_P53
- MDAPCA2B
- NCIH660
- PC3
- VCAP

LTL_longitudinal
Gene = MAP3K13
Target = HALLMARK_ADIPOGENESIS
ENST00000448876.5 (up) = -0.369288305630608
ENST00000265026.8 (down) = 0.433830842877641



PDX_longitudinal
Gene = MAP3K13
Target = HALLMARK_ADIPOGENESIS
ENST00000448876.5 (up) = -0.186378392219709
ENST00000265026.8 (down) = 0.264822230082927



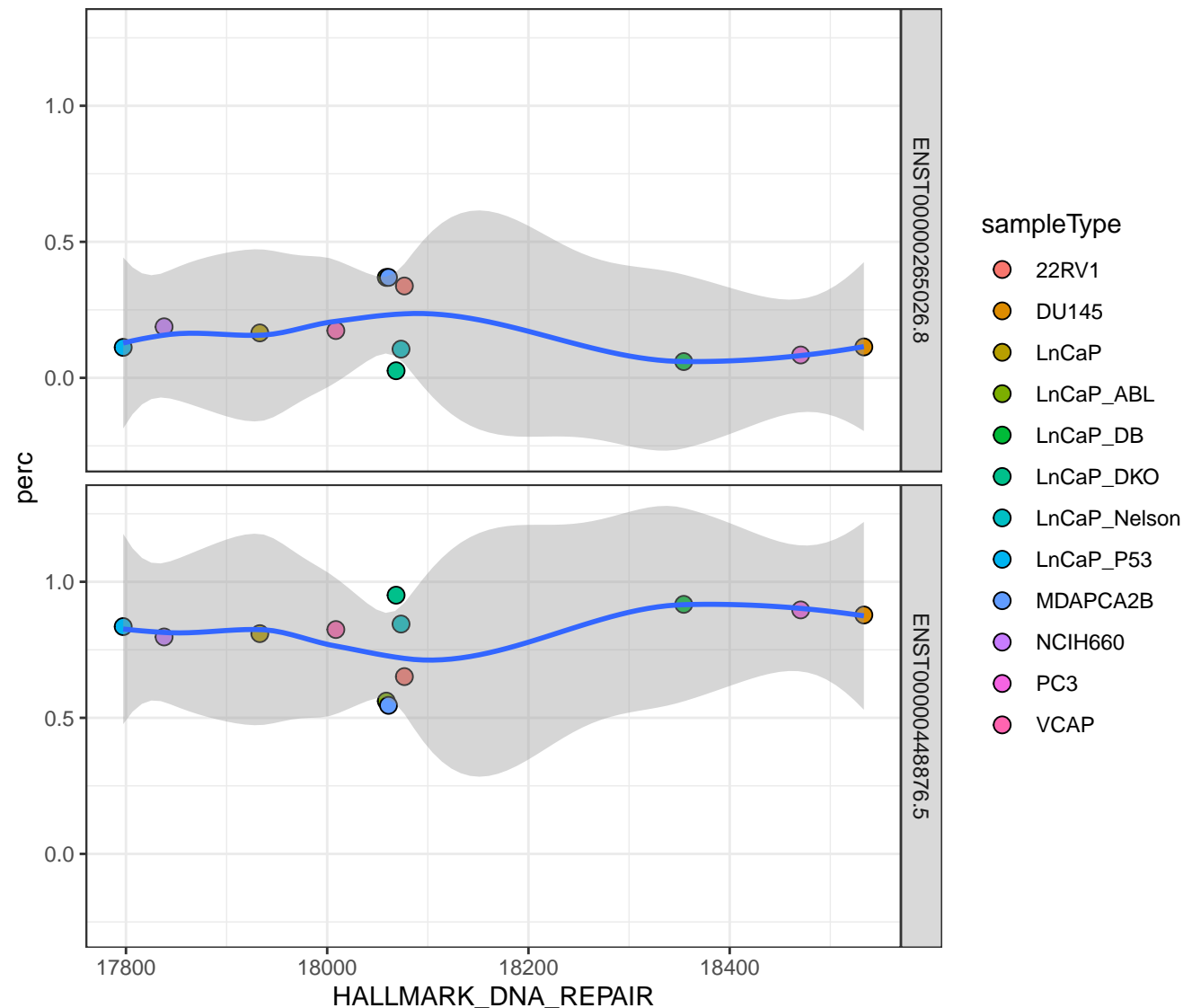
CellLine

Gene = MAP3K13

Target = HALLMARK_DNA_REPAIR

ENST00000448876.5 (up) = 0.301661963528069

ENST00000265026.8 (down) = -0.277657446445252



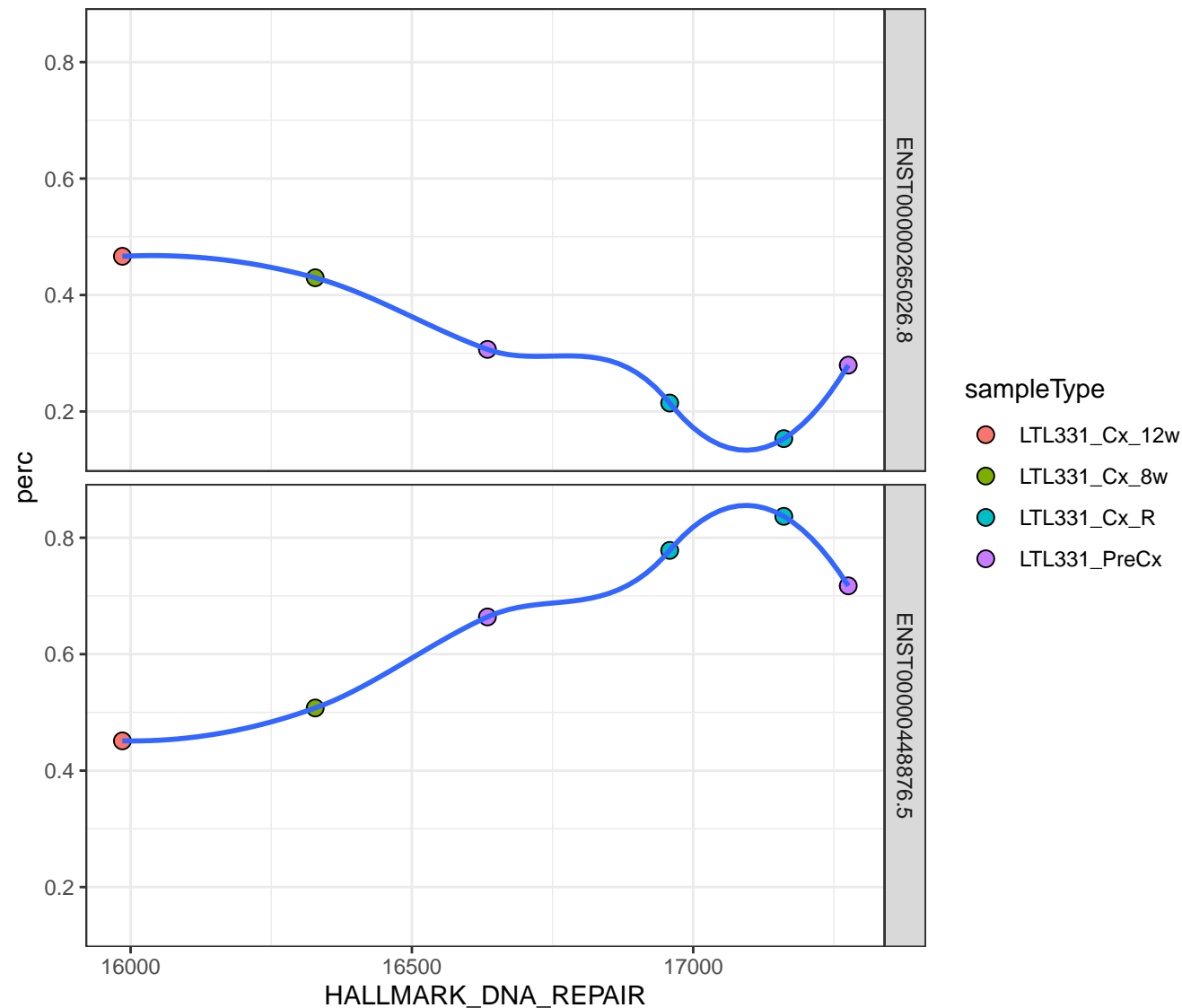
LTL_longitudinal

Gene = MAP3K13

Target = HALLMARK_DNA_REPAIR

ENST00000448876.5 (up) = 0.919989033281344

ENST00000265026.8 (down) = -0.888681875646317



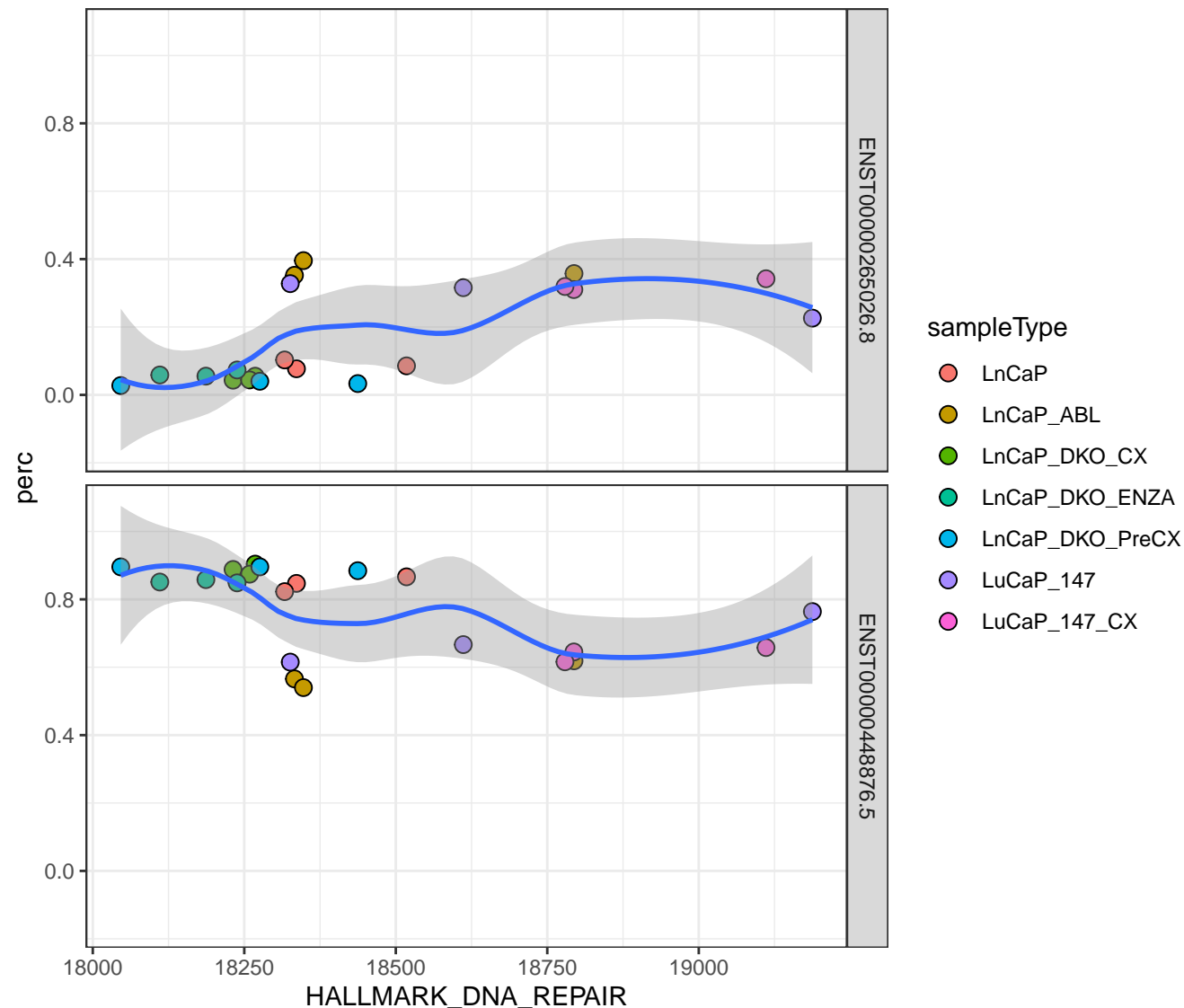
PDX_longitudinal

Gene = MAP3K13

Target = HALLMARK_DNA_REPAIR

ENST00000448876.5 (up) = -0.490169693867597

ENST00000265026.8 (down) = 0.604153635992937



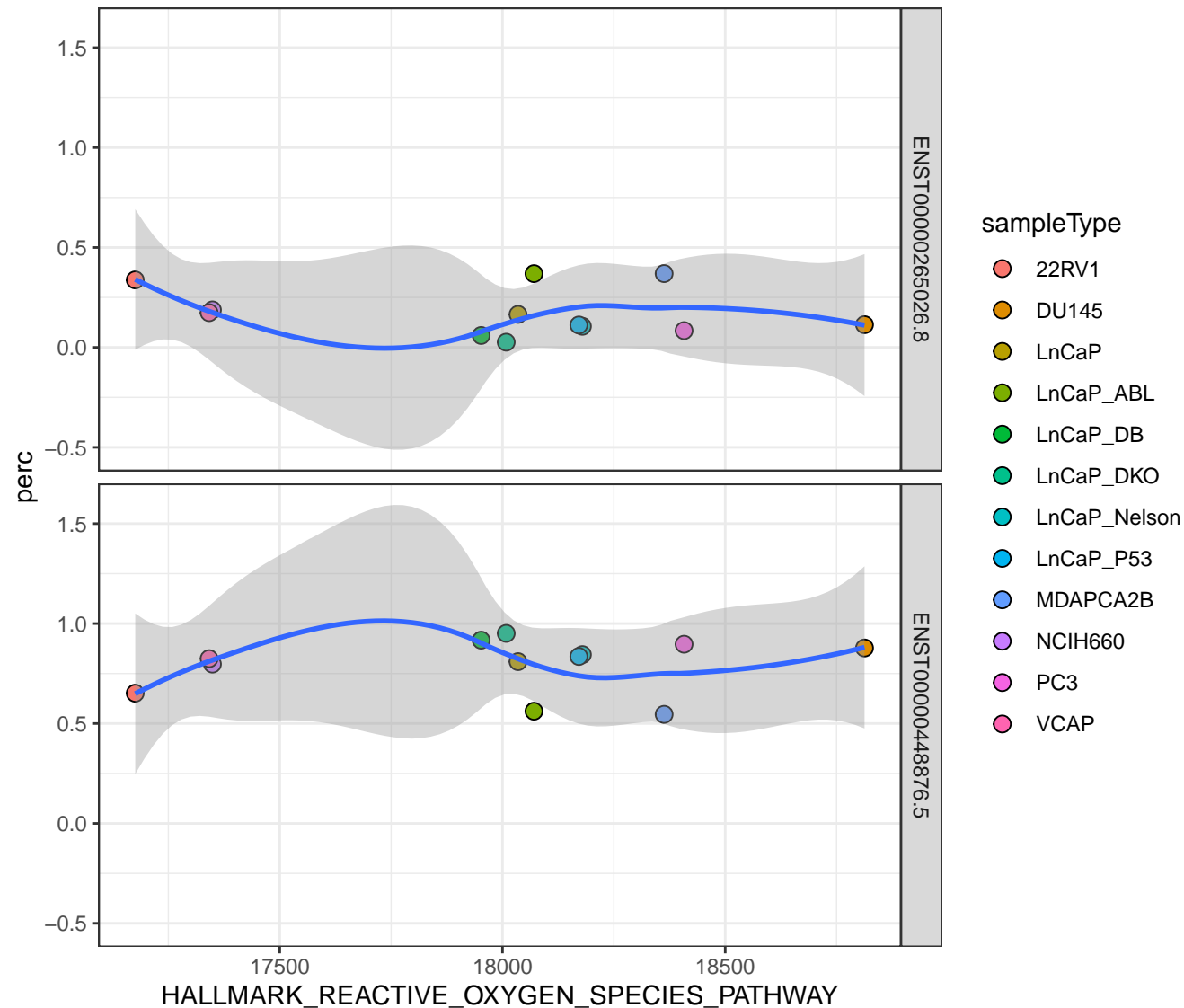
CellLine

Gene = MAP3K13

Target = HALLMARK_REACTIVE_OXYGEN_SPECIES_PATHWAY

ENST00000448876.5 (up) = 0.155322401111631

ENST00000265026.8 (down) = -0.25685366333952



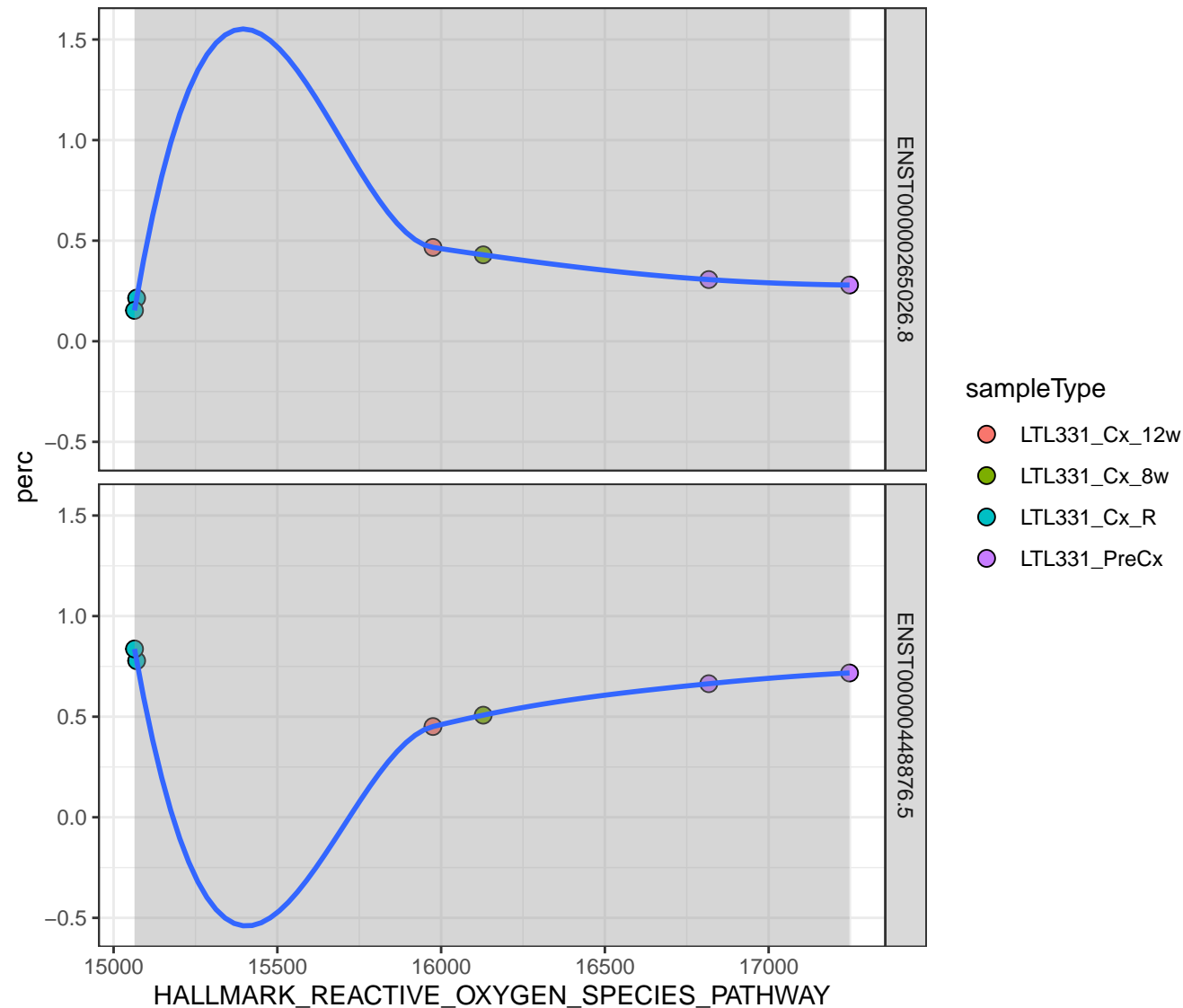
LTL_longitudinal

Gene = MAP3K13

Target = HALLMARK_REACTIVE_OXYGEN_SPECIES_PATHWAY

ENST00000448876.5 (up) = -0.317450526756331

ENST00000265026.8 (down) = 0.383084137511882



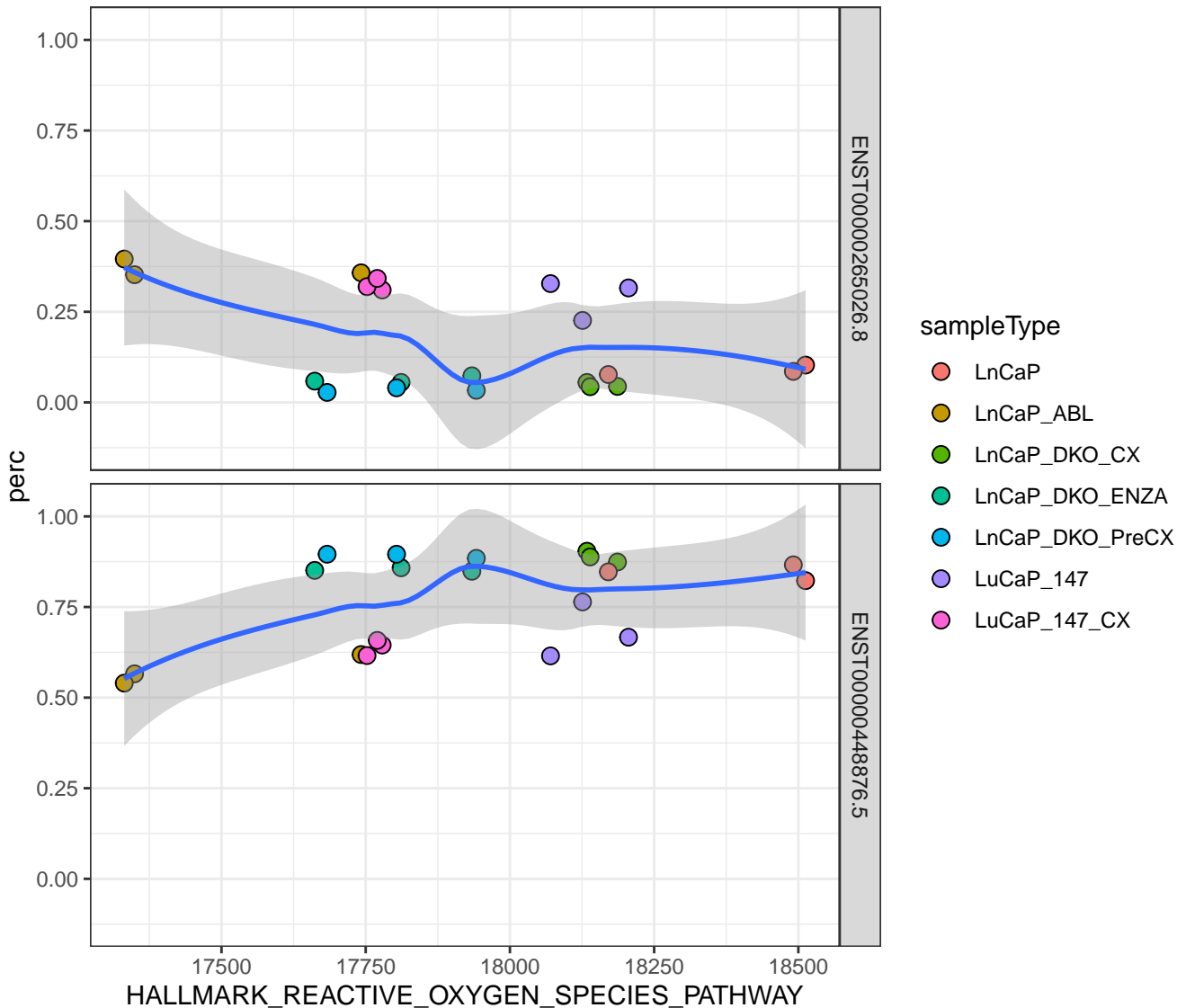
PDX_longitudinal

Gene = MAP3K13

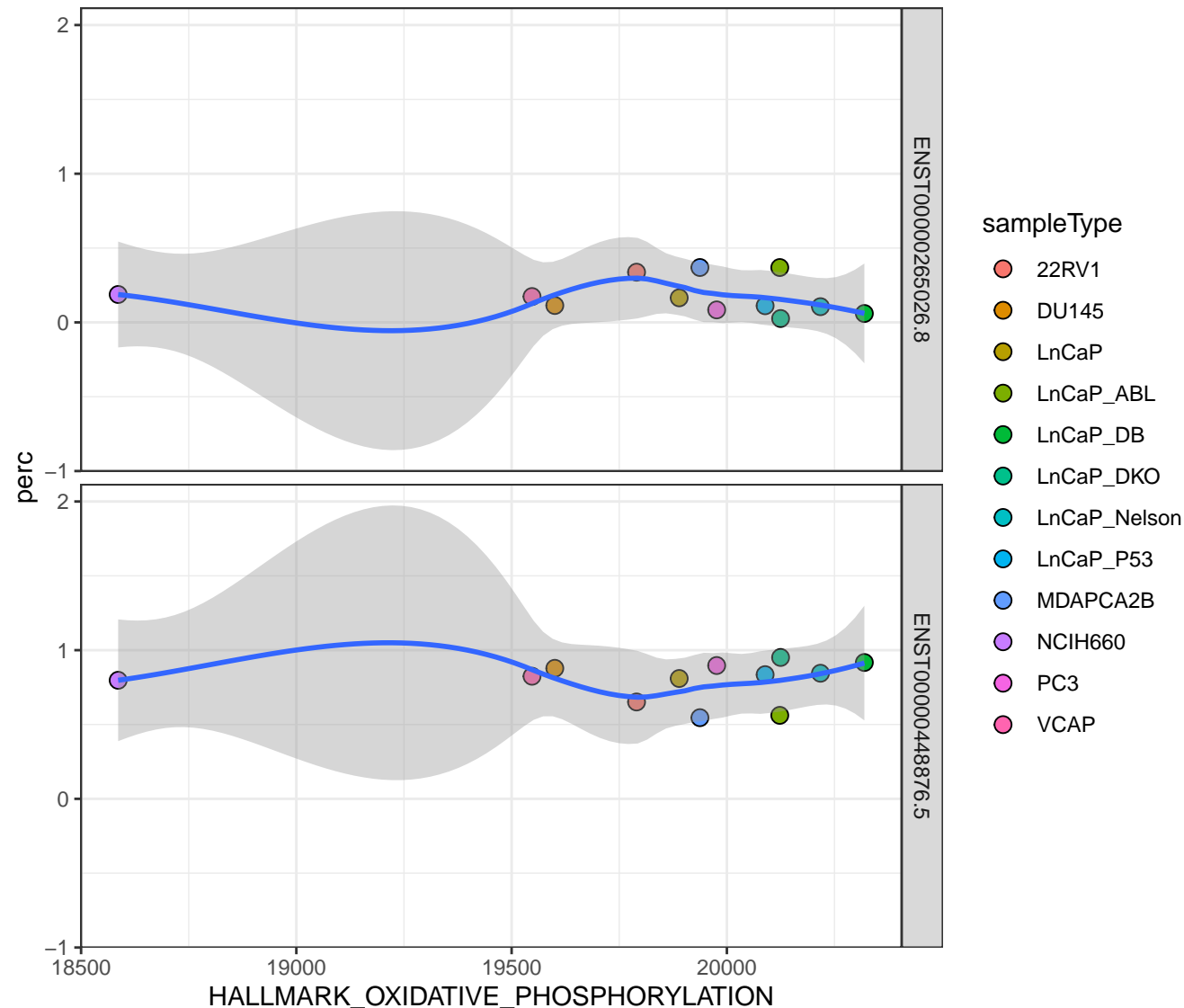
Target = HALLMARK_REACTIVE_OXYGEN_SPECIES_PATHWAY

ENST00000448876.5 (up) = 0.498831703118158

ENST00000265026.8 (down) = -0.425890867229829



CellLine
 Gene = MAP3K13
 Target = HALLMARK_OXIDATIVE_PHOSPHORYLATION
 ENST00000448876.5 (up) = 0.0468942864441836
 ENST00000265026.8 (down) = -0.143808013521439



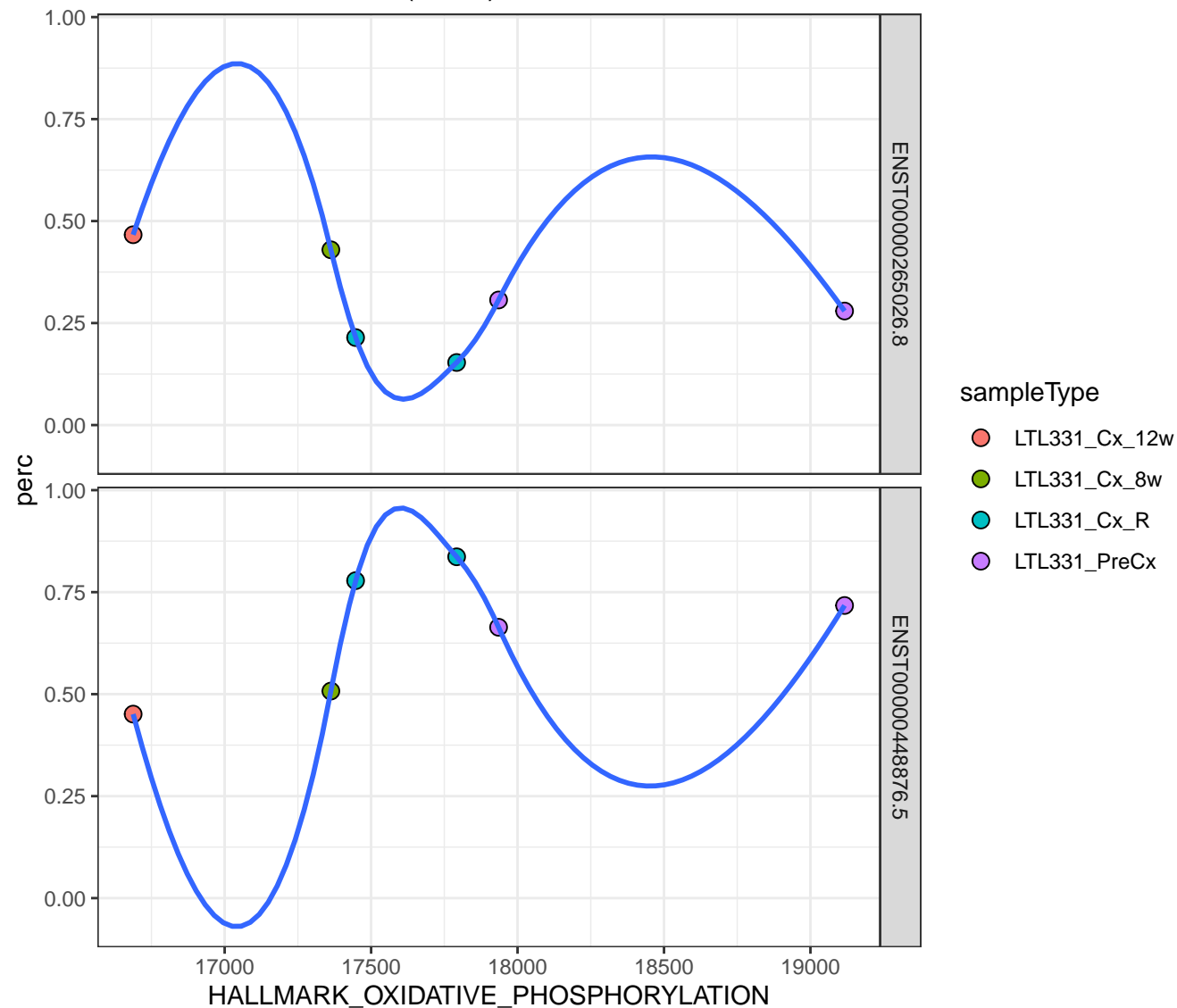
LTL_longitudinal

Gene = MAP3K13

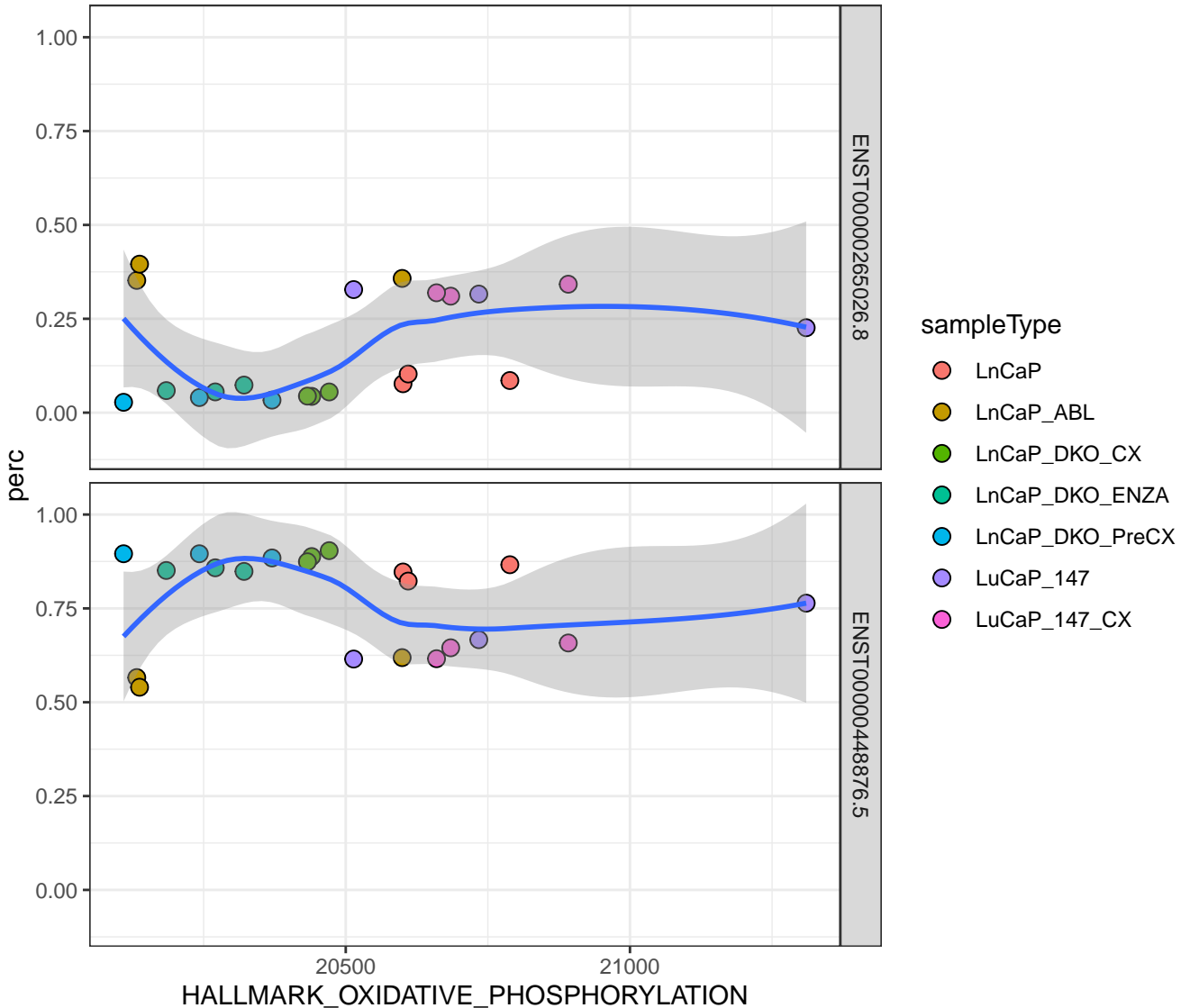
Target = HALLMARK_OXIDATIVE_PHOSPHORYLATION

ENST00000448876.5 (up) = 0.539842308260934

ENST00000265026.8 (down) = -0.475209698591984



PDX_longitudinal
 Gene = MAP3K13
 Target = HALLMARK_OXIDATIVE_PHOSPHORYLATION
 ENST00000448876.5 (up) = -0.14804958841048
 ENST00000265026.8 (down) = 0.280585342284792



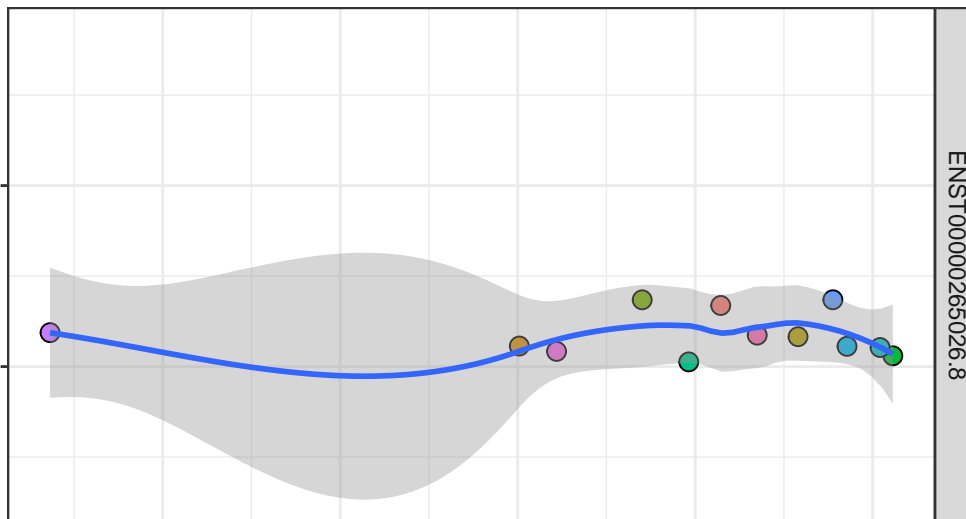
CellLine

Gene = MAP3K13

Target = HALLMARK_FATTY_ACID_METABOLISM

ENST00000448876.5 (up) = -0.0416634320516786

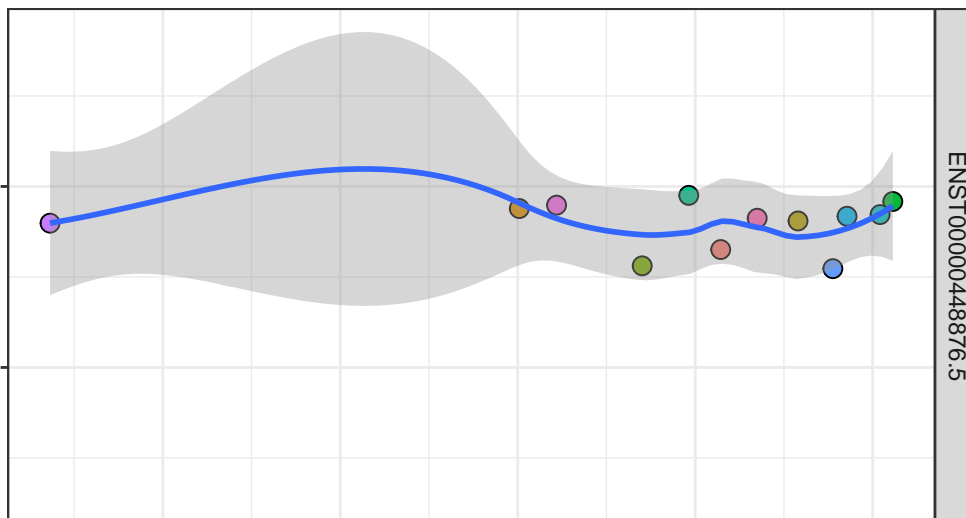
ENST00000265026.8 (down) = -0.0305872891418502



ENST00000265026.8

sampleType

- 22RV1
- DU145
- LnCaP
- LnCaP_ABL
- LnCaP_DB
- LnCaP_DKO
- LnCaP_Nelson
- LnCaP_P53
- MDAPCA2B
- NCIH660
- PC3
- VCAP



ENST00000448876.5

HALLMARK_FATTY_ACID_METABOLISM

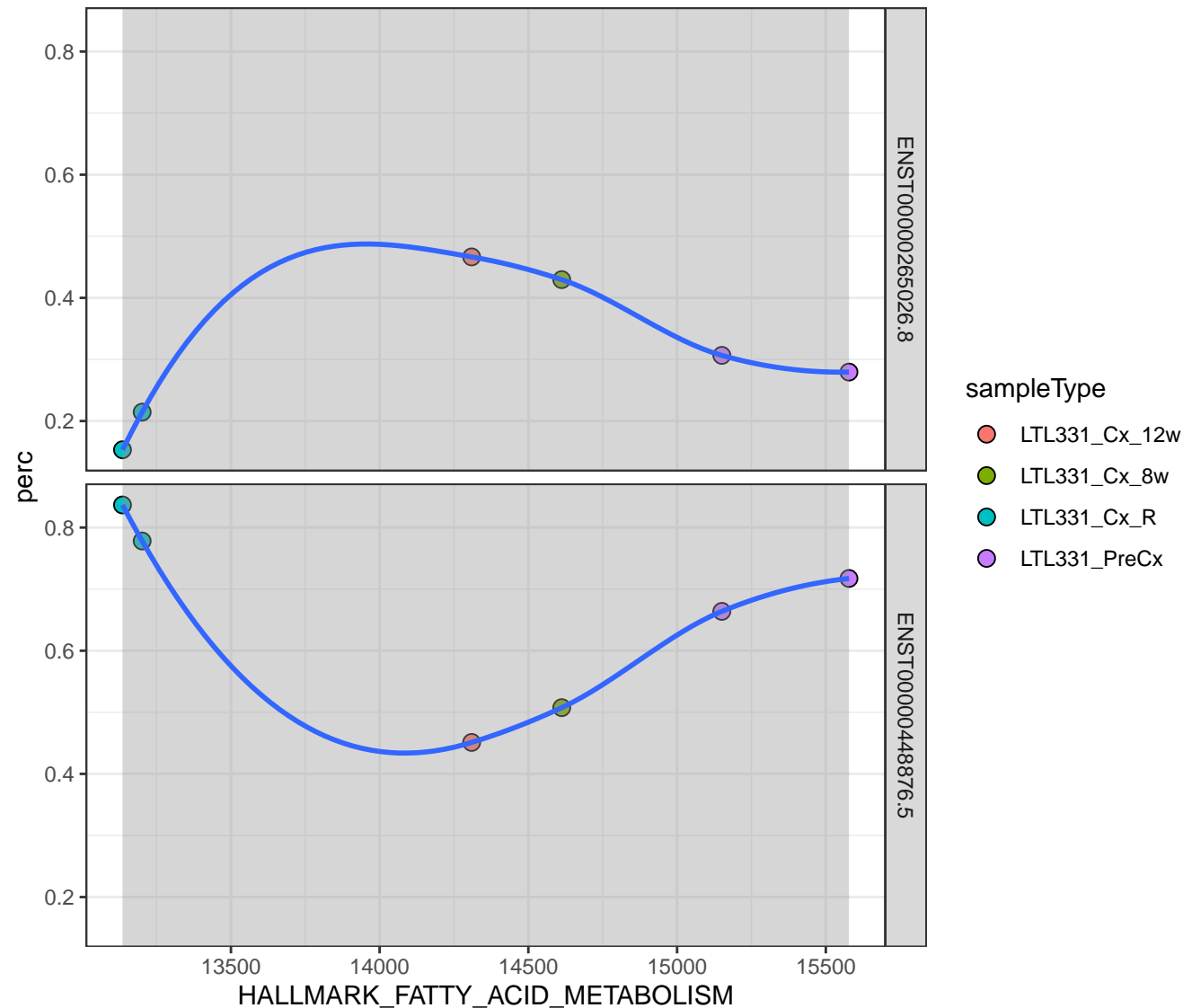
LTL_longitudinal

Gene = MAP3K13

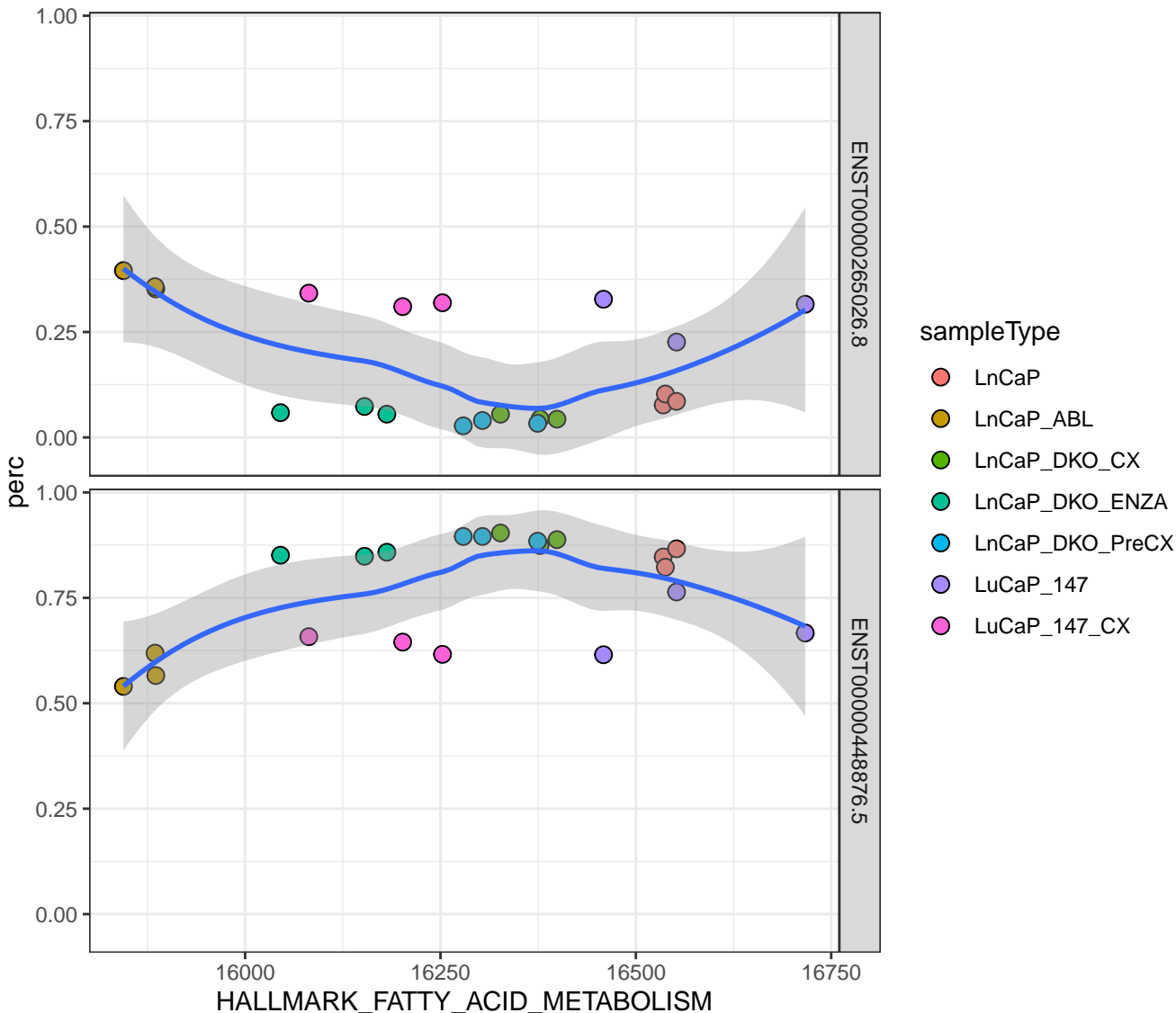
Target = HALLMARK_FATTY_ACID_METABOLISM

ENST00000448876.5 (up) = -0.405535233789492

ENST00000265026.8 (down) = 0.46935273248466



PDX_longitudinal
 Gene = MAP3K13
 Target = HALLMARK_FATTY_ACID_METABOLISM
 ENST00000448876.5 (up) = 0.432673666783885
 ENST00000265026.8 (down) = -0.367074972952484



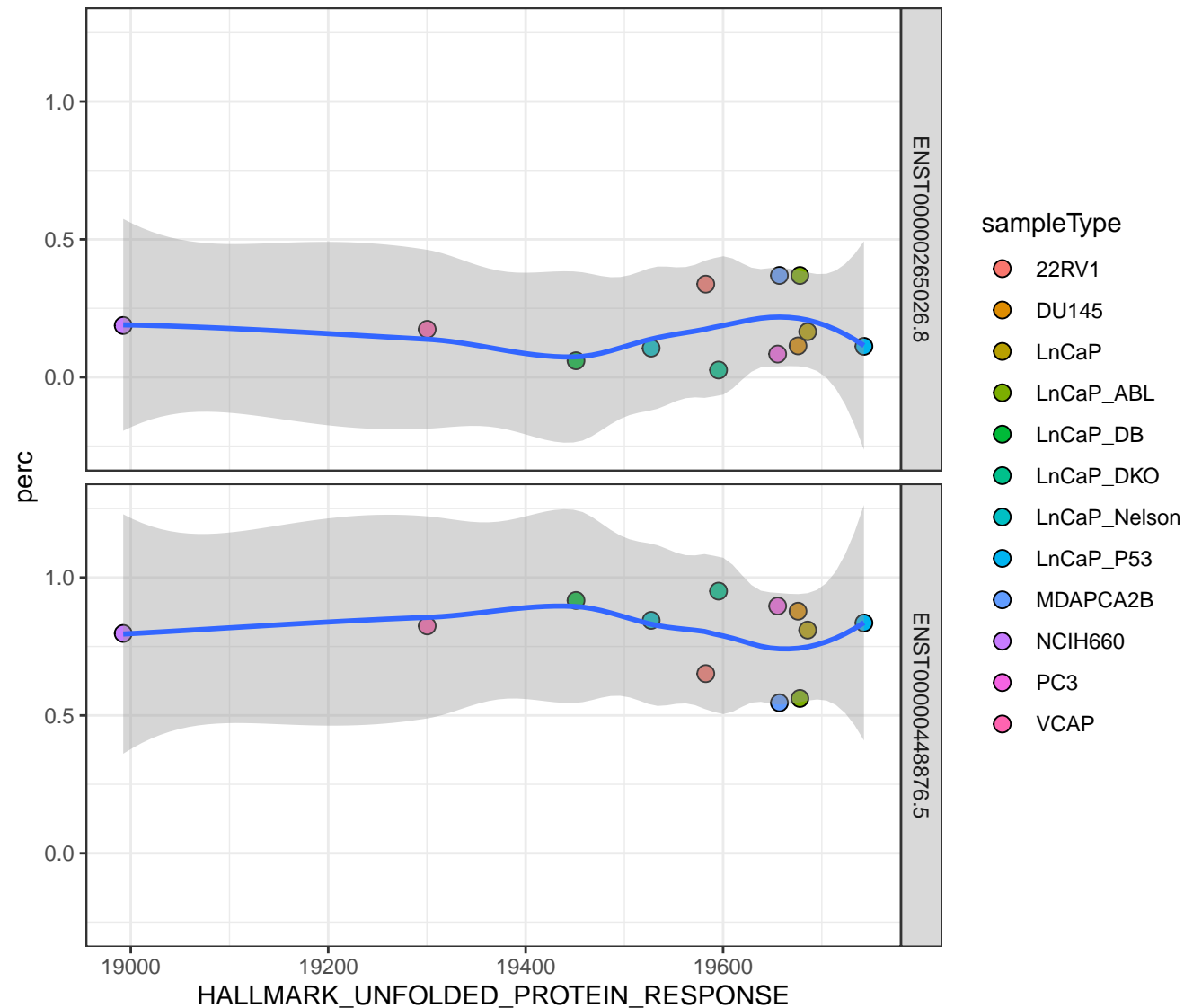
CellLine

Gene = MAP3K13

Target = HALLMARK_UNFOLDED_PROTEIN_RESPONSE

ENST00000448876.5 (up) = -0.143830390658915

ENST00000265026.8 (down) = 0.0697426245010765



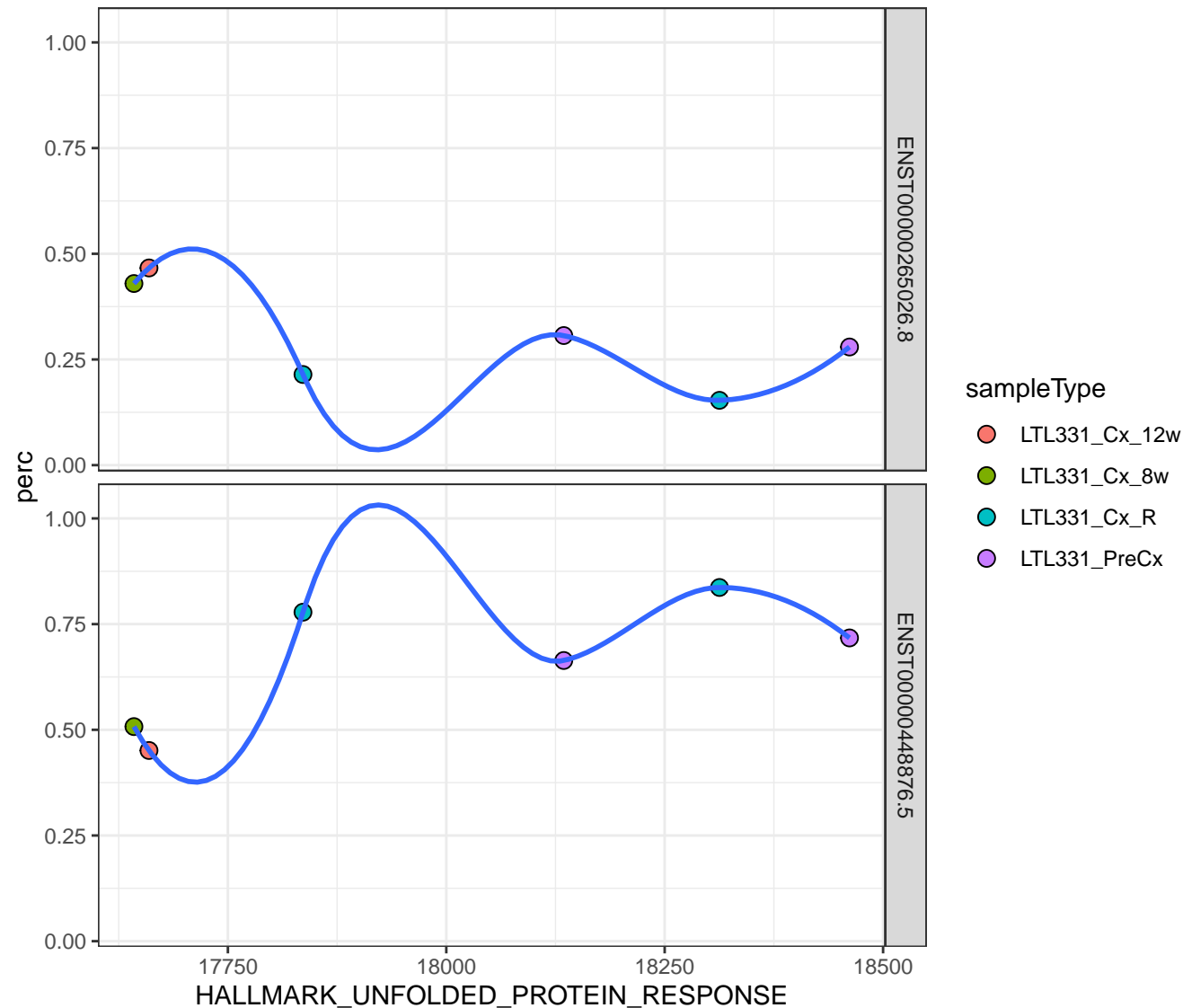
LTL_longitudinal

Gene = MAP3K13

Target = HALLMARK_UNFOLDED_PROTEIN_RESPONSE

ENST00000448876.5 (up) = 0.716975398889894

ENST00000265026.8 (down) = -0.685743574228836



PDX_longitudinal
 Gene = MAP3K13
 Target = HALLMARK_UNFOLDED_PROTEIN_RESPONSE
 ENST00000448876.5 (up) = 0.568762490920985
 ENST00000265026.8 (down) = -0.518198110500274

