

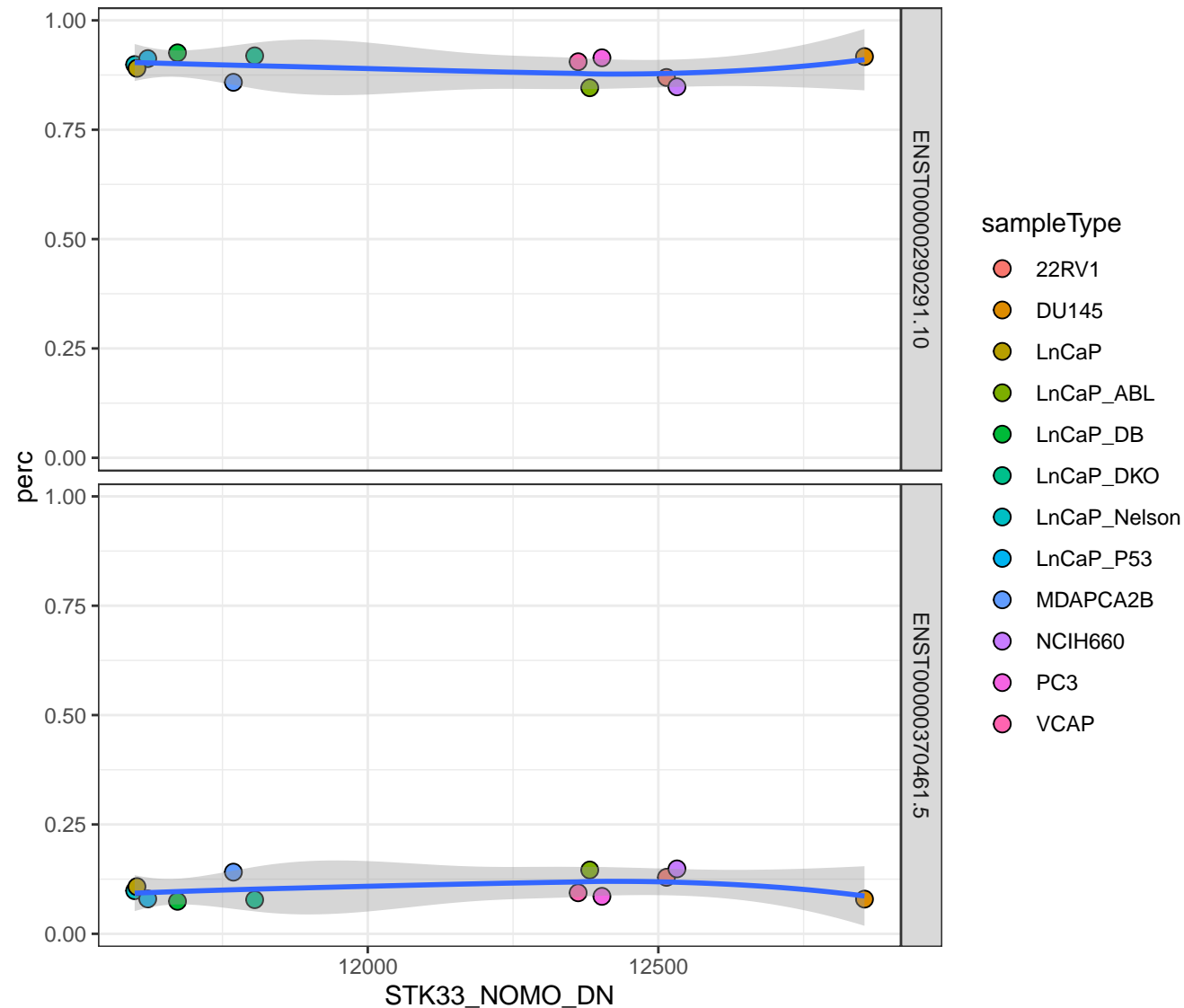
CellLine

Gene = OGFR

Target = STK33_NOMO_DN

ENST00000290291.10 (up) = -0.24480572651673

ENST00000370461.5 (down) = 0.245840418583767



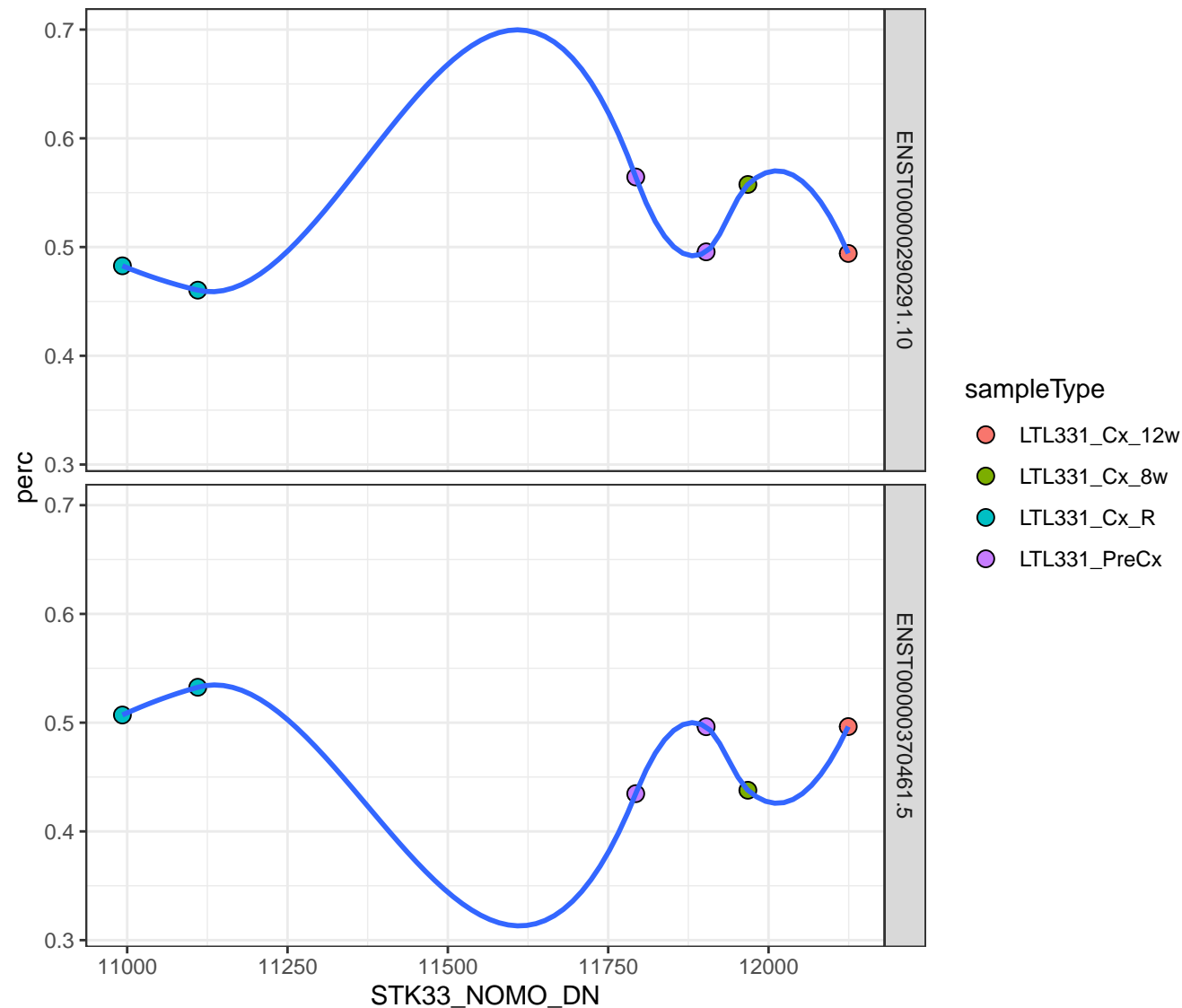
LTL_longitudinal

Gene = OGFR

Target = STK33_NOMO_DN

ENST00000290291.10 (up) = 0.562268872991786

ENST00000370461.5 (down) = -0.577122620521706



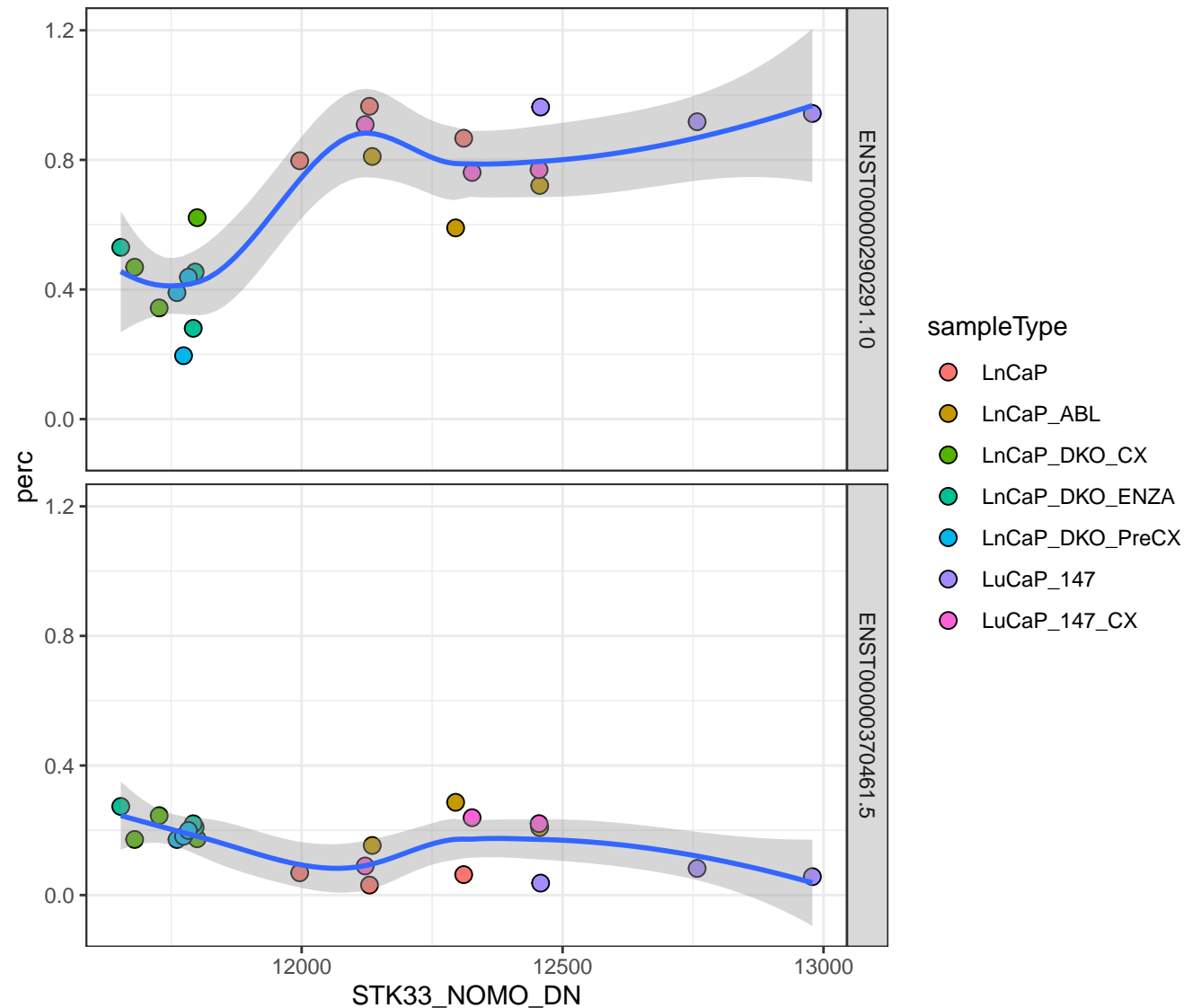
PDX_longitudinal

Gene = OGFR

Target = STK33_NOMO_DN

ENST00000290291.10 (up) = 0.765360584483024

ENST00000370461.5 (down) = -0.446855132677861



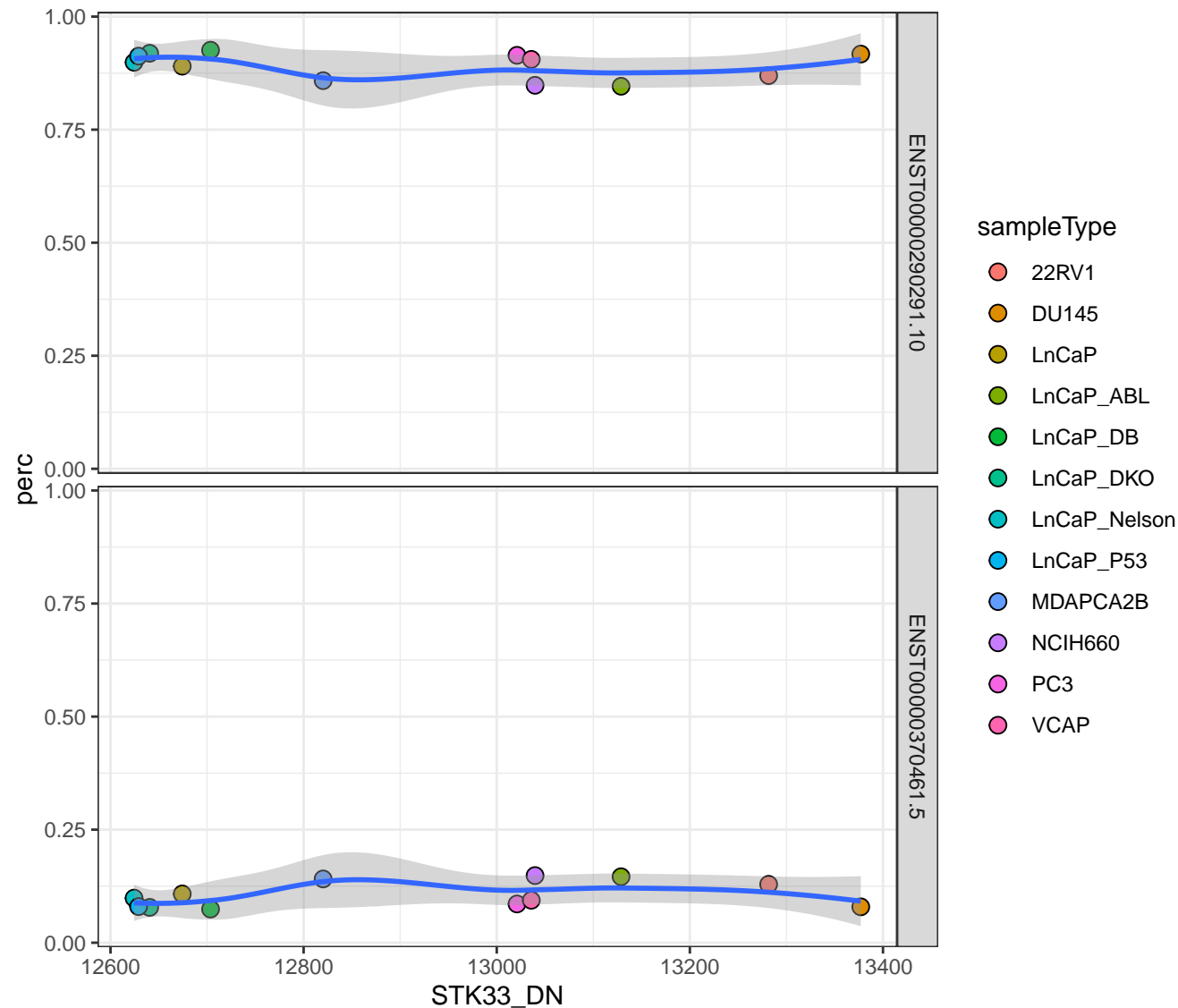
CellLine

Gene = OGFR

Target = STK33_DN

ENST00000290291.10 (up) = -0.312062508044866

ENST00000370461.5 (down) = 0.316242247728238



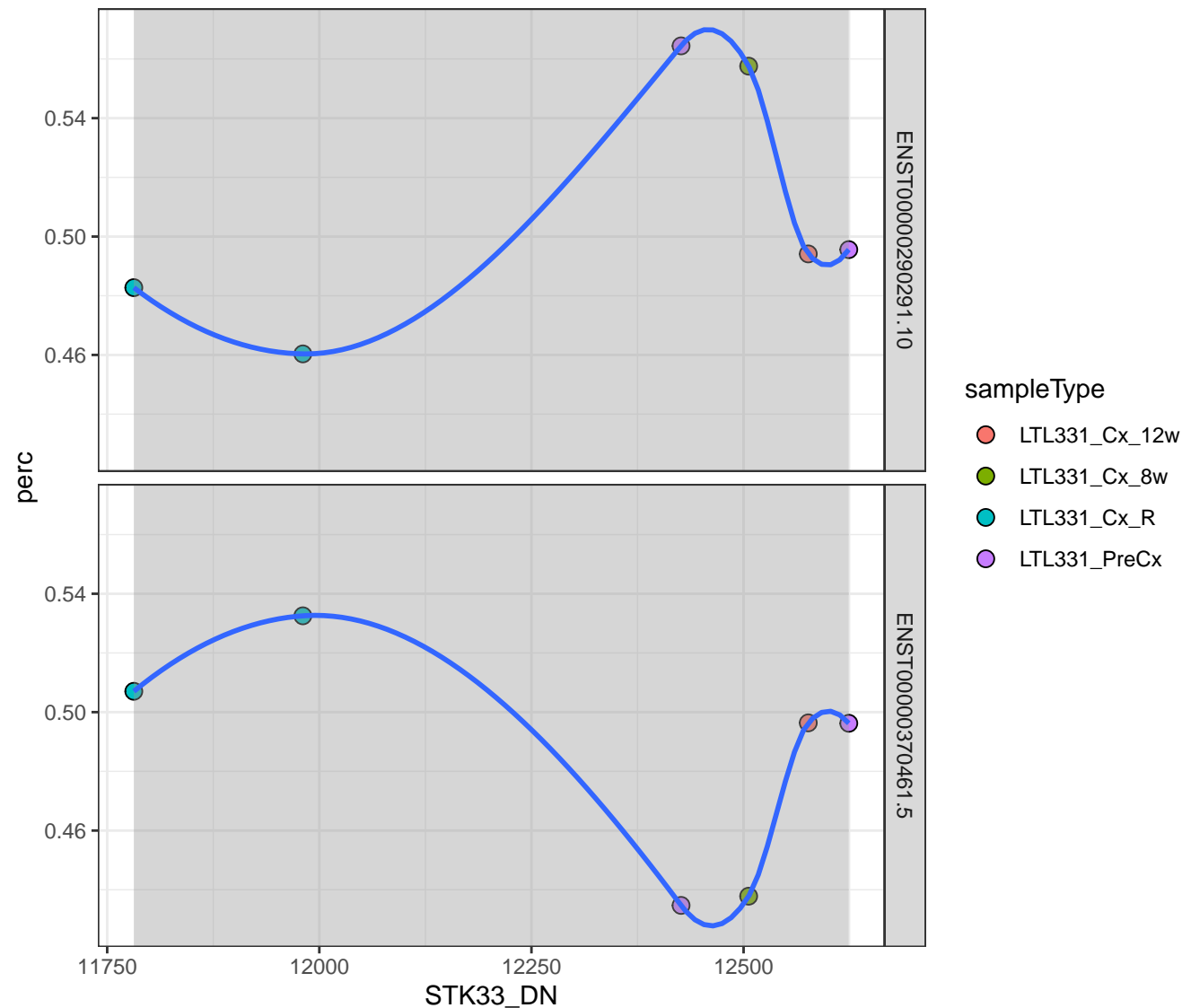
LTL_longitudinal

Gene = OGFR

Target = STK33_DN

ENST00000290291.10 (up) = 0.512361644100001

ENST00000370461.5 (down) = -0.521240934955602



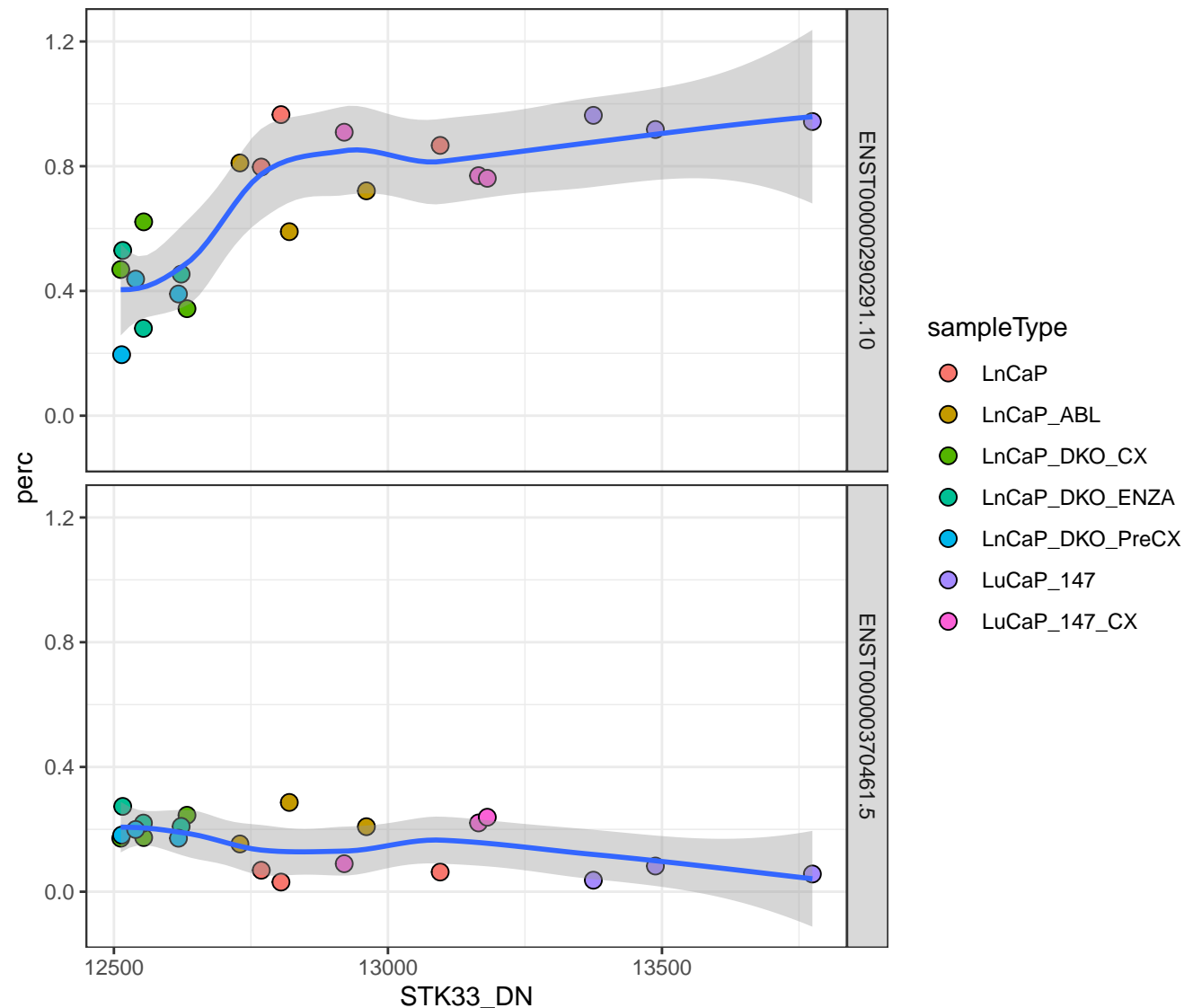
PDX_longitudinal

Gene = OGFR

Target = STK33_DN

ENST00000290291.10 (up) = 0.752570673300076

ENST00000370461.5 (down) = -0.507865245143817



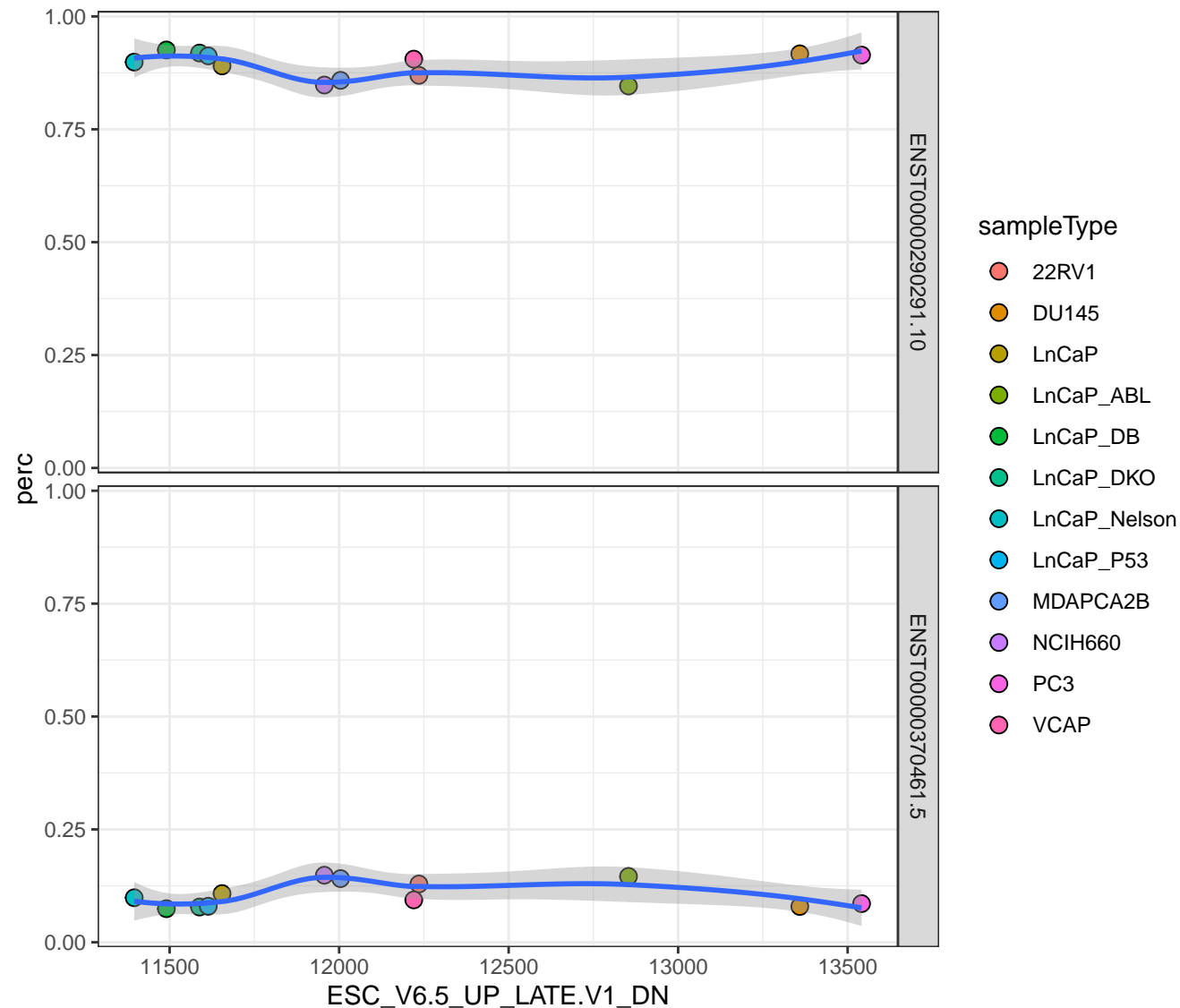
CellLine

Gene = OGFR

Target = ESC_V6.5_UP_LATE.V1_DN

ENST00000290291.10 (up) = -0.0480288487191668

ENST00000370461.5 (down) = 0.0467917028200017



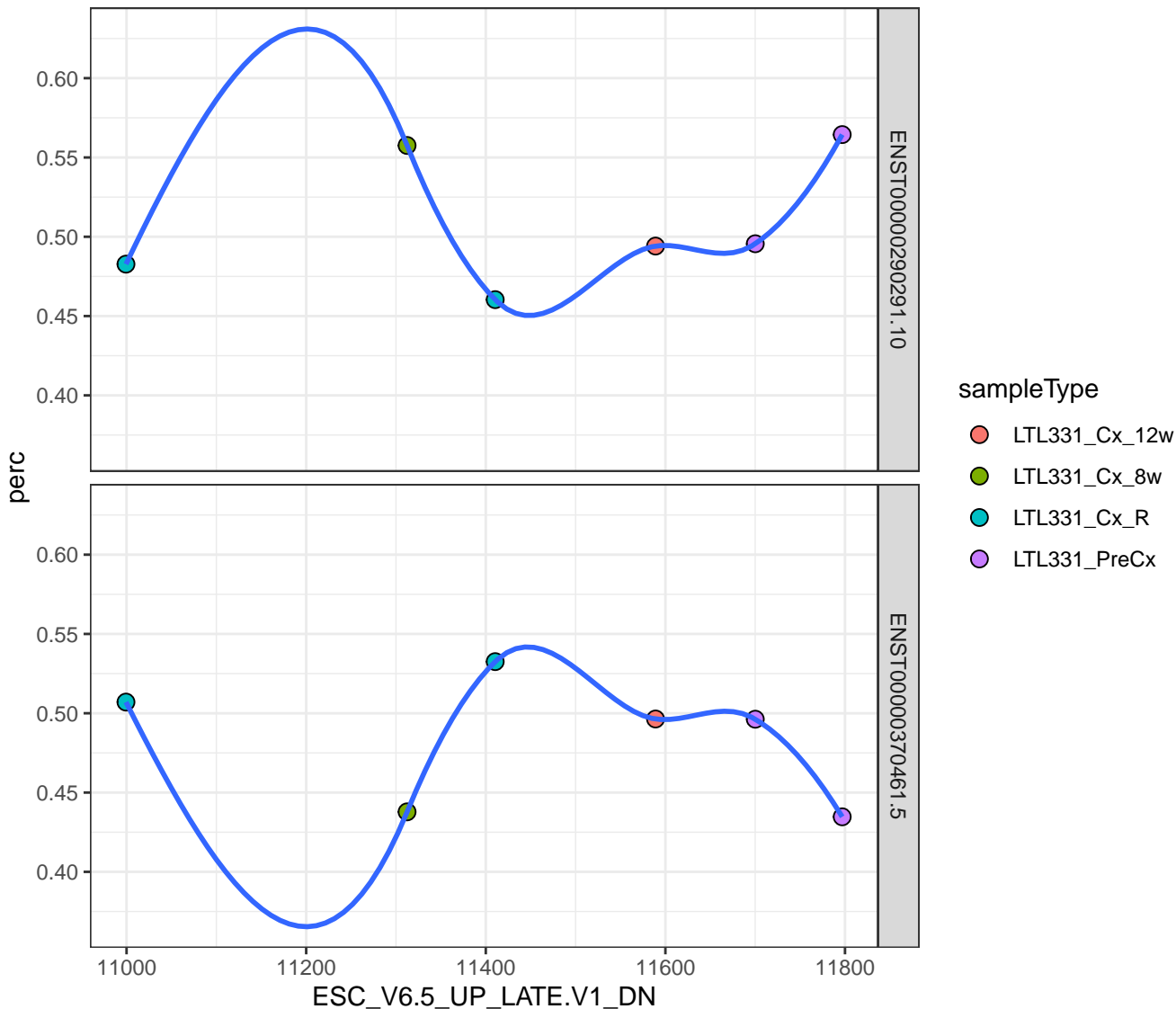
LTL_longitudinal

Gene = OGFR

Target = ESC_V6.5_UP_LATE.V1_DN

ENST00000290291.10 (up) = 0.339973521460579

ENST00000370461.5 (down) = -0.318850556205994



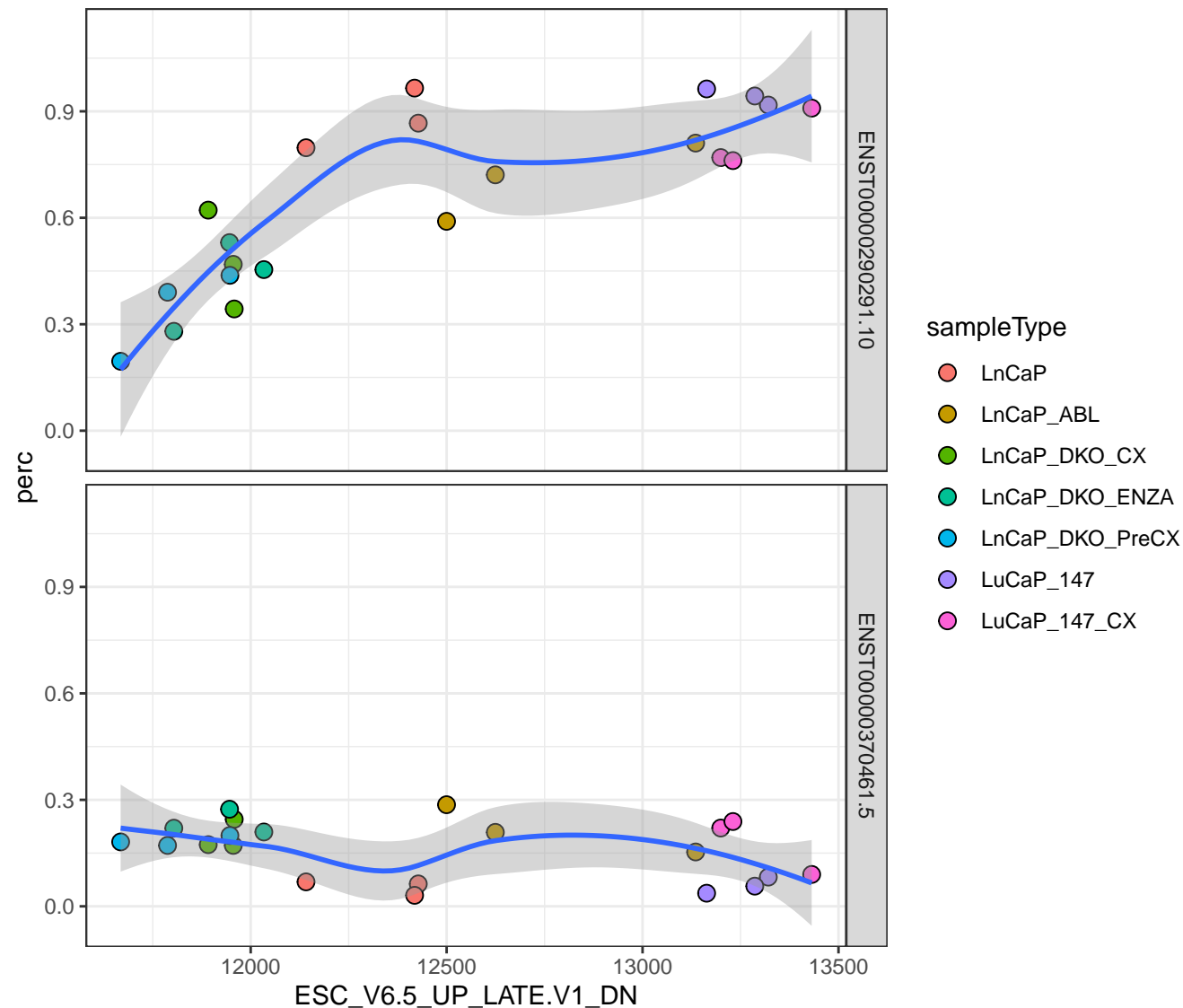
PDX_longitudinal

Gene = OGFR

Target = ESC_V6.5_UP_LATE.V1_DN

ENST00000290291.10 (up) = 0.817038546038309

ENST00000370461.5 (down) = -0.380329332797699



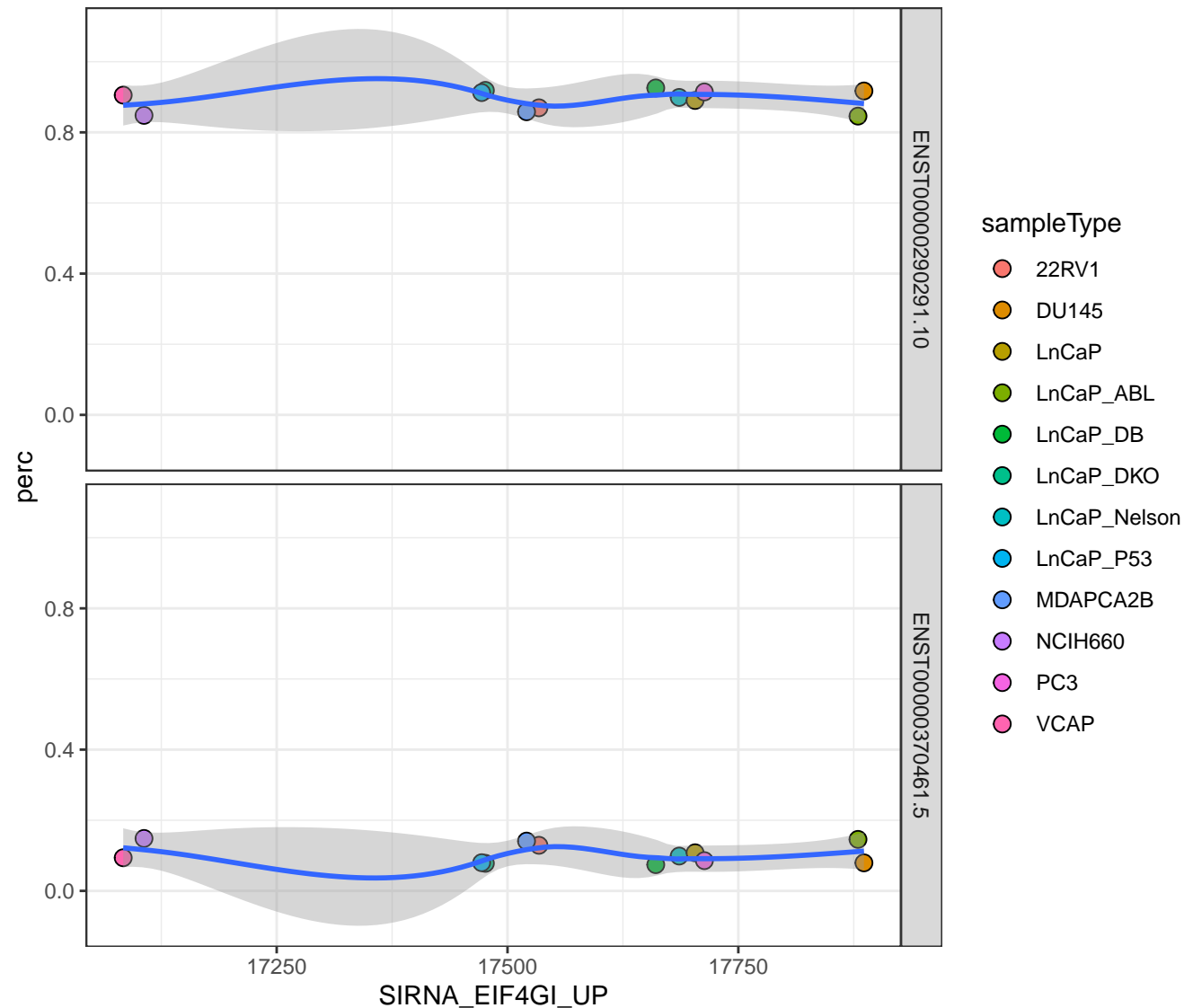
CellLine

Gene = OGFR

Target = SIRNA_EIF4GI_UP

ENST00000290291.10 (up) = 0.149575234829331

ENST00000370461.5 (down) = -0.169882825423634



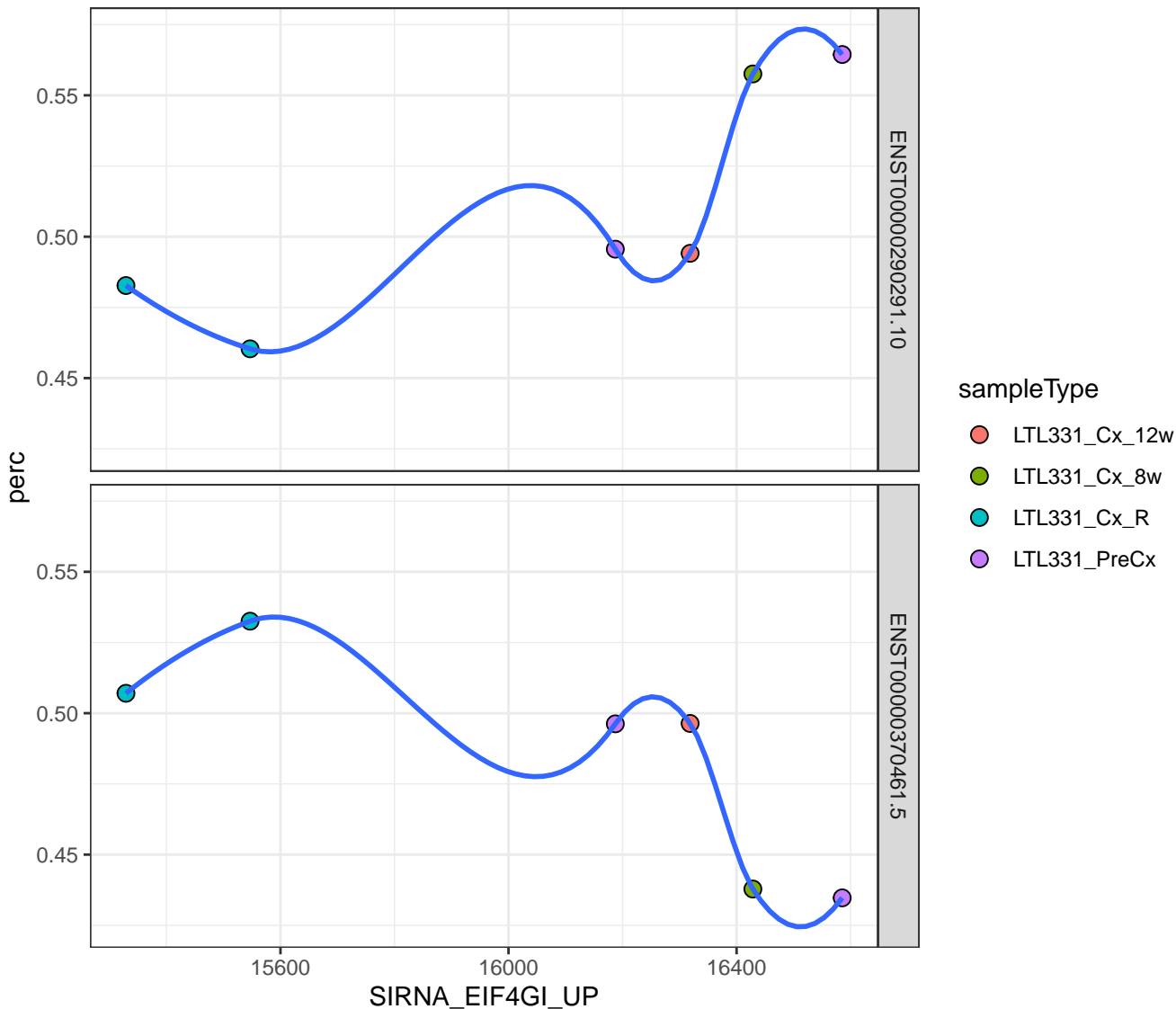
LTL_longitudinal

Gene = OGFR

Target = SIRNA_EIF4GI_UP

ENST00000290291.10 (up) = 0.798945240231133

ENST00000370461.5 (down) = -0.798413566195734



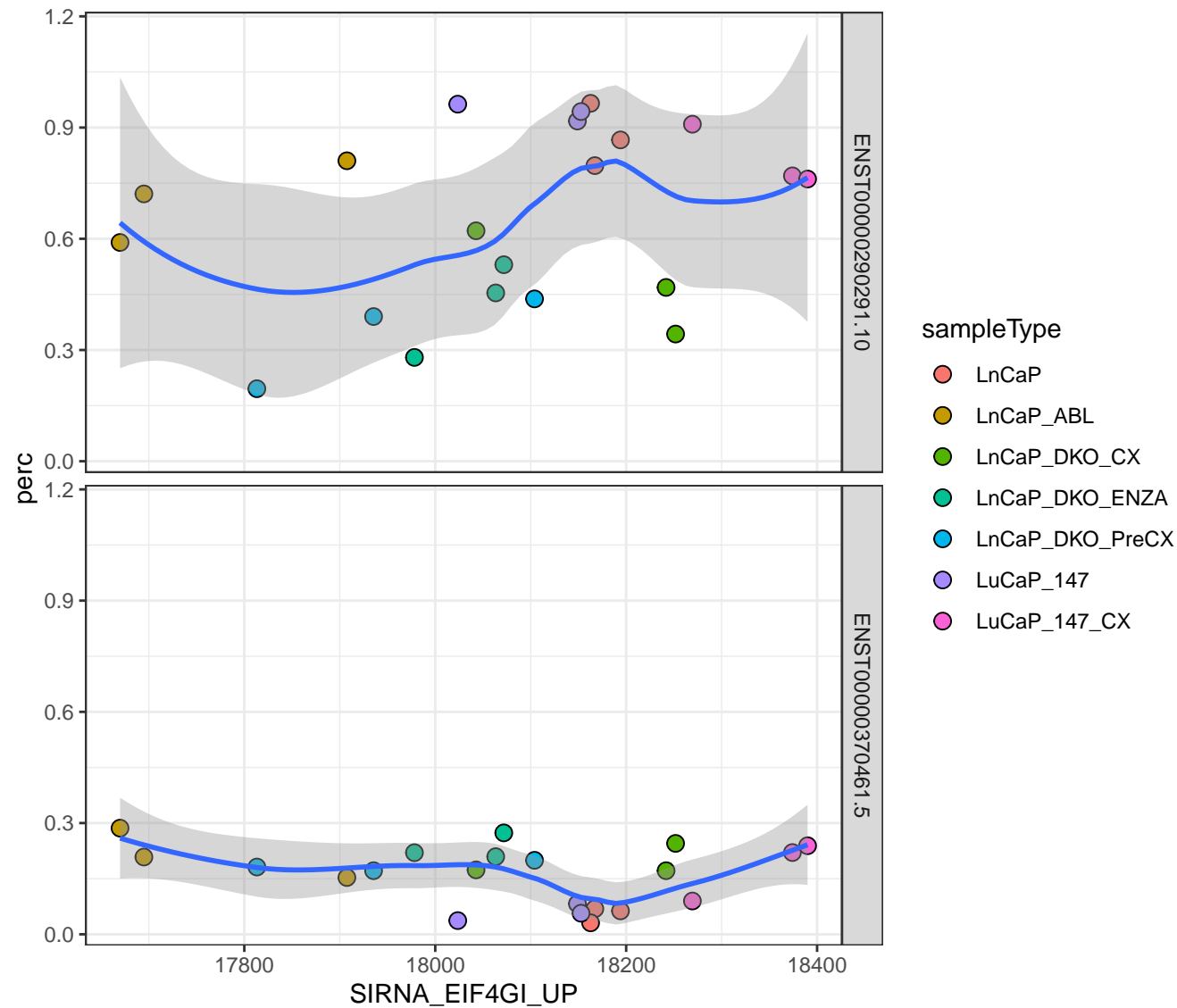
PDX_longitudinal

Gene = OGFR

Target = SIRNA_EIF4GI_UP

ENST00000290291.10 (up) = 0.3027213553631

ENST00000370461.5 (down) = -0.237039991719546



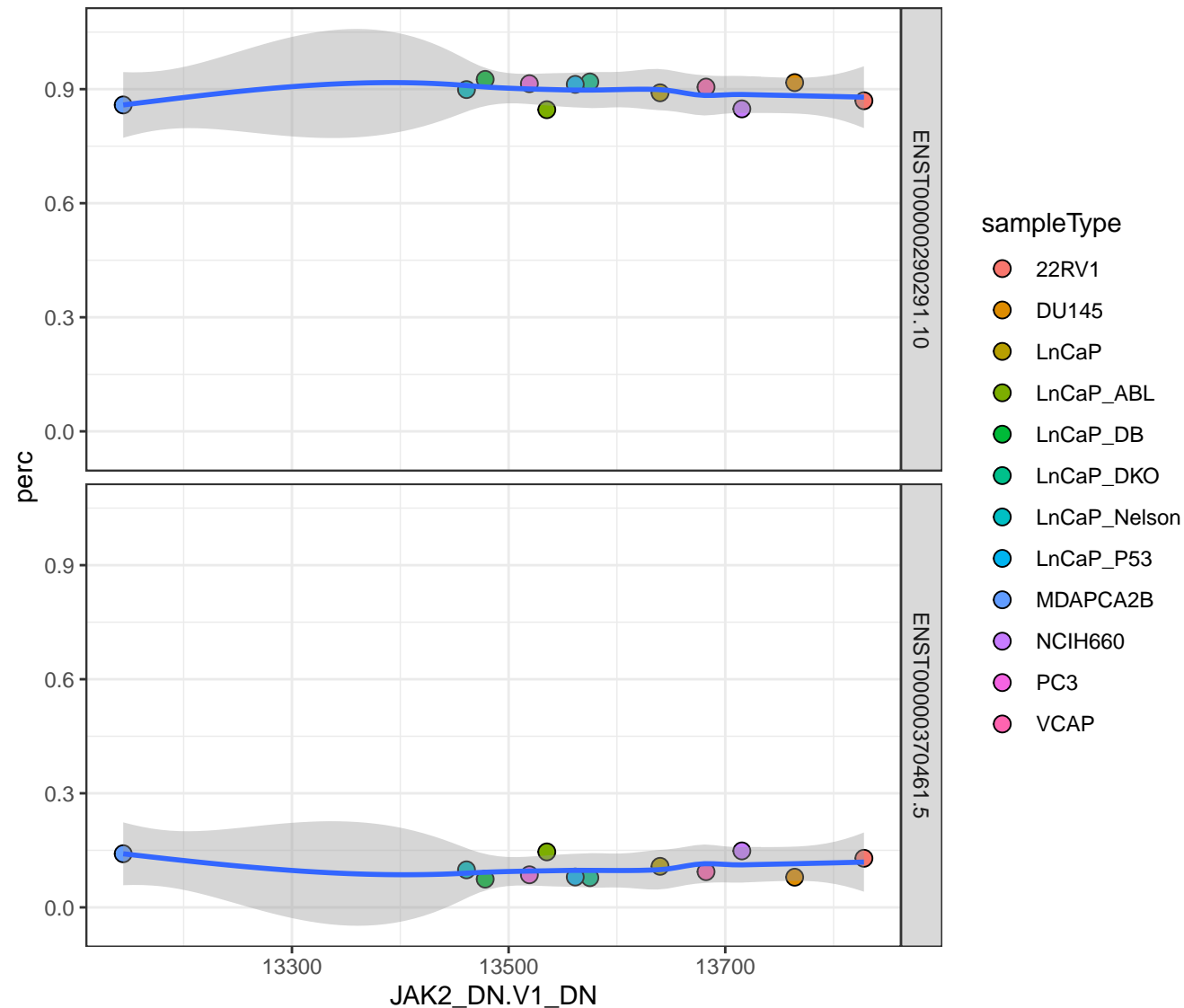
CellLine

Gene = OGFR

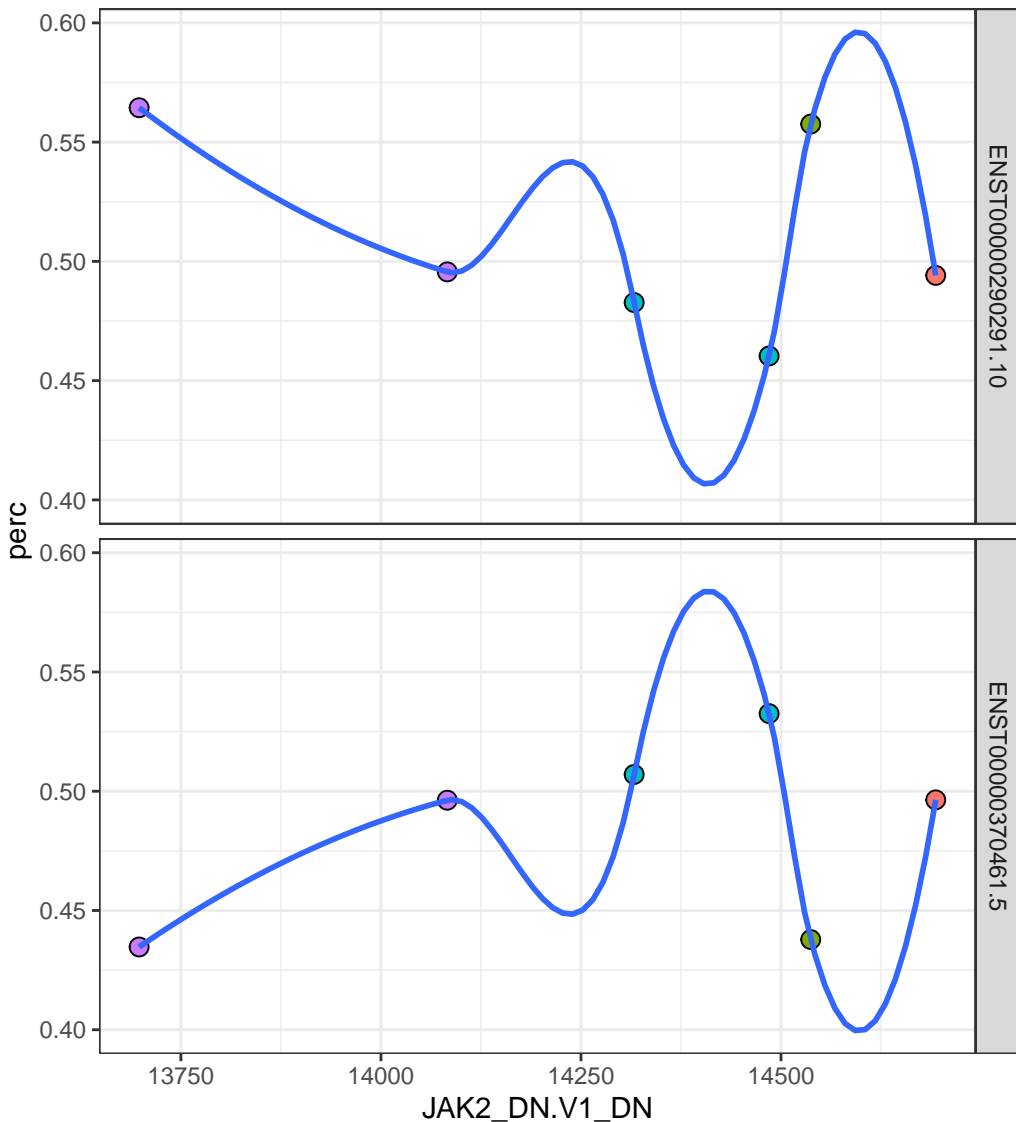
Target = JAK2_DN.V1_DN

ENST00000370461.5 (up) = -0.102291998128652

ENST00000290291.10 (down) = 0.0872343839895024



LTL_longitudinal
Gene = OGFR
Target = JAK2_DN.V1_DN
ENST00000370461.5 (up) = 0.425299166142721
ENST00000290291.10 (down) = -0.448677644088997



sampleType

- LTL331_Cx_12w
- LTL331_Cx_8w
- LTL331_Cx_R
- LTL331_PreCx

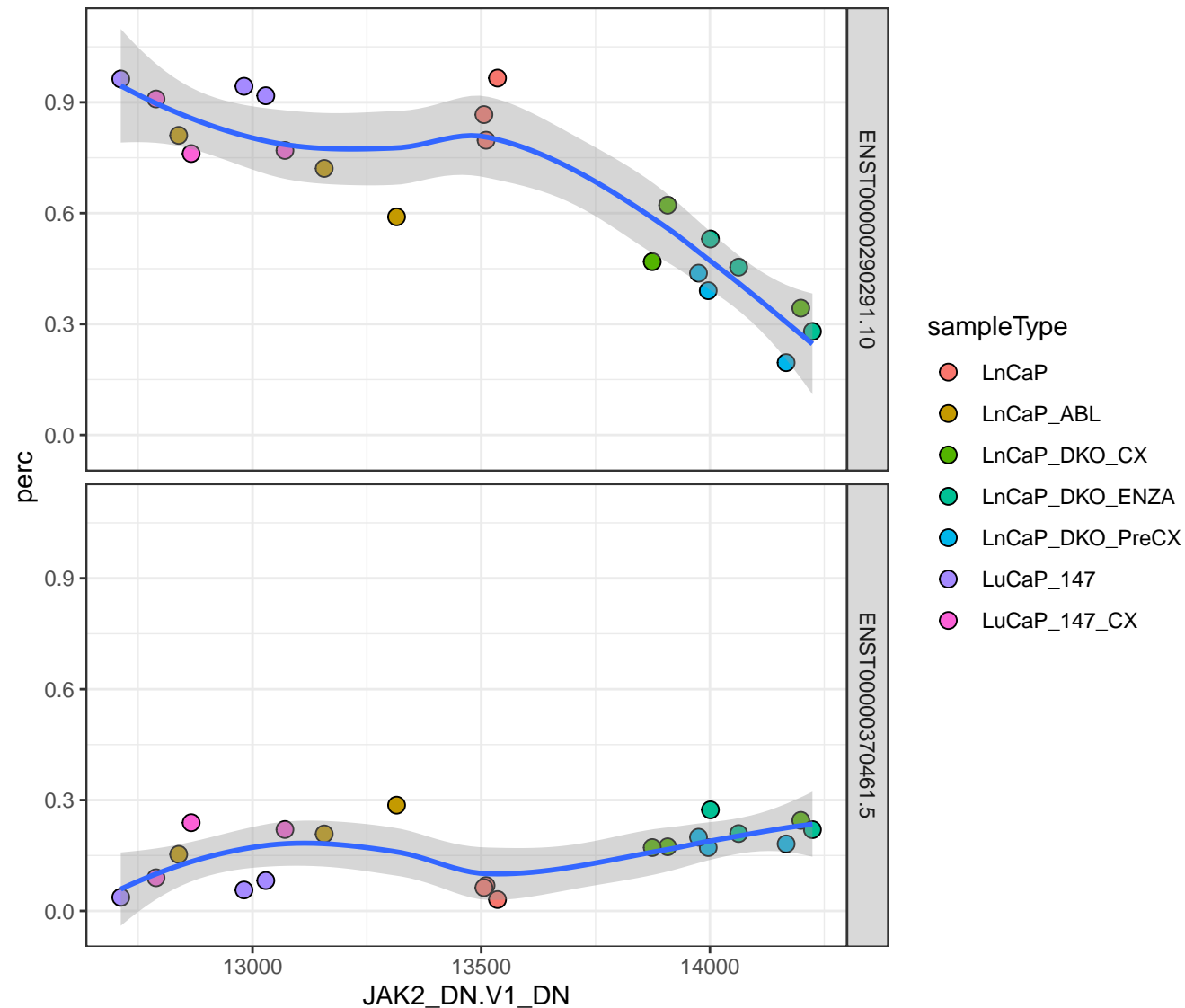
PDX_longitudinal

Gene = OGFR

Target = JAK2_DN.V1_DN

ENST00000370461.5 (up) = 0.420348161348527

ENST00000290291.10 (down) = -0.854006099434207



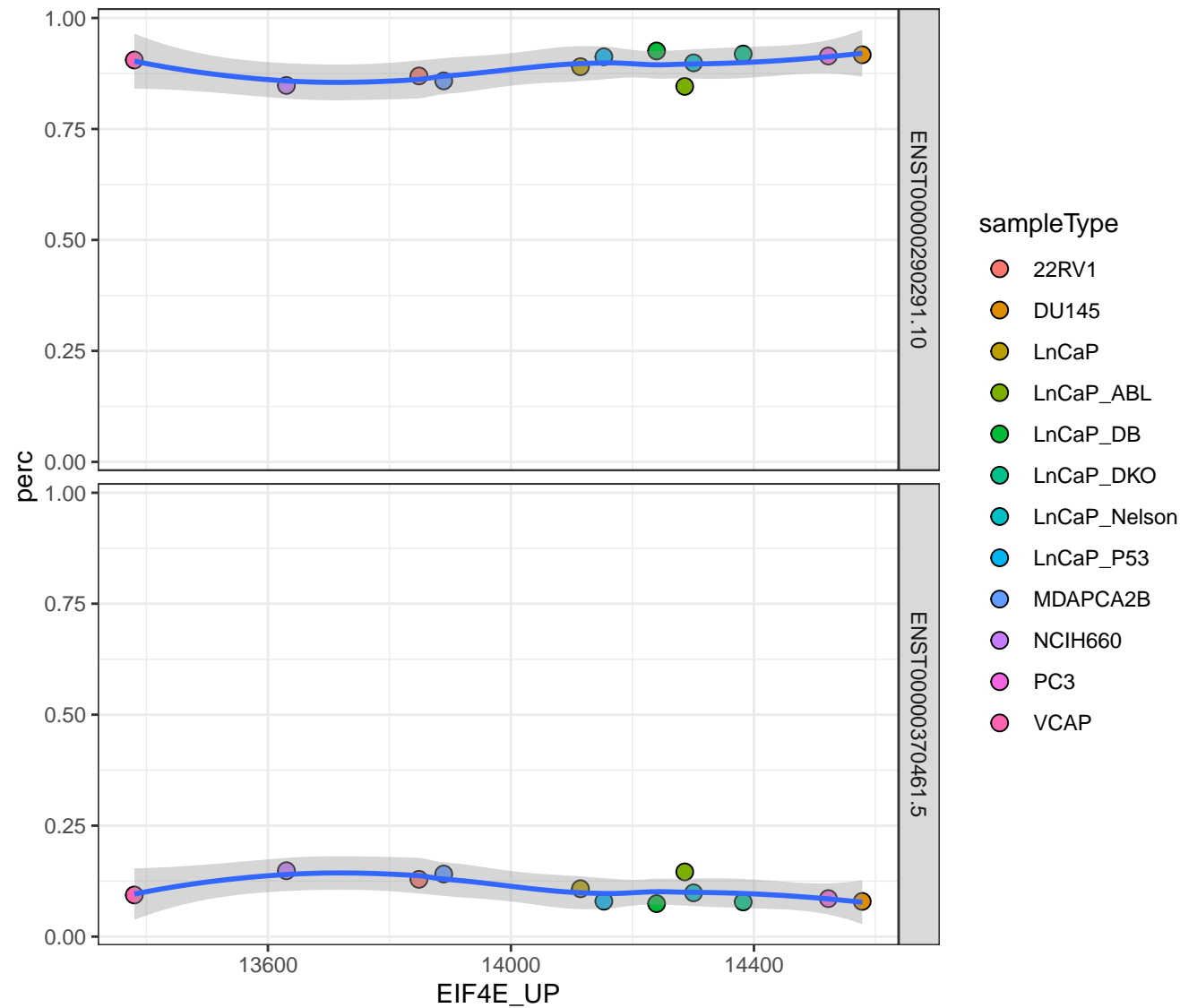
CellLine

Gene = OGFR

Target = EIF4E_UP

ENST00000290291.10 (up) = 0.444898777628031

ENST00000370461.5 (down) = -0.471641510122368



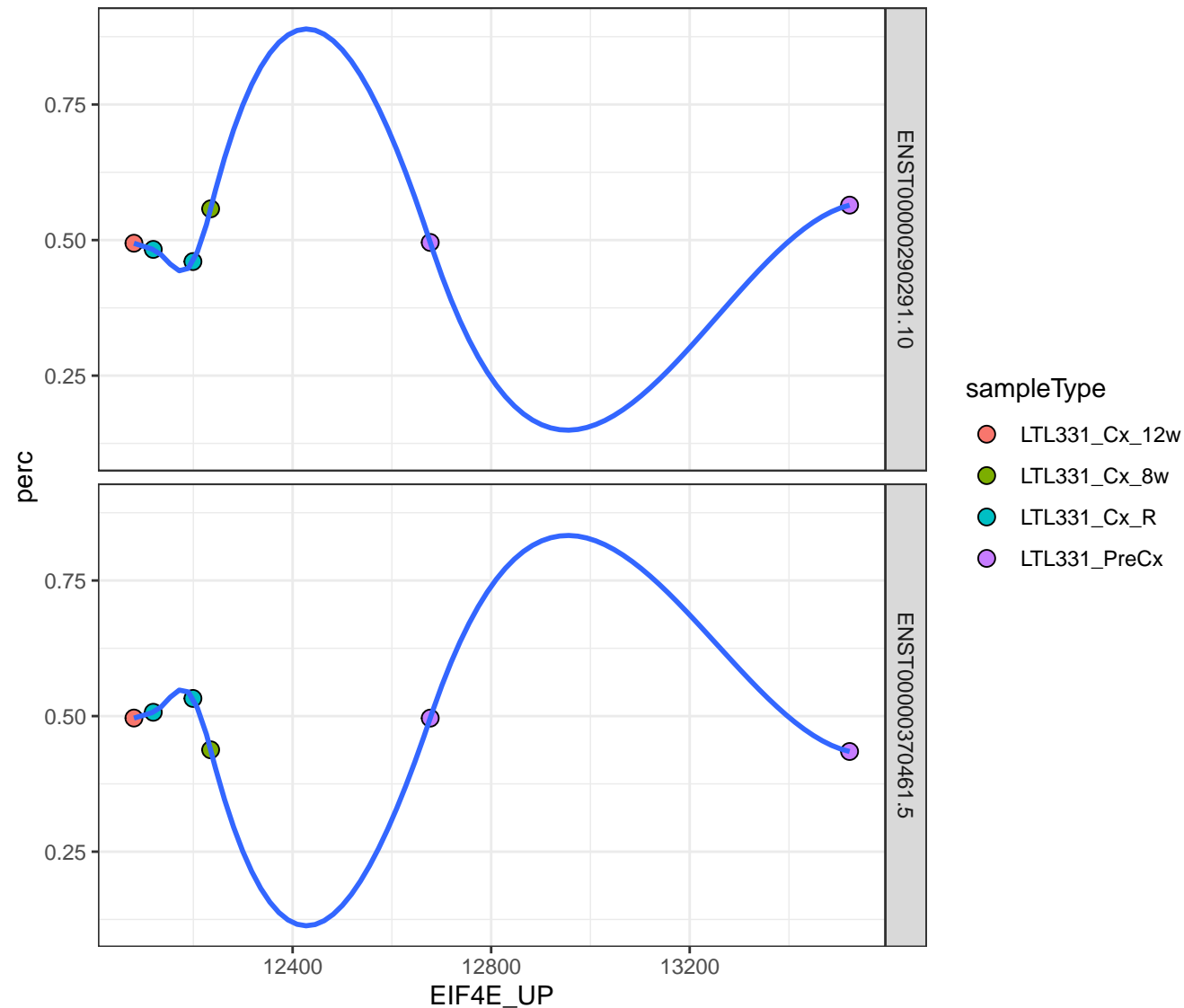
LTL_longitudinal

Gene = OGFR

Target = EIF4E_UP

ENST00000290291.10 (up) = 0.615287199730235

ENST00000370461.5 (down) = -0.587958667981171



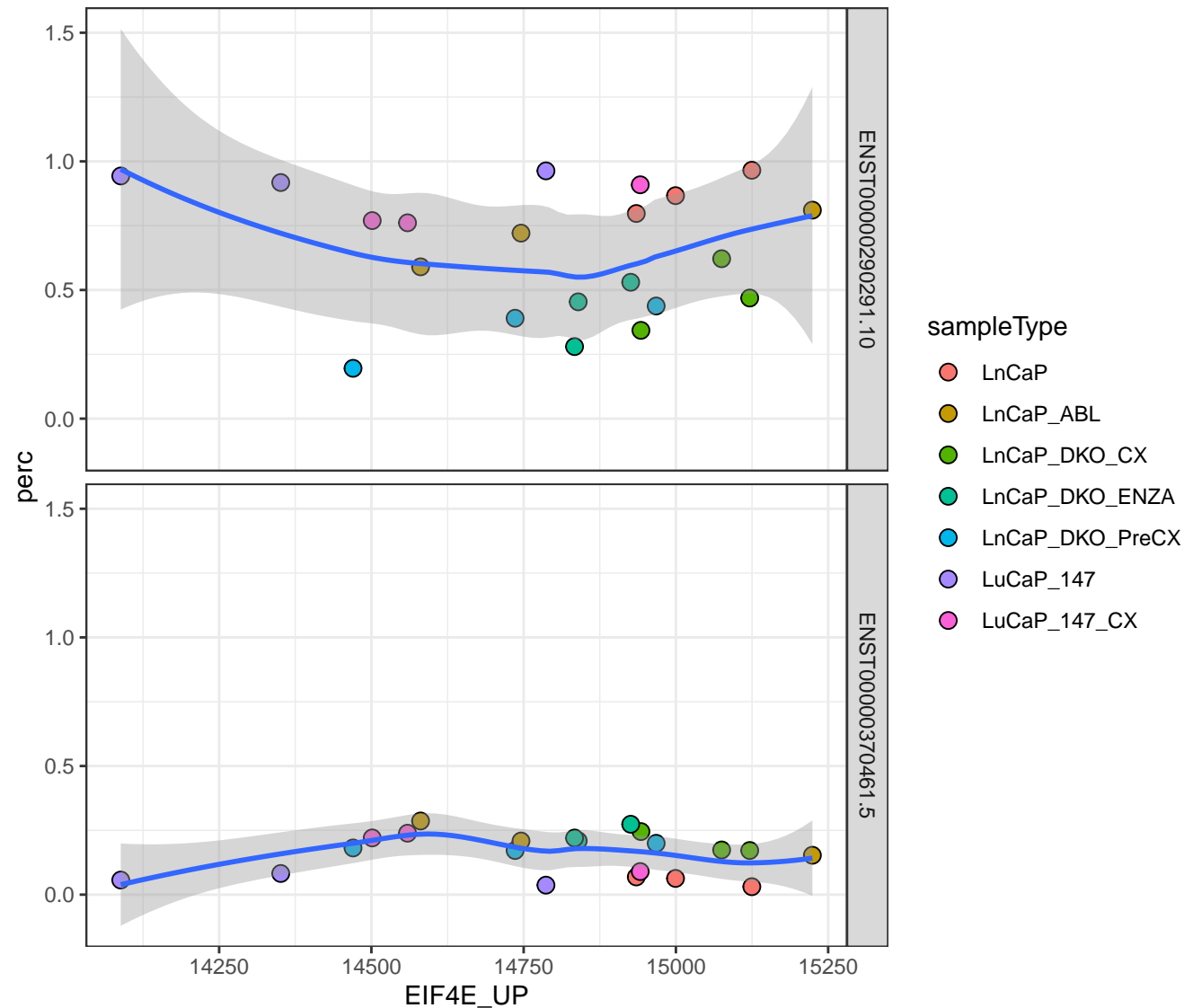
PDX_longitudinal

Gene = OGFR

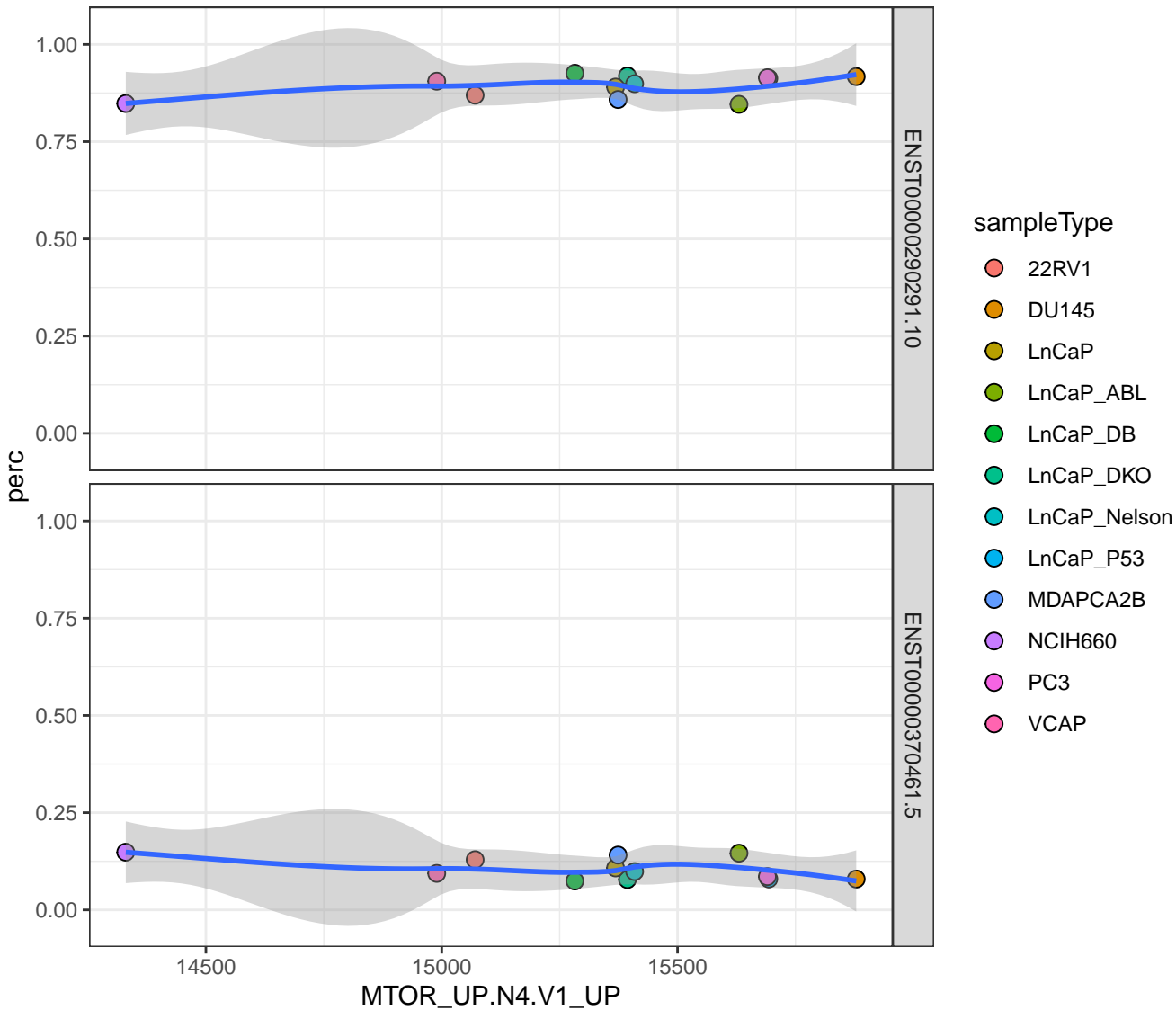
Target = EIF4E_UP

ENST00000290291.10 (up) = -0.0933450547289267

ENST00000370461.5 (down) = -0.0153831869817492



CellLine
 Gene = OGFR
 Target = MTOR_UP.N4.V1_UP
 ENST00000290291.10 (up) = 0.455255045185016
 ENST00000370461.5 (down) = -0.48905992967045



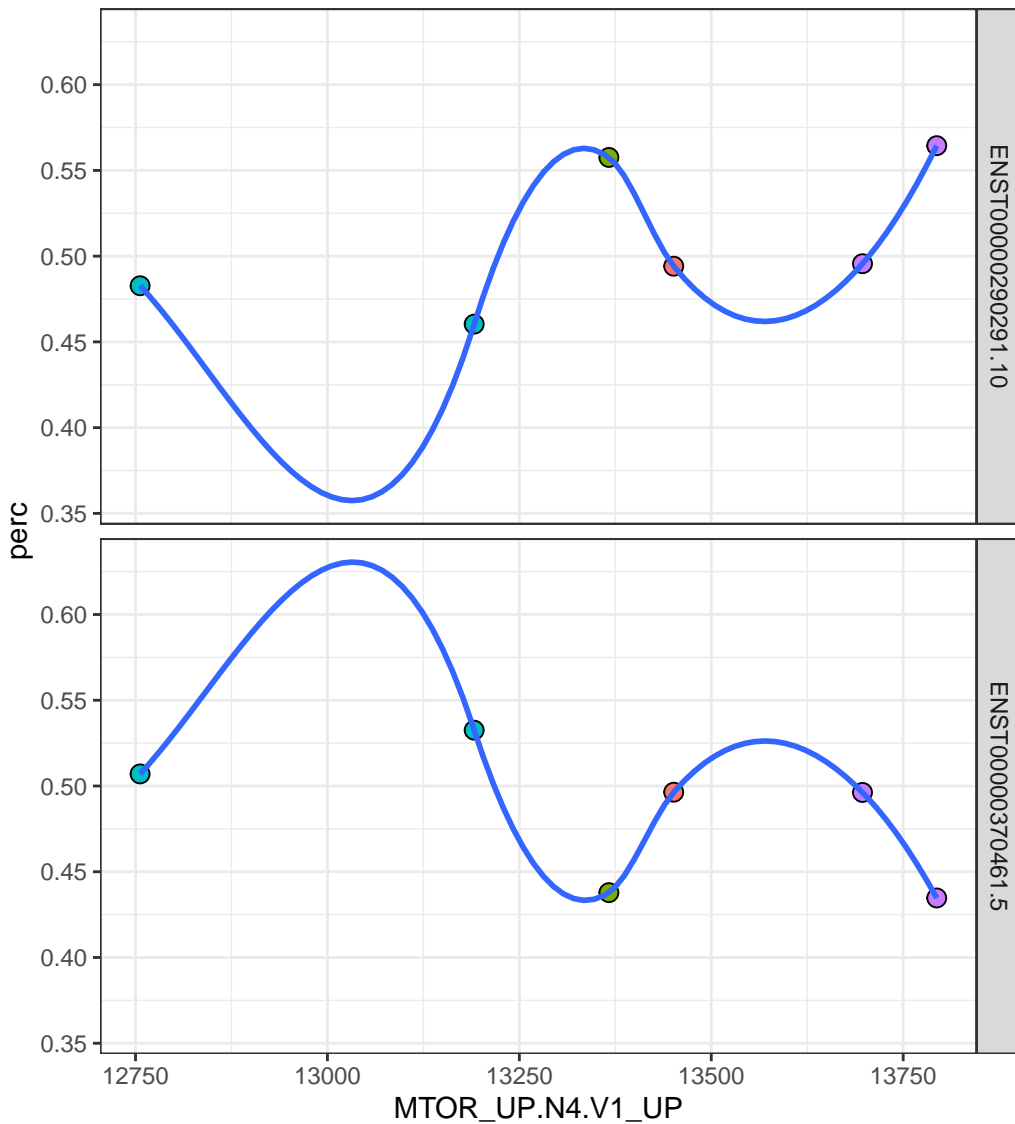
LTL_longitudinal

Gene = OGFR

Target = MTOR_UP.N4.V1_UP

ENST00000290291.10 (up) = 0.538808381983965

ENST00000370461.5 (down) = -0.522757787959842



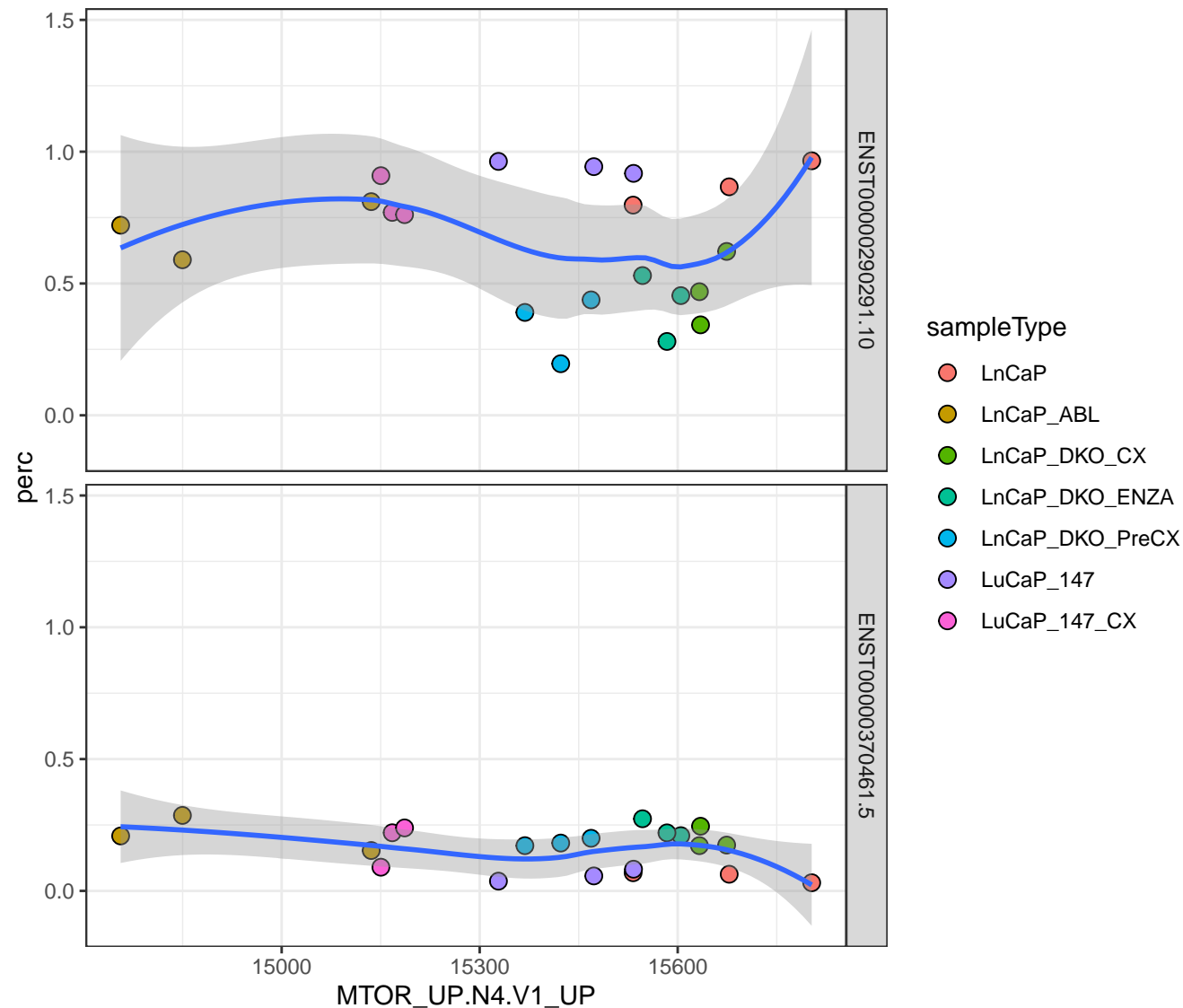
PDX_longitudinal

Gene = OGFR

Target = MTOR_UP.N4.V1_UP

ENST00000290291.10 (up) = -0.145017391332887

ENST00000370461.5 (down) = -0.329417260257756



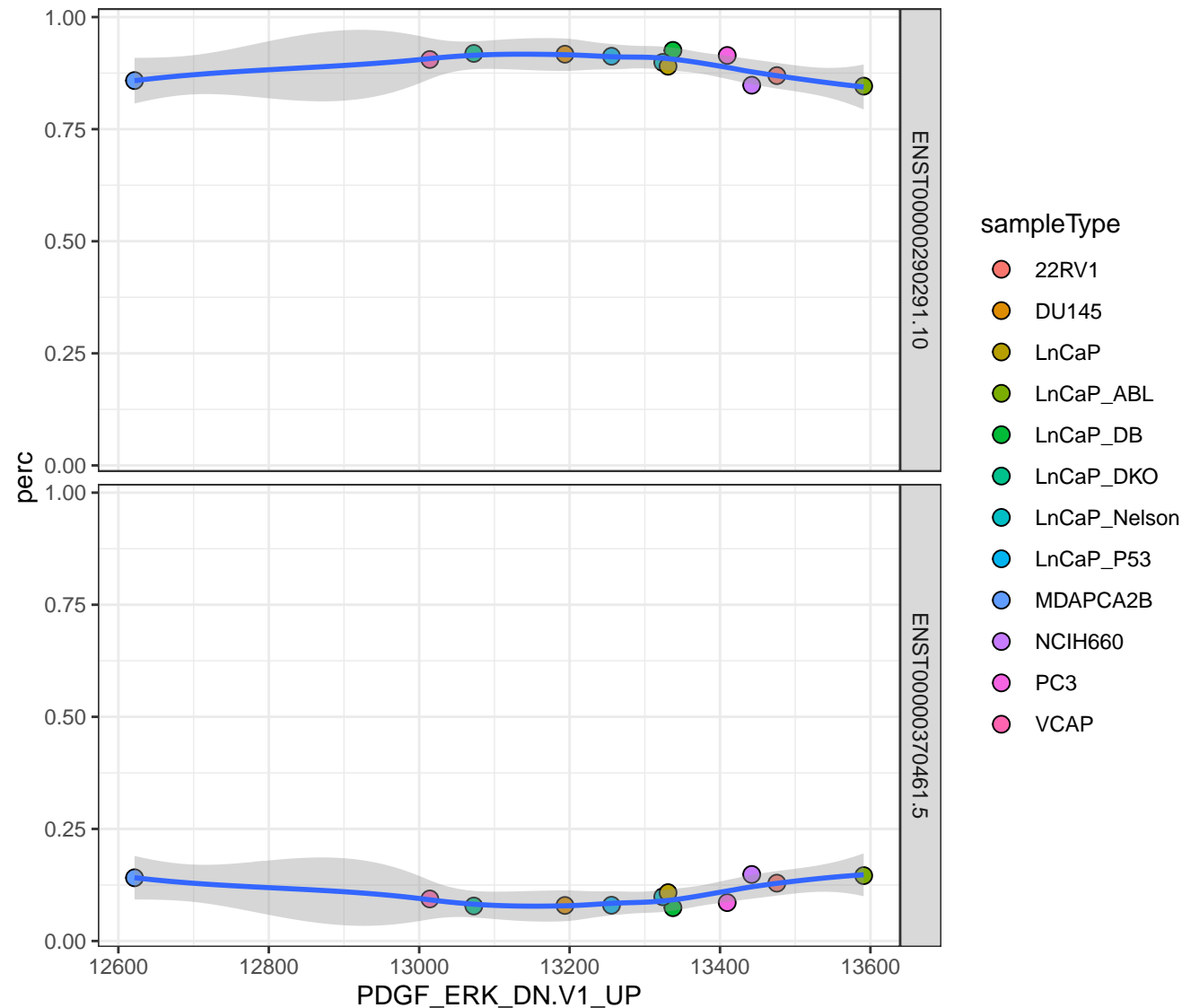
CellLine

Gene = OGFR

Target = PDGF_ERK_DN.V1_UP

ENST00000290291.10 (up) = -0.126121370439868

ENST00000370461.5 (down) = 0.0959590255891028



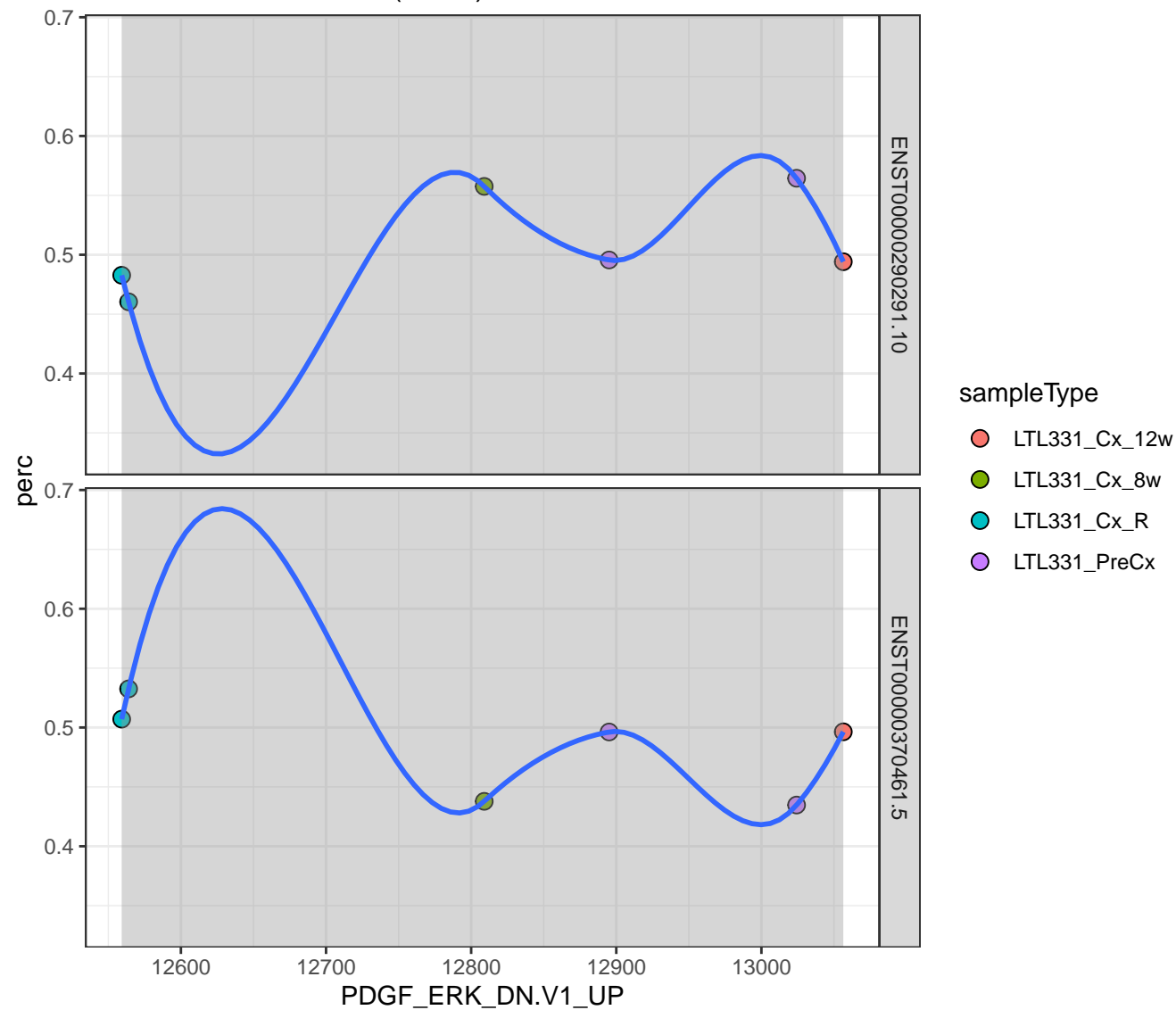
LTL_longitudinal

Gene = OGFR

Target = PDGF_ERK_DN.V1_UP

ENST00000290291.10 (up) = 0.557247600717925

ENST00000370461.5 (down) = -0.563529810791444



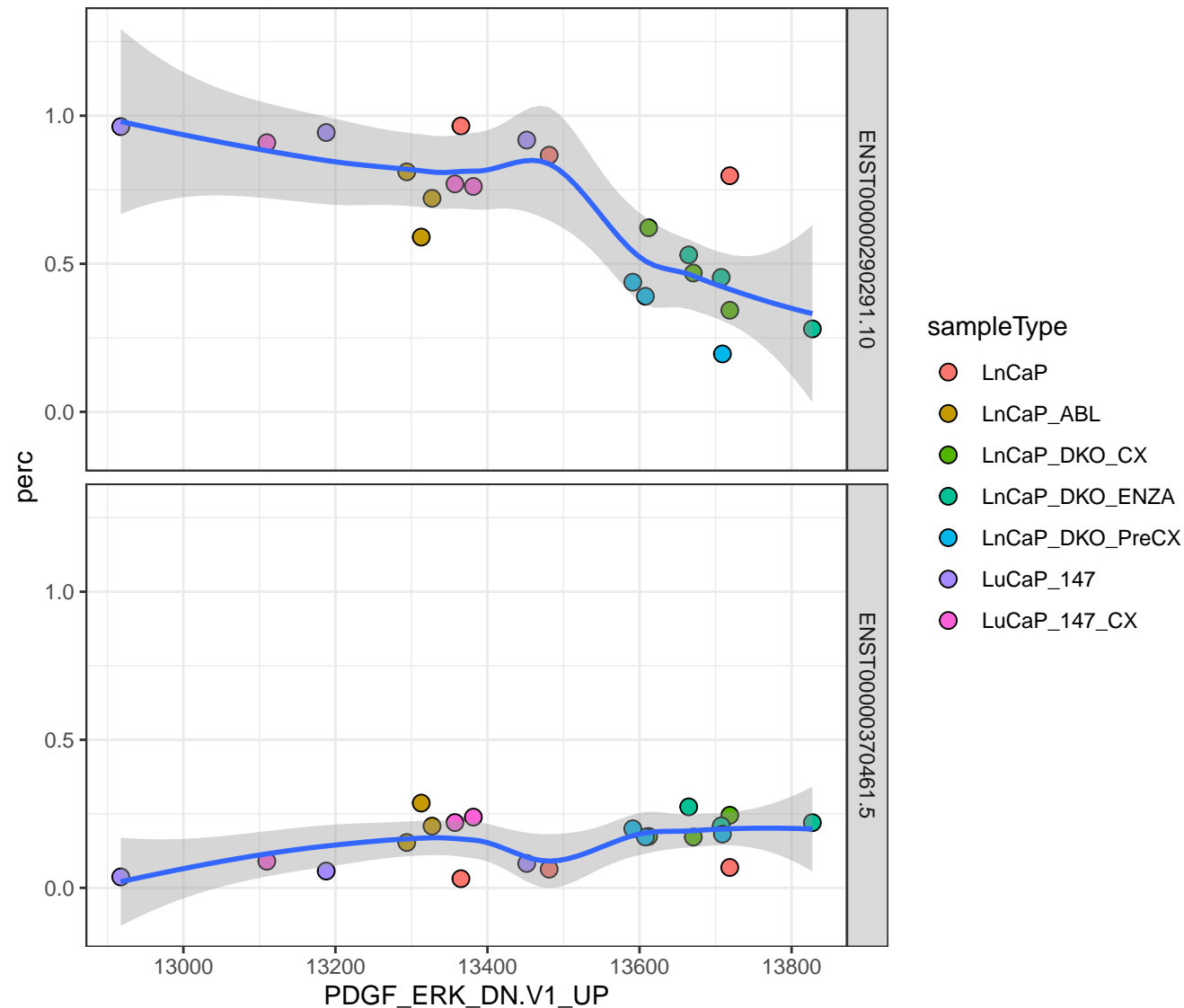
PDX_longitudinal

Gene = OGFR

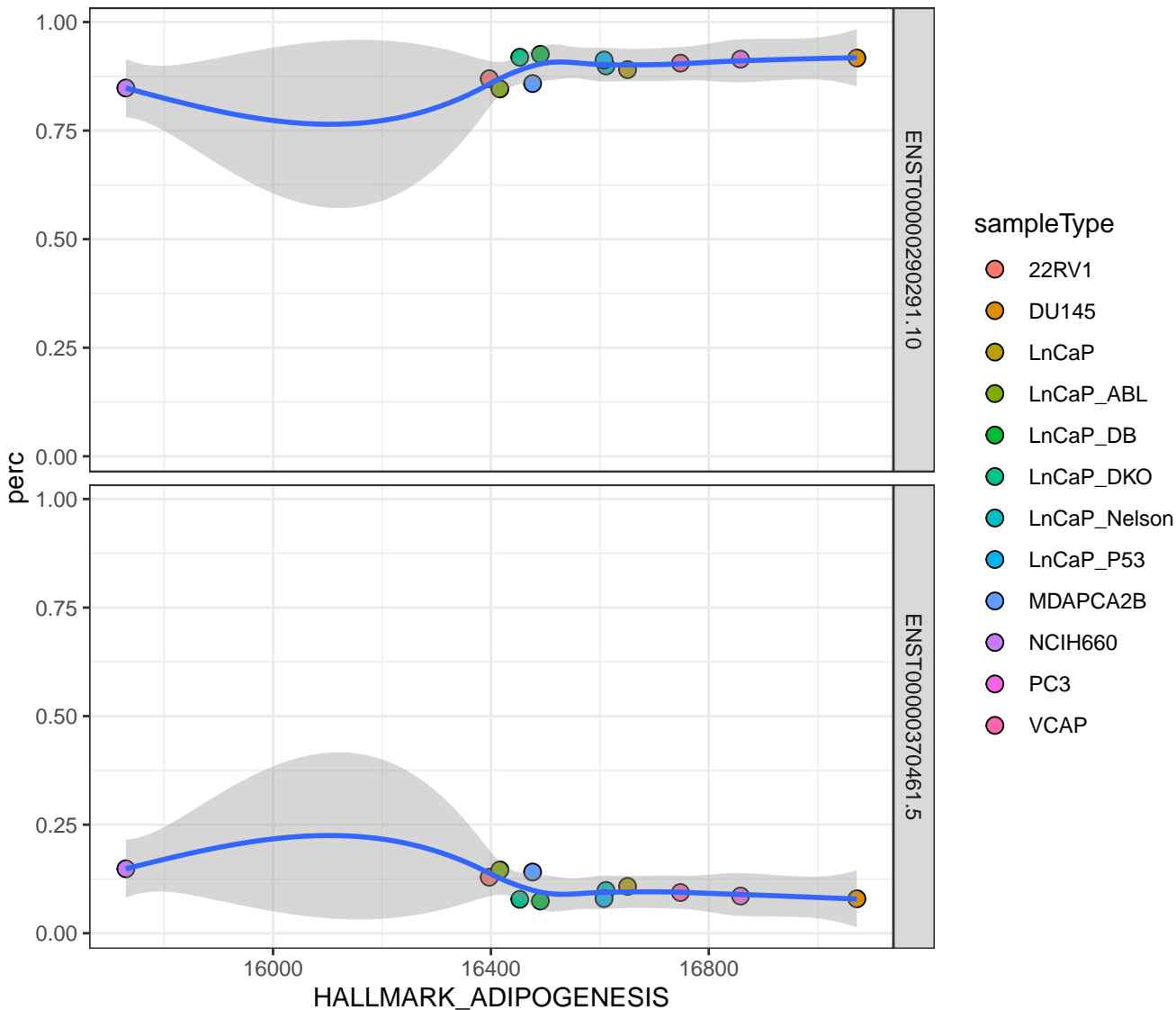
Target = PDGF_ERK_DN.V1_UP

ENST00000290291.10 (up) = -0.771849248668779

ENST00000370461.5 (down) = 0.439713718781772



CellLine
 Gene = OGFR
 Target = HALLMARK_ADIPOGENESIS
 ENST00000290291.10 (up) = 0.651144518924058
 ENST00000370461.5 (down) = -0.650303230356284



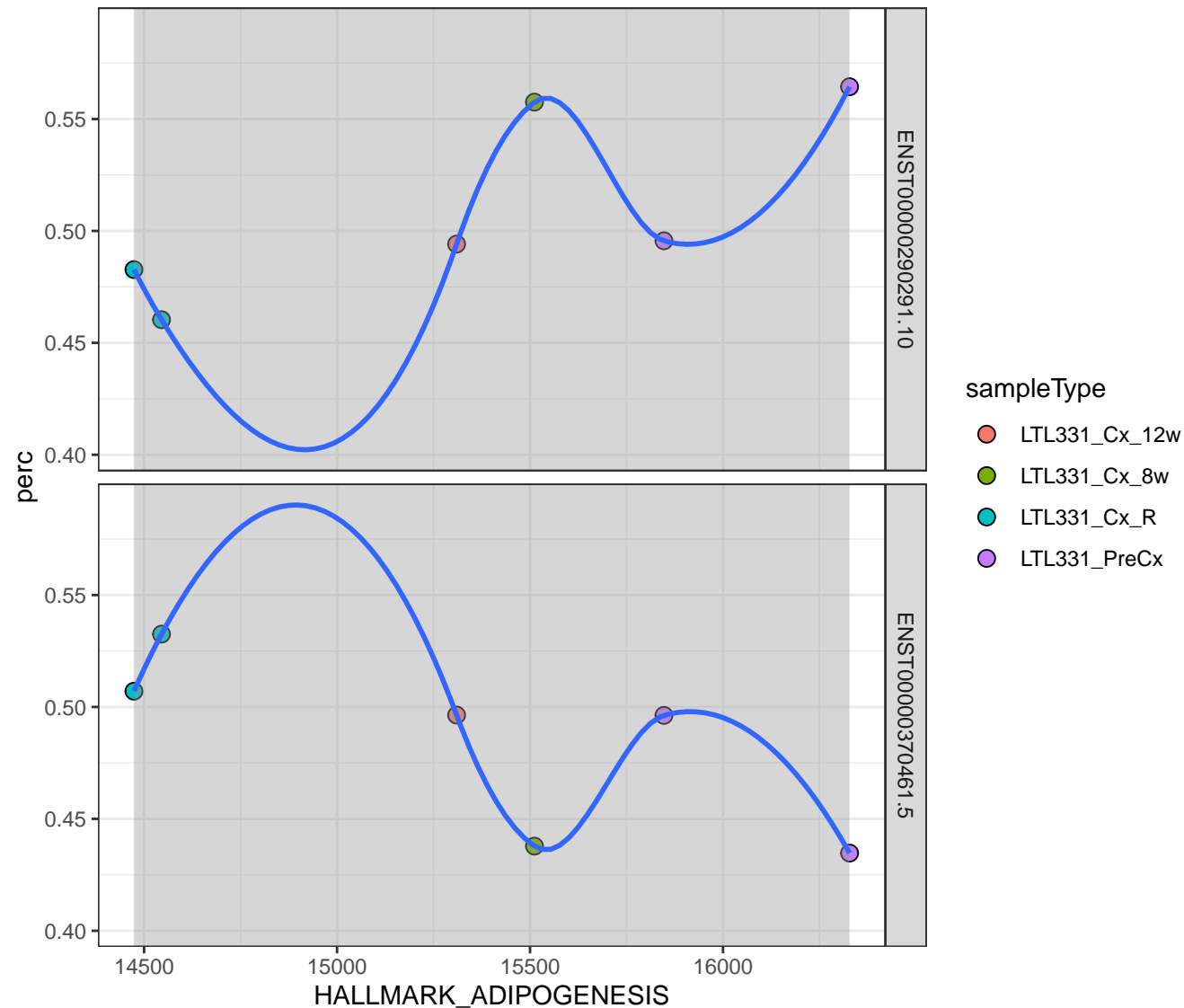
LTL_longitudinal

Gene = OGFR

Target = HALLMARK_ADIPOGENESIS

ENST00000290291.10 (up) = 0.770601922019127

ENST00000370461.5 (down) = -0.763433117221376



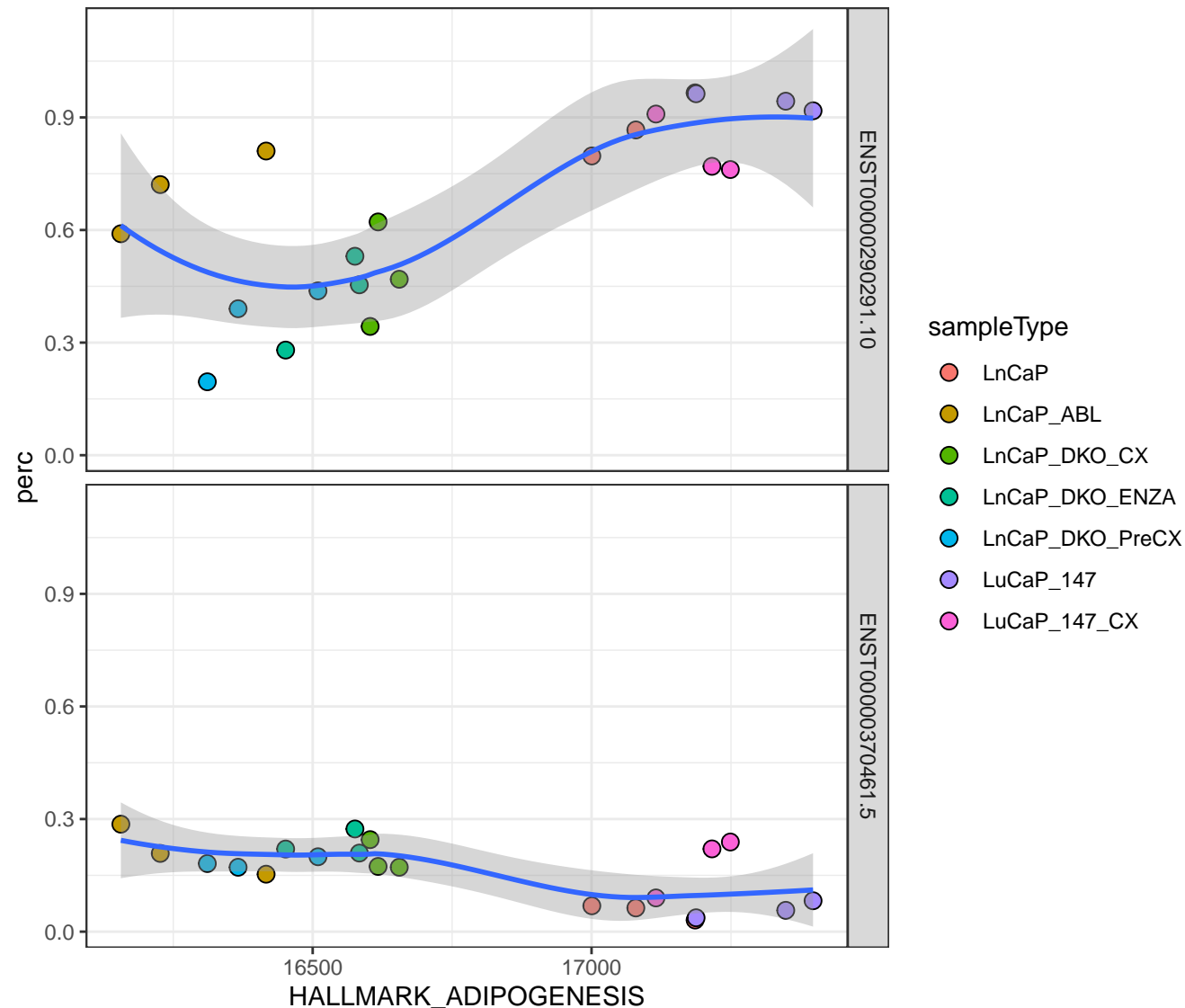
PDX_longitudinal

Gene = OGFR

Target = HALLMARK_ADIPOGENESIS

ENST00000290291.10 (up) = 0.743388683209129

ENST00000370461.5 (down) = -0.648805235606019



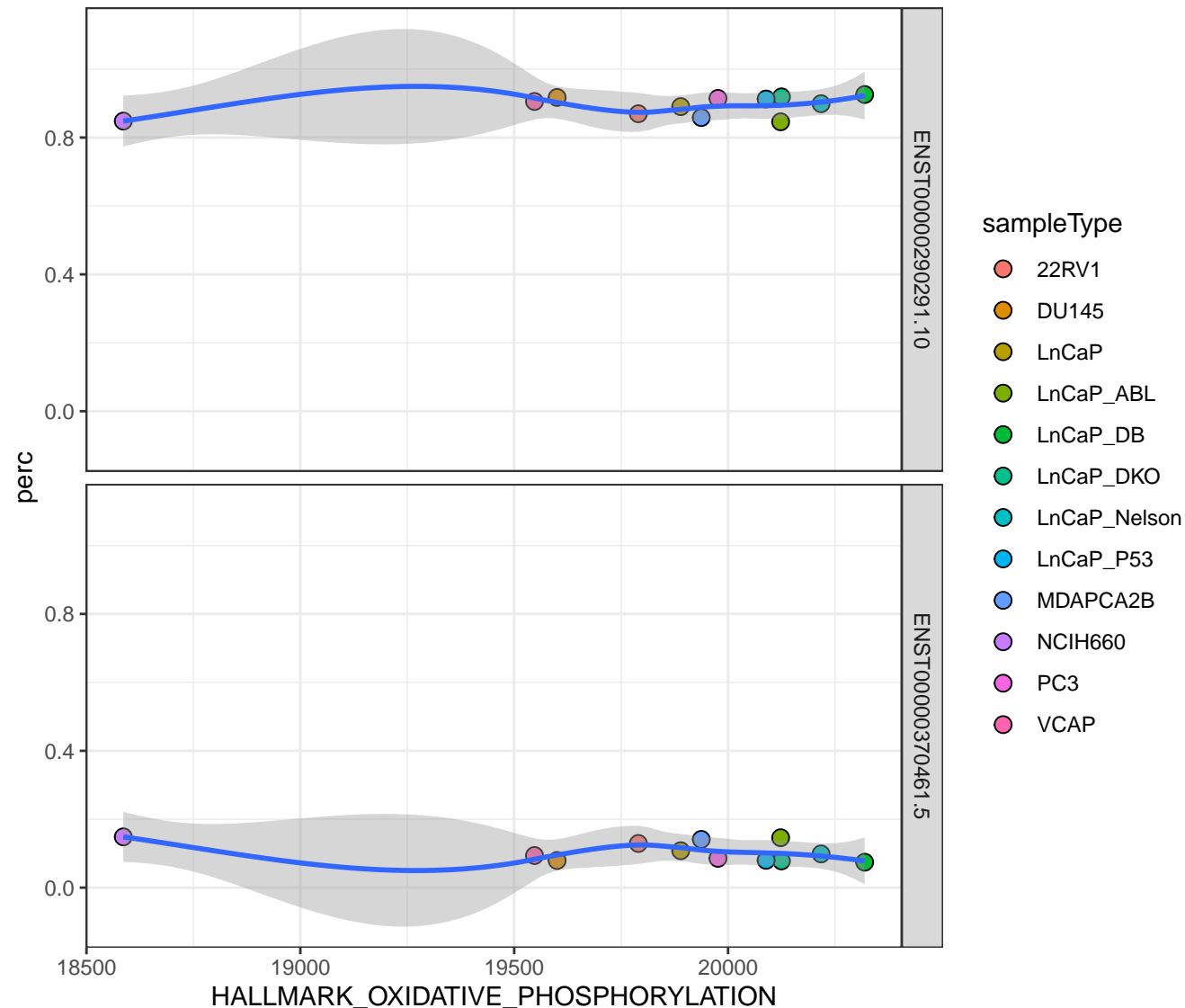
CellLine

Gene = OGFR

Target = HALLMARK_OXIDATIVE_PHOSPHORYLATION

ENST00000290291.10 (up) = 0.435835867362236

ENST00000370461.5 (down) = -0.447463700014896



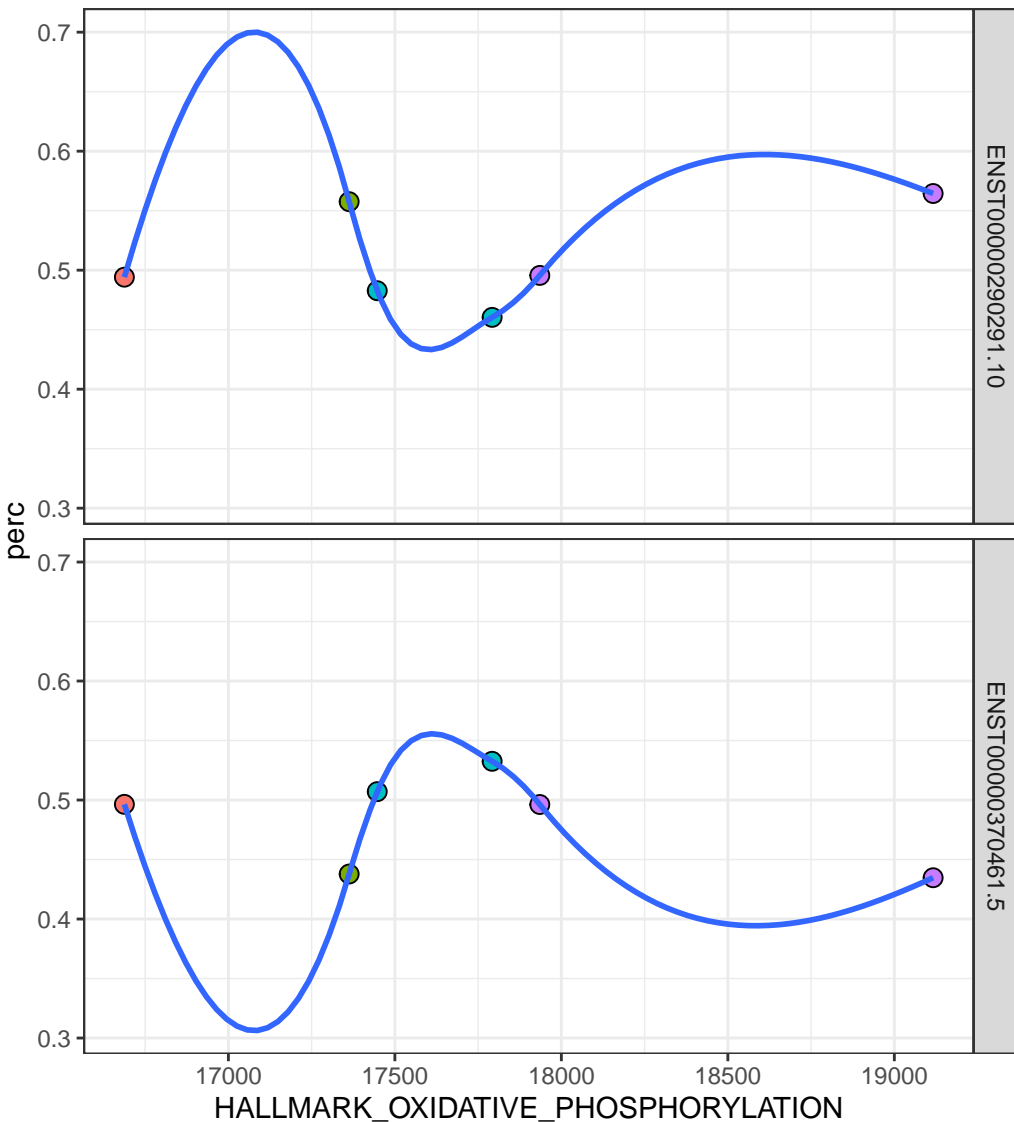
LTL_longitudinal

Gene = OGFR

Target = HALLMARK_OXIDATIVE_PHOSPHORYLATION

ENST00000290291.10 (up) = 0.446873405973876

ENST00000370461.5 (down) = -0.410026047732745



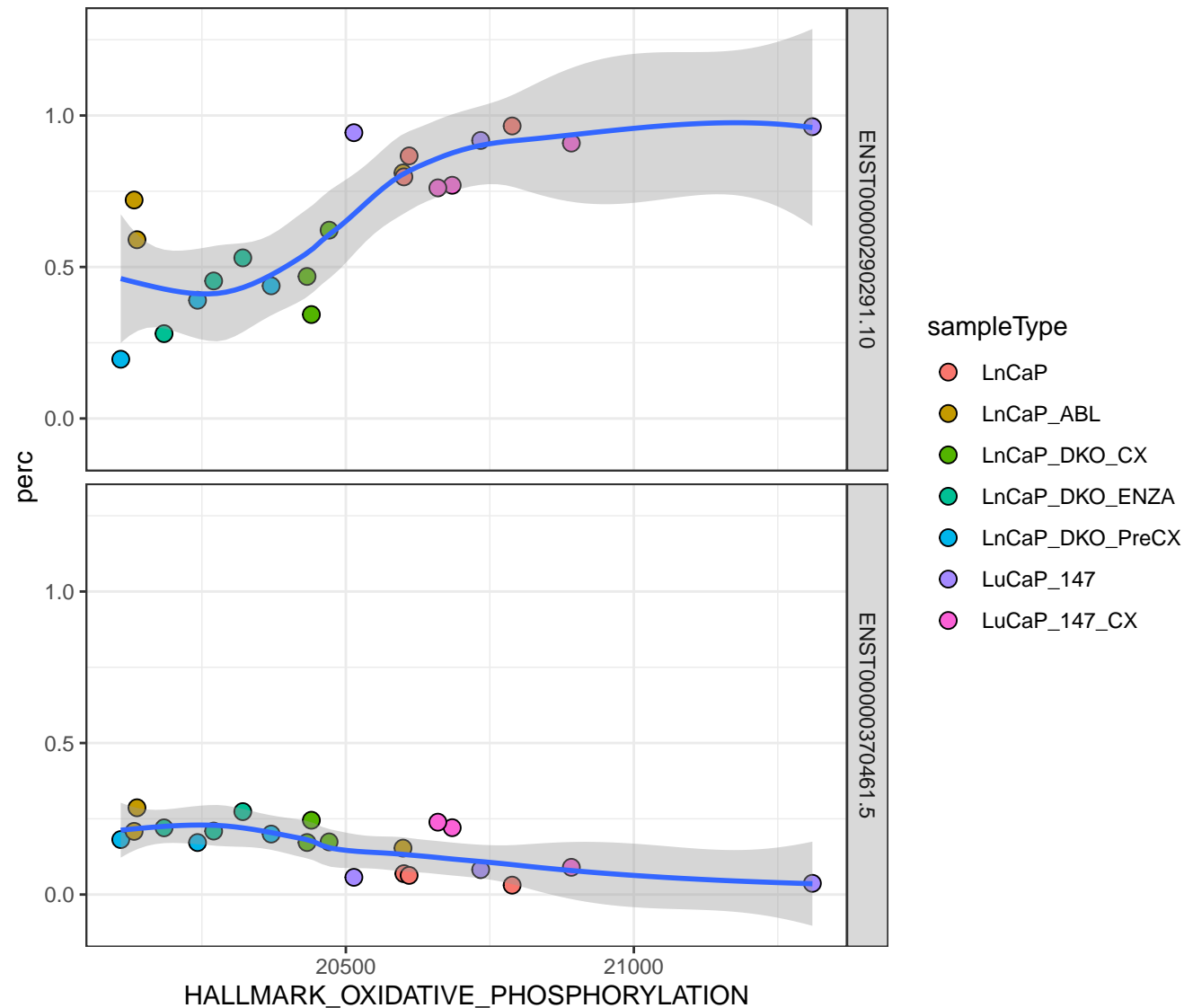
PDX_longitudinal

Gene = OGFR

Target = HALLMARK_OXIDATIVE_PHOSPHORYLATION

ENST00000290291.10 (up) = 0.75879460869403

ENST00000370461.5 (down) = -0.674372354052035



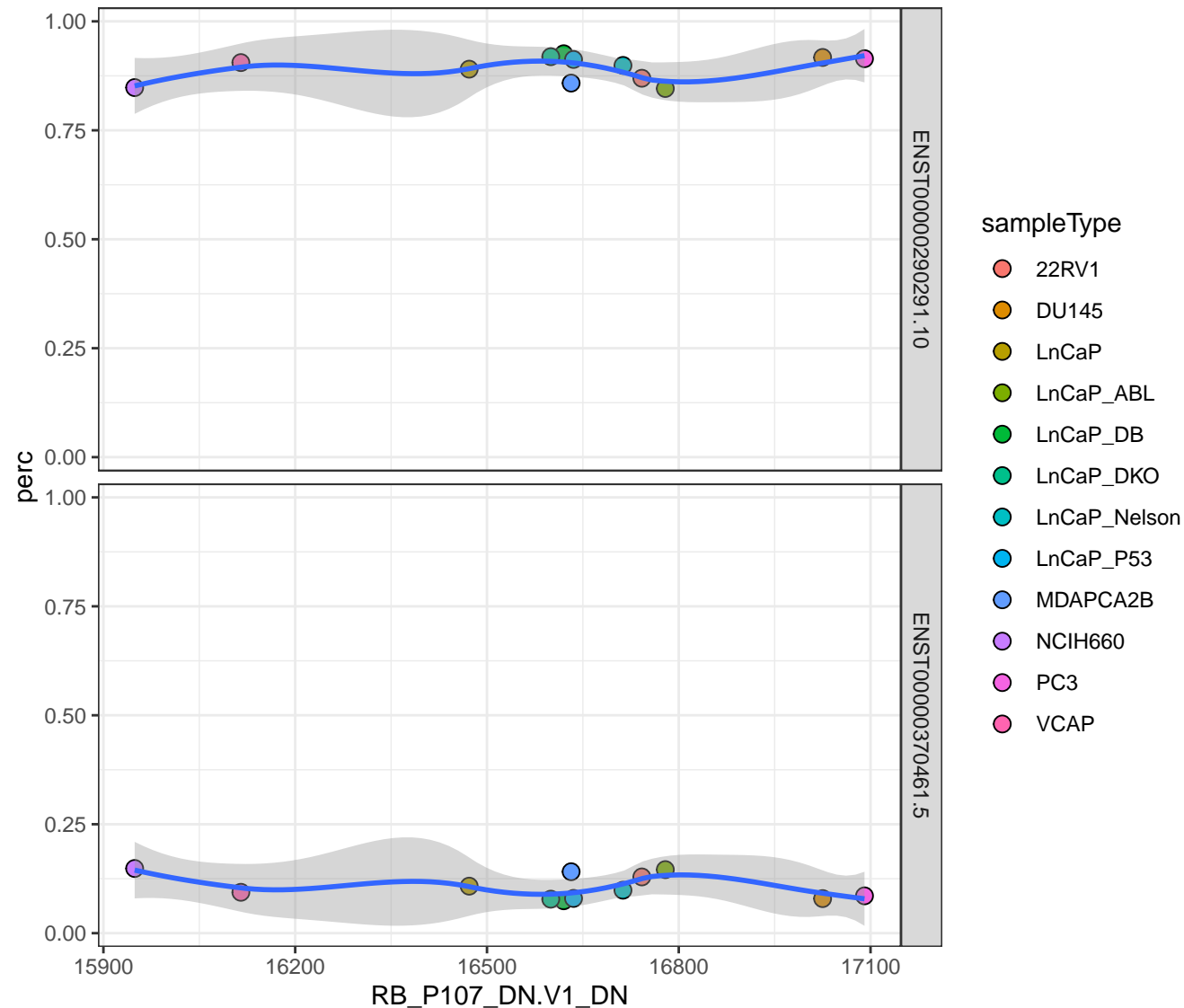
CellLine

Gene = OGFR

Target = RB_P107_DN.V1_DN

ENST00000290291.10 (up) = 0.32304034354629

ENST00000370461.5 (down) = -0.333640251661457



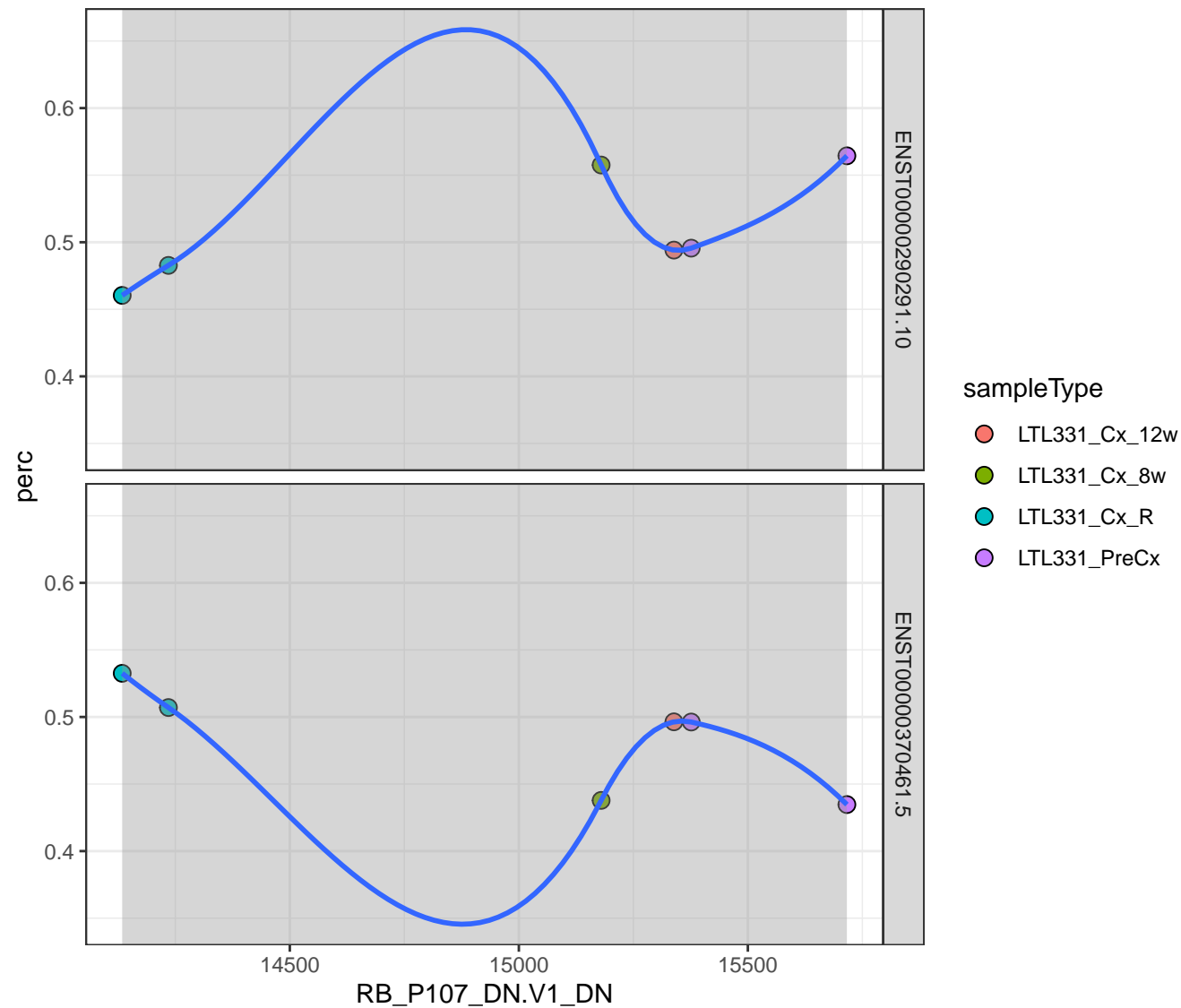
LTL_longitudinal

Gene = OGFR

Target = RB_P107_DN.V1_DN

ENST00000290291.10 (up) = 0.729766376714301

ENST00000370461.5 (down) = -0.733996746917397



PDX_longitudinal

Gene = OGFR

Target = RB_P107_DN.V1_DN

ENST00000290291.10 (up) = -0.133257202706242

ENST00000370461.5 (down) = -0.21935028495045

