

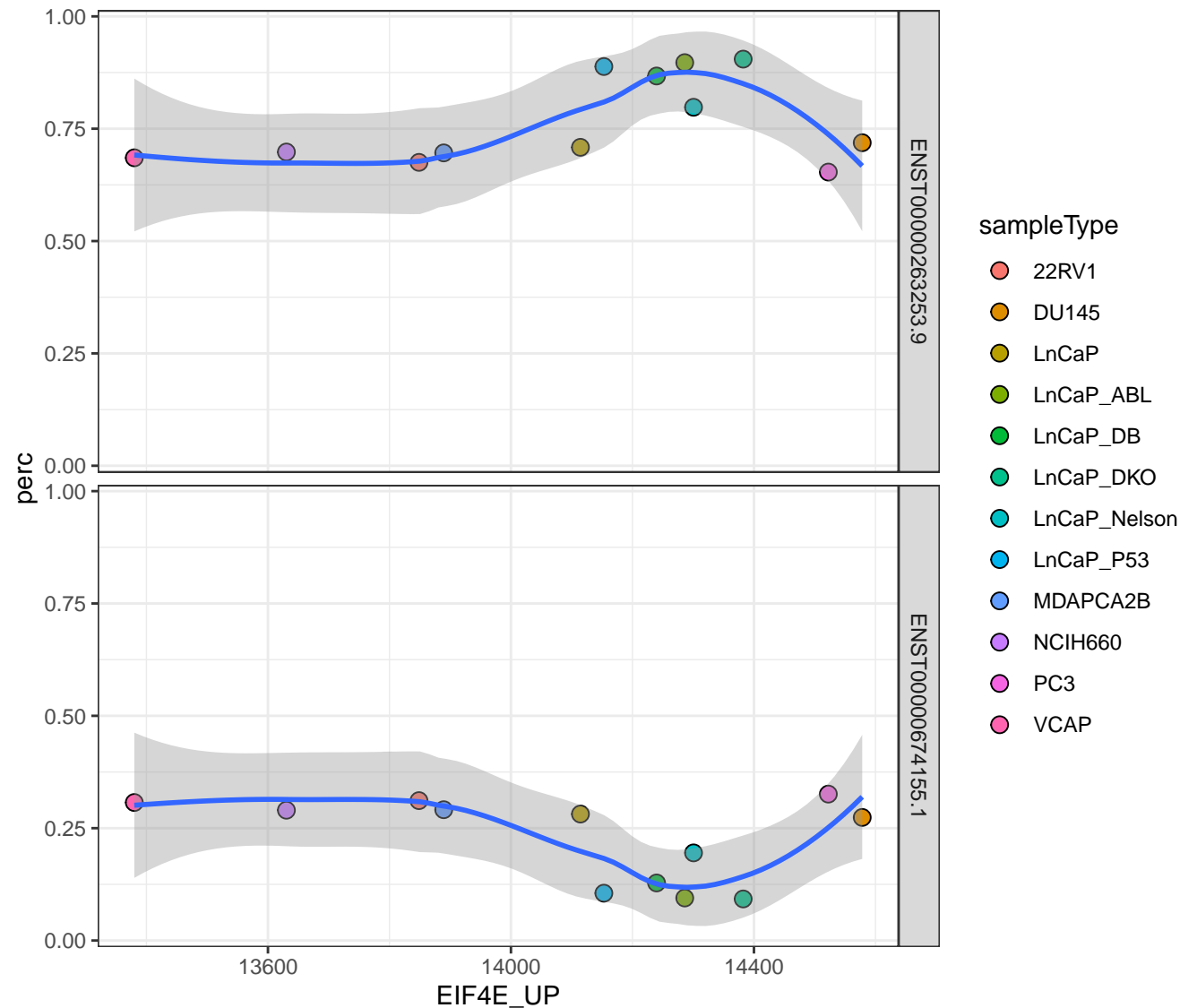
CellLine

Gene = EP300

Target = EIF4E_UP

ENST00000674155.1 (up) = -0.390592386393865

ENST00000263253.9 (down) = 0.379855125921124



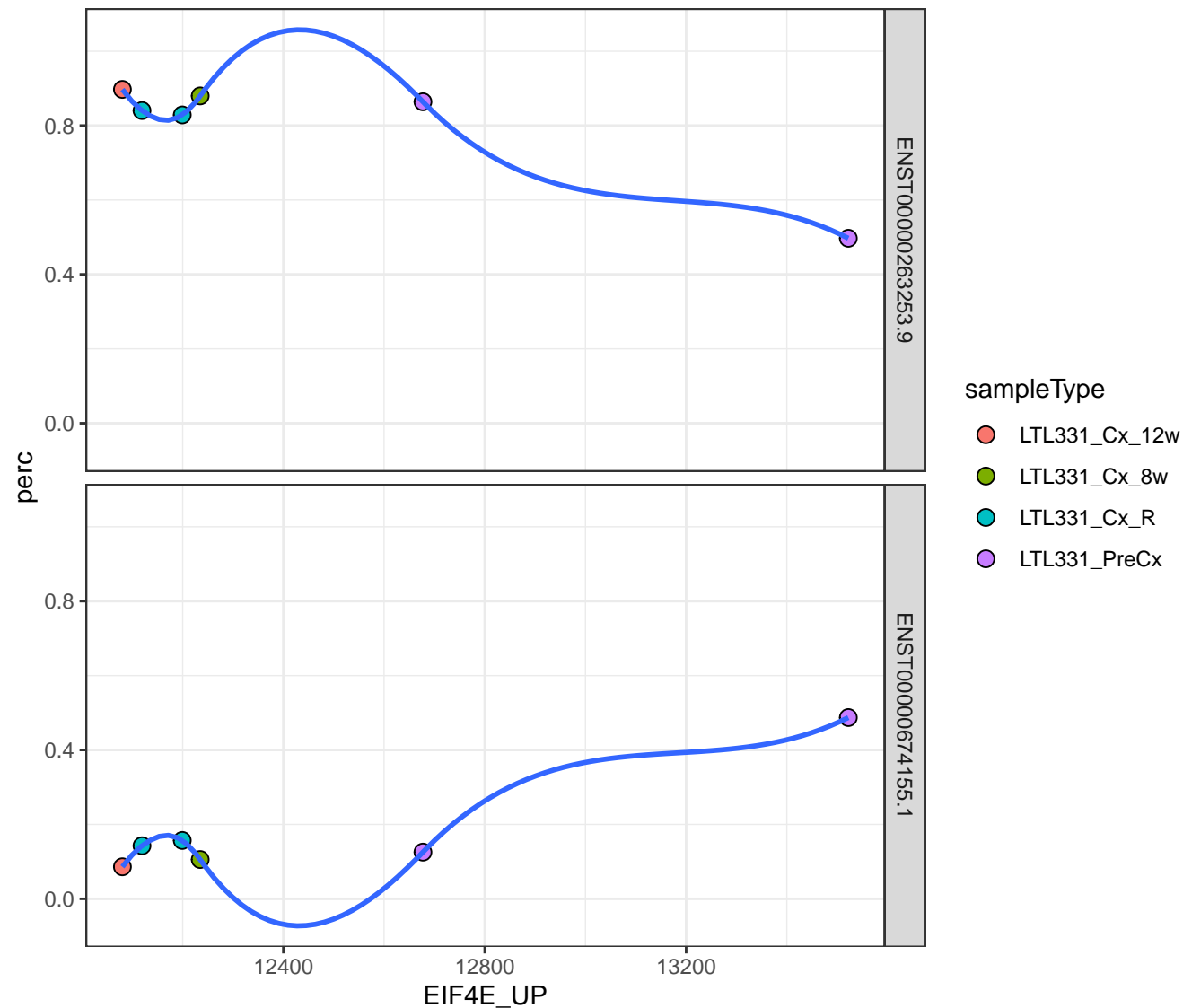
LTL_longitudinal

Gene = EP300

Target = EIF4E_UP

ENST00000674155.1 (up) = 0.917532725433417

ENST00000263253.9 (down) = -0.912123703079173



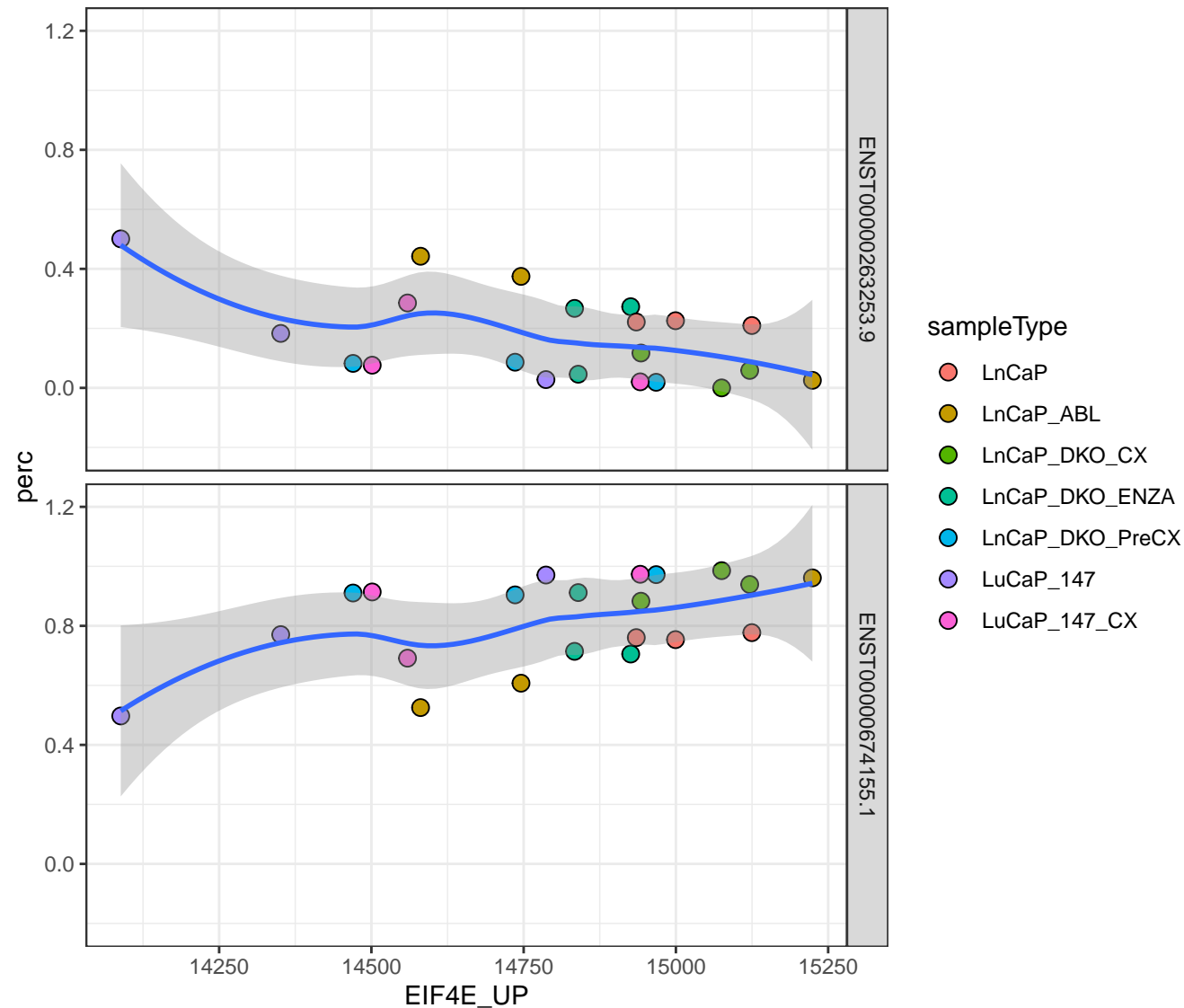
PDX_longitudinal

Gene = EP300

Target = EIF4E_UP

ENST00000674155.1 (up) = 0.514978333663679

ENST00000263253.9 (down) = -0.512318013455265



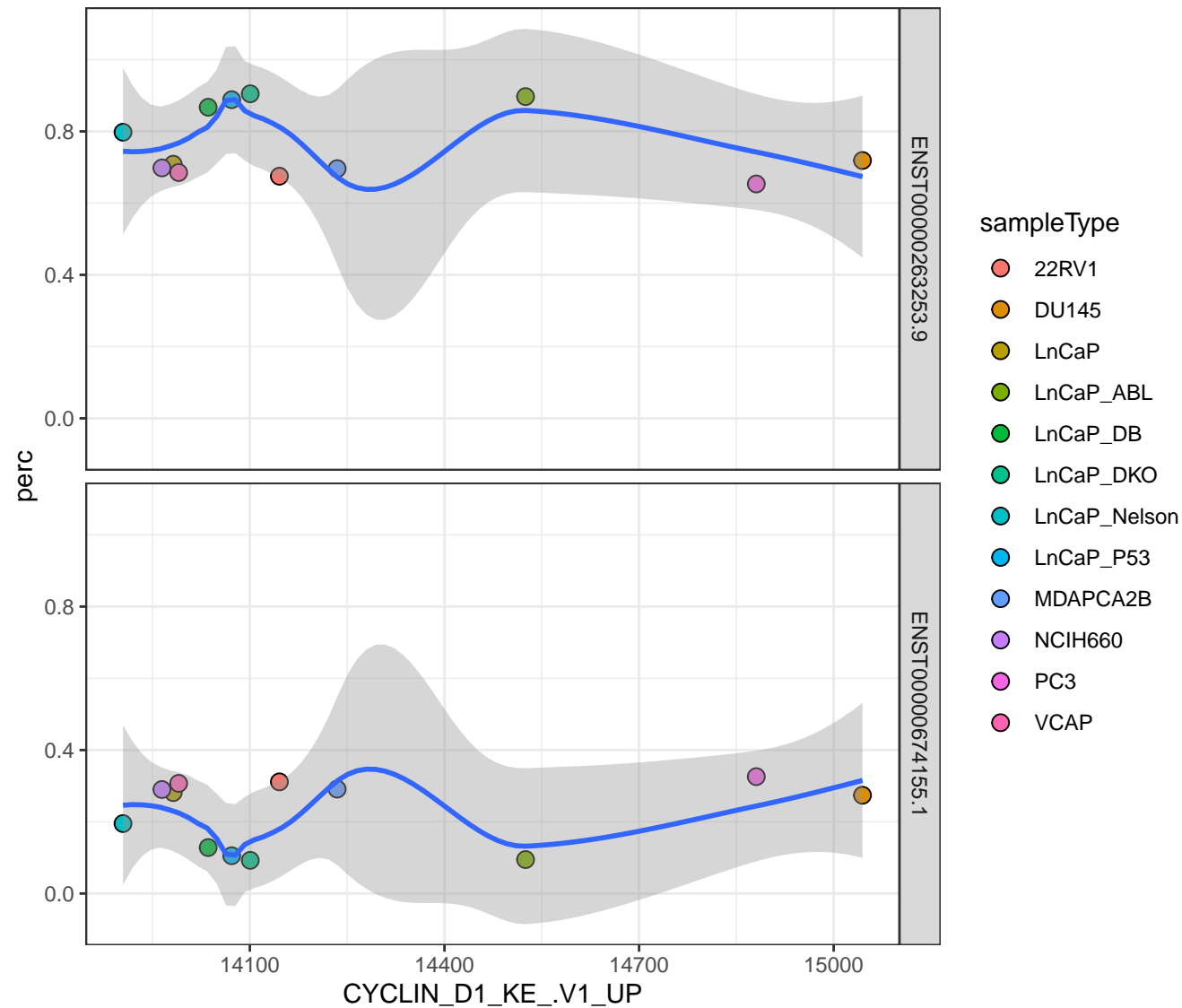
CellLine

Gene = EP300

Target = CYCLIN_D1_KE_.V1_UP

ENST00000674155.1 (up) = 0.189269618953381

ENST00000263253.9 (down) = -0.200943335001907



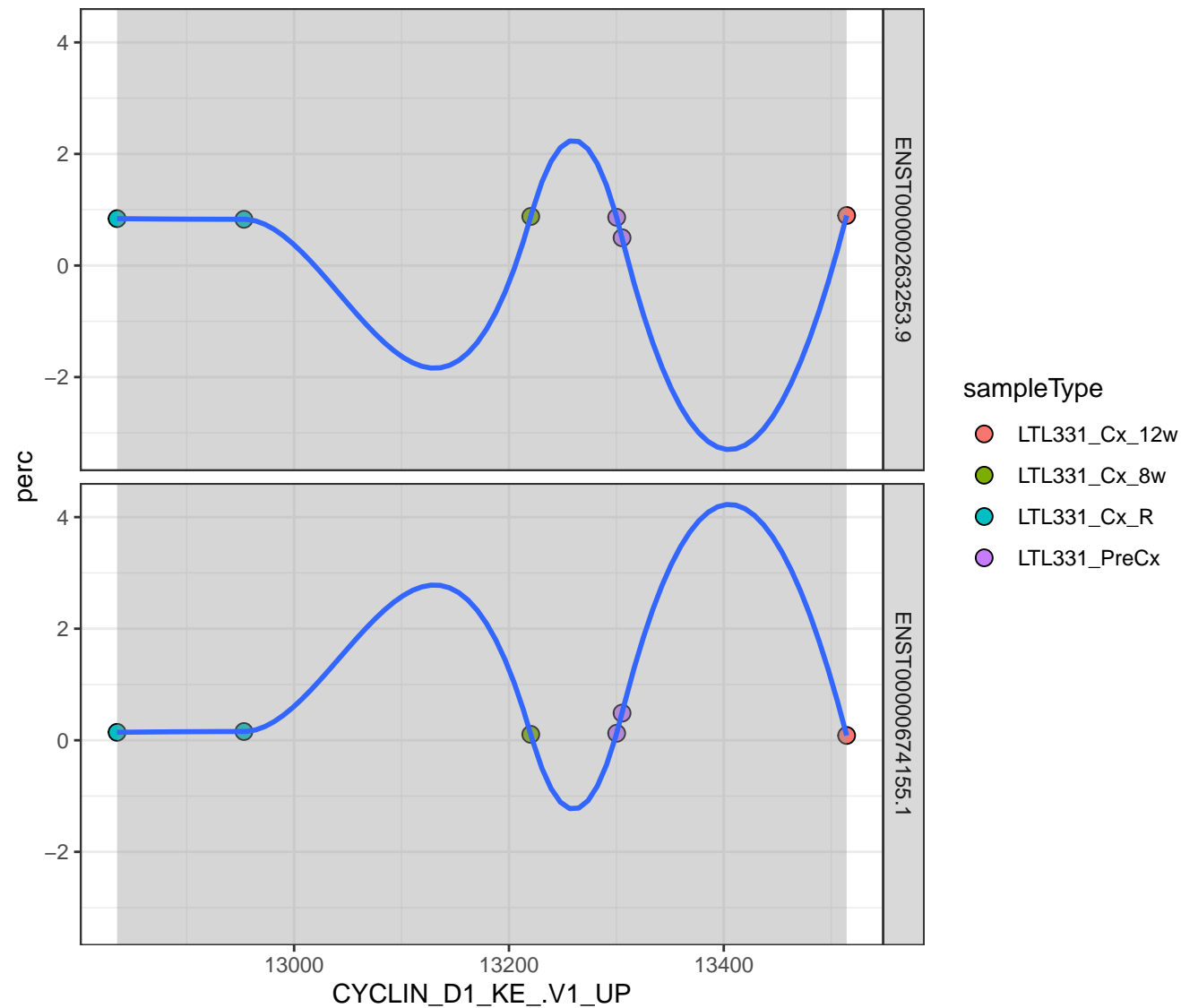
LTL_longitudinal

Gene = EP300

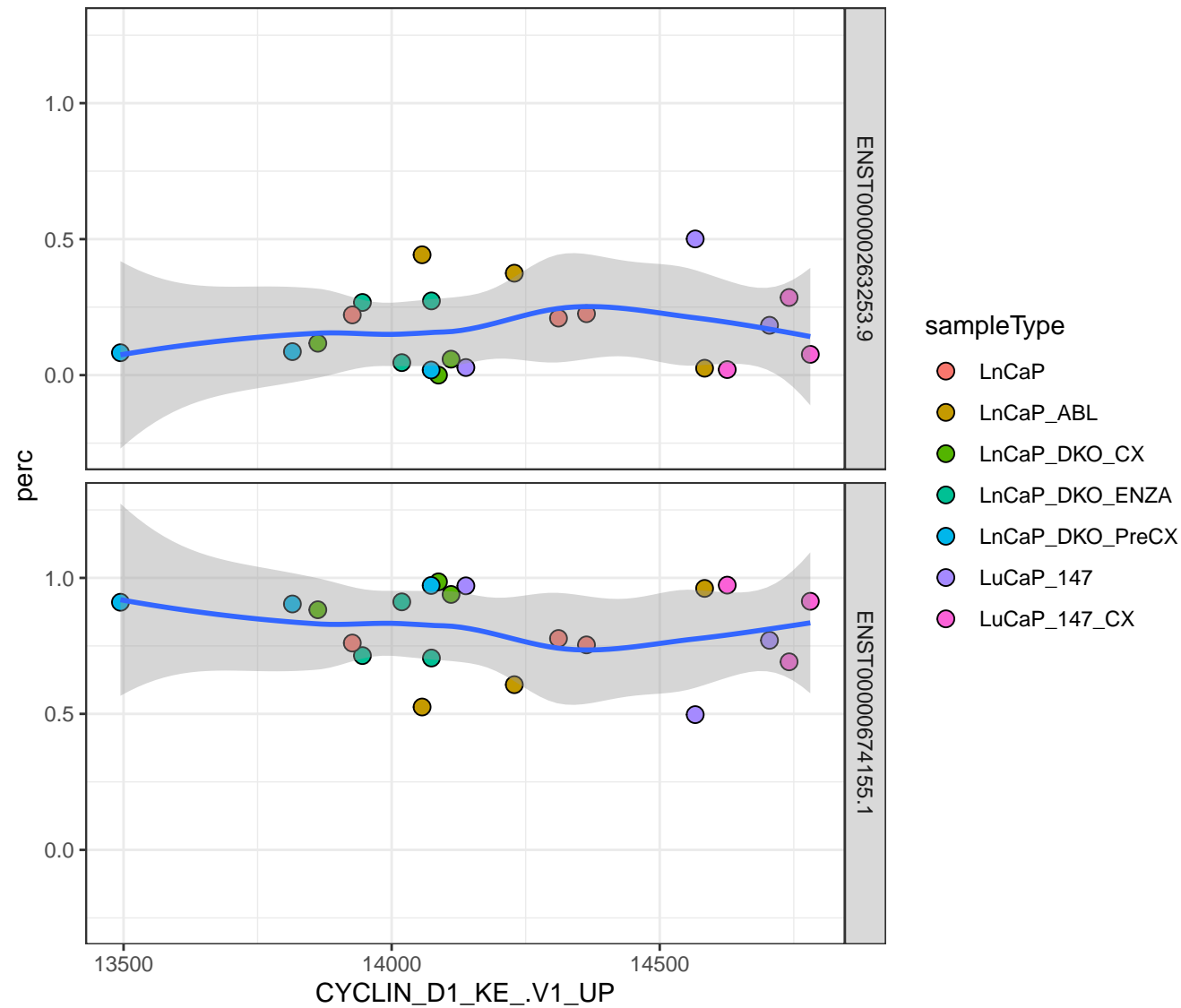
Target = CYCLIN_D1_KE_.V1_UP

ENST00000674155.1 (up) = 0.0818445675424813

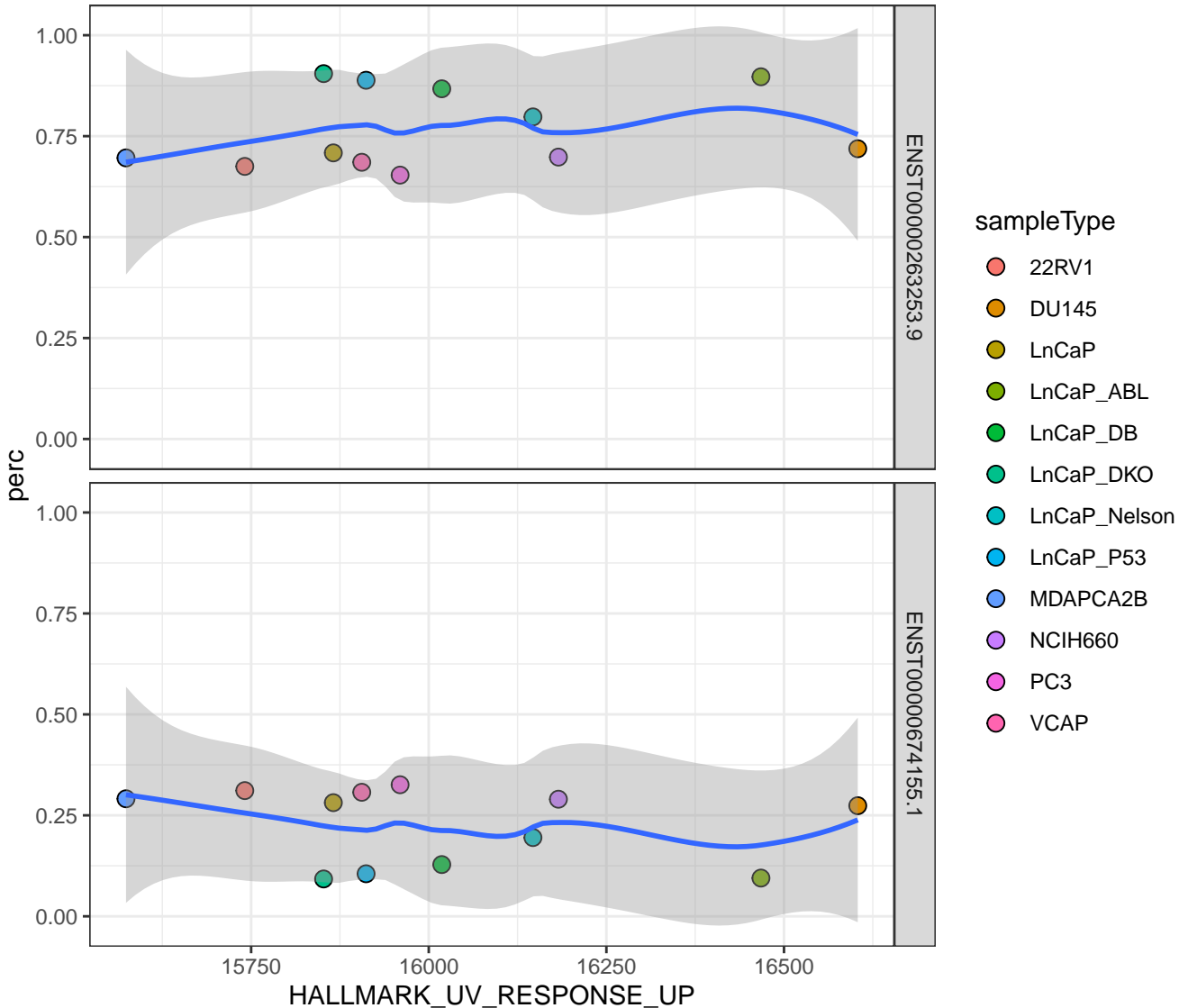
ENST00000263253.9 (down) = -0.0793444482034467



ENST00000263253.9 (down) = 0.134263146669855



CellLine
 Gene = EP300
 Target = HALLMARK_UV_RESPONSE_UP
 ENST00000674155.1 (up) = -0.21565072467497
 ENST00000263253.9 (down) = 0.218676778758411



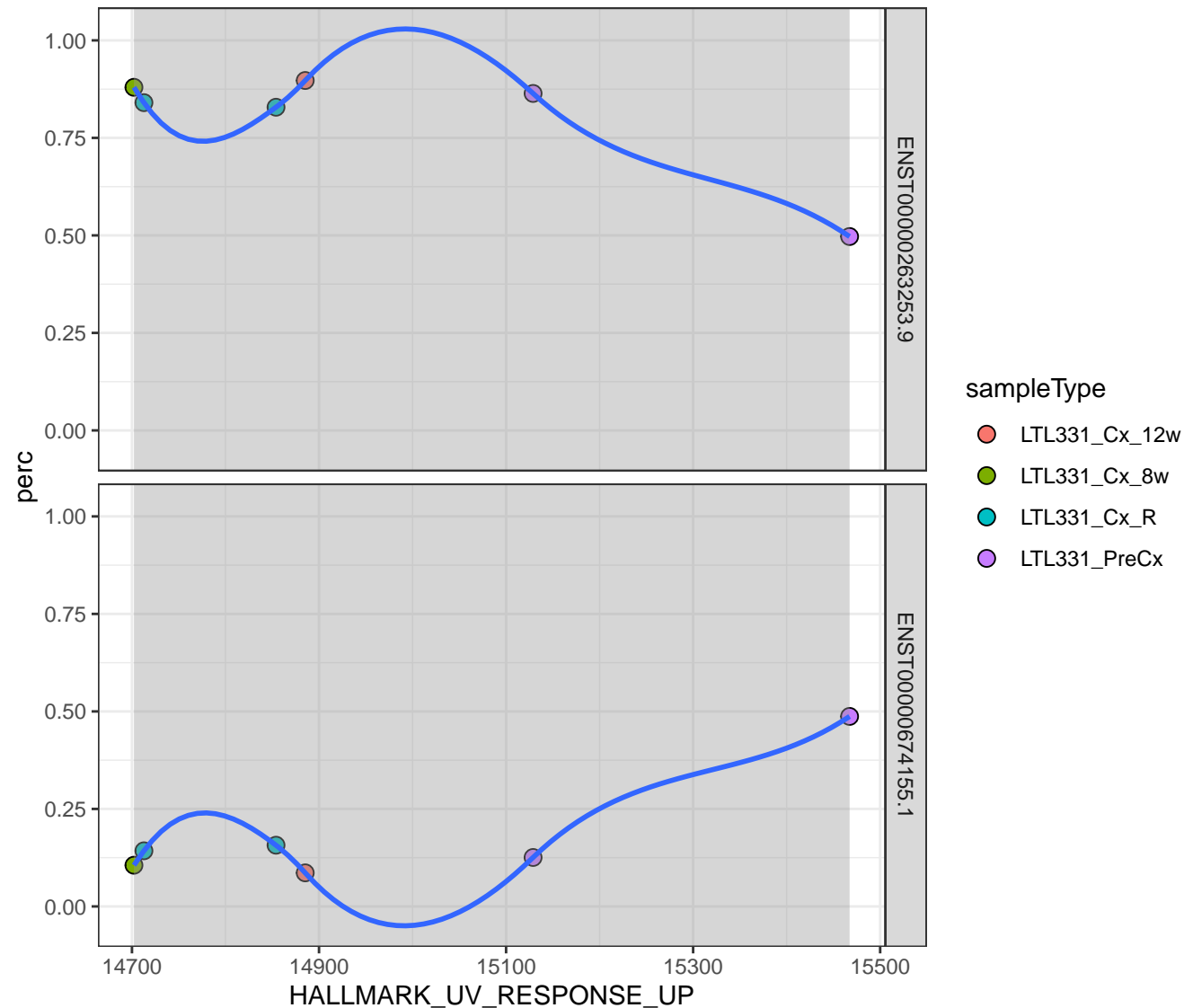
LTL_longitudinal

Gene = EP300

Target = HALLMARK_UV_RESPONSE_UP

ENST00000674155.1 (up) = 0.835040362954504

ENST00000263253.9 (down) = -0.829040803038265



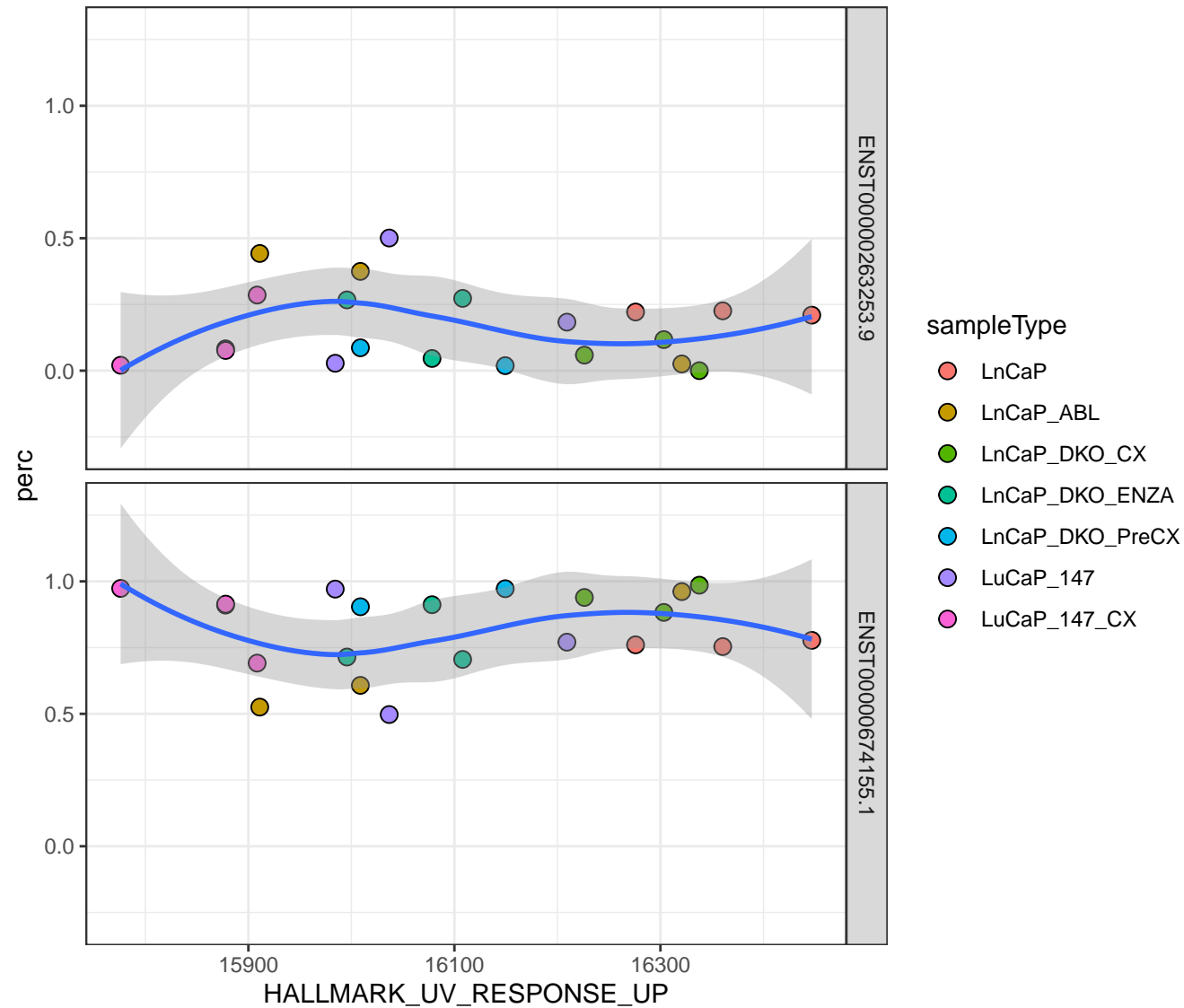
PDX_longitudinal

Gene = EP300

Target = HALLMARK_UV_RESPONSE_UP

ENST00000674155.1 (up) = 0.123855811517058

ENST00000263253.9 (down) = -0.13006816642344



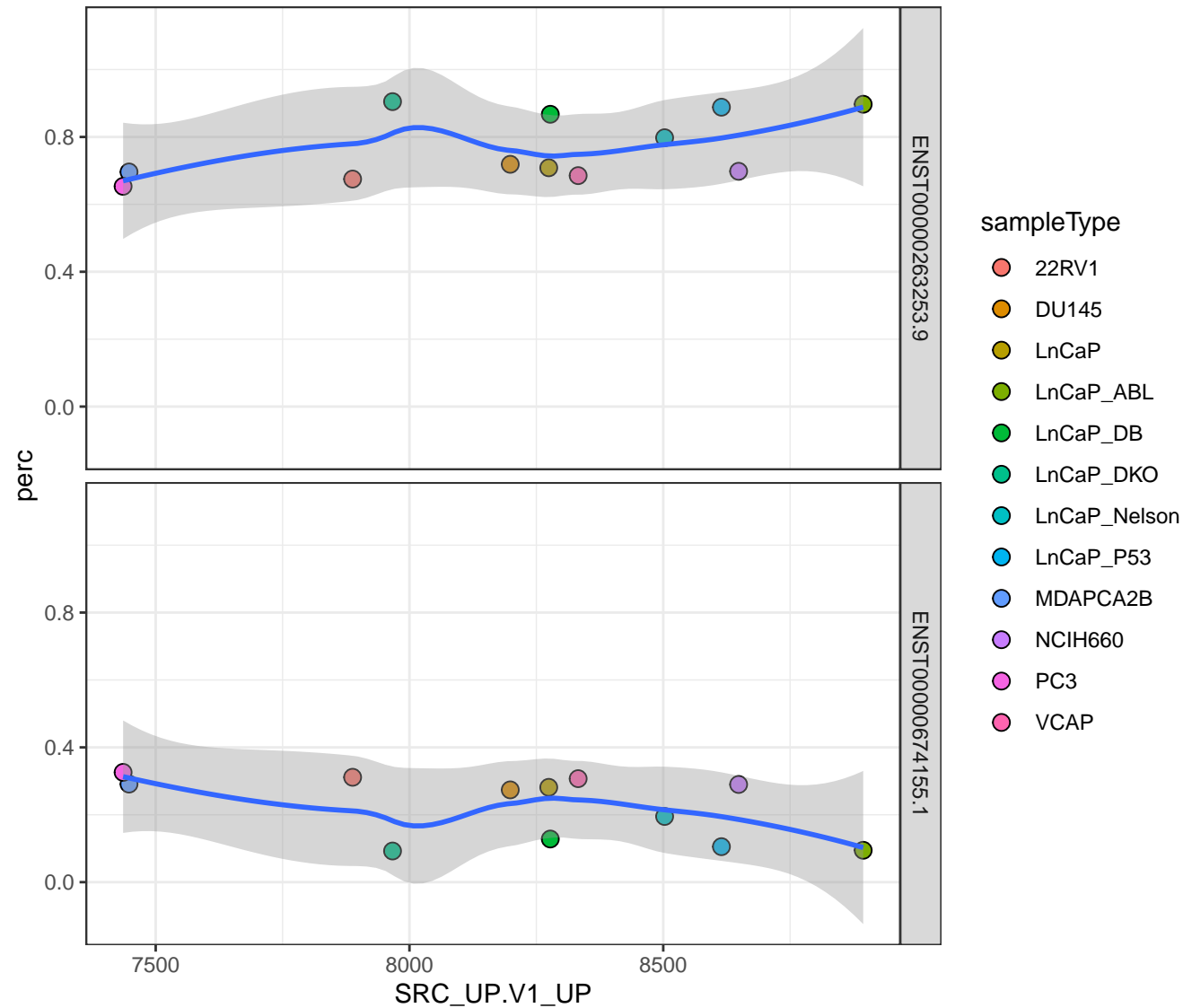
CellLine

Gene = EP300

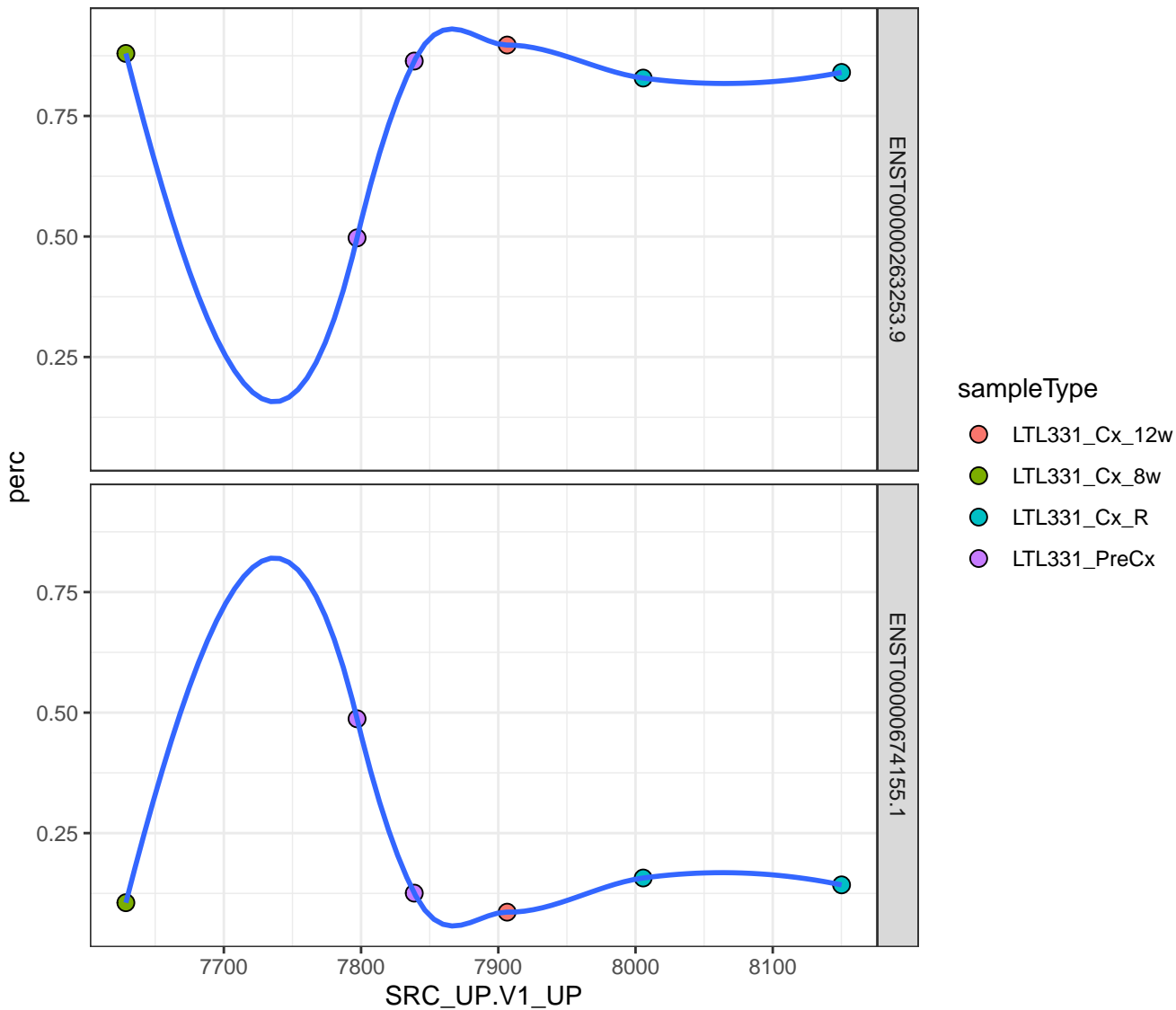
Target = SRC_UP.V1_UP

ENST00000674155.1 (up) = -0.498927588916119

ENST00000263253.9 (down) = 0.507503726558673



LTL_longitudinal
Gene = EP300
Target = SRC_UP.V1_UP
ENST00000674155.1 (up) = -0.149355585619234
ENST00000263253.9 (down) = 0.143522040847543



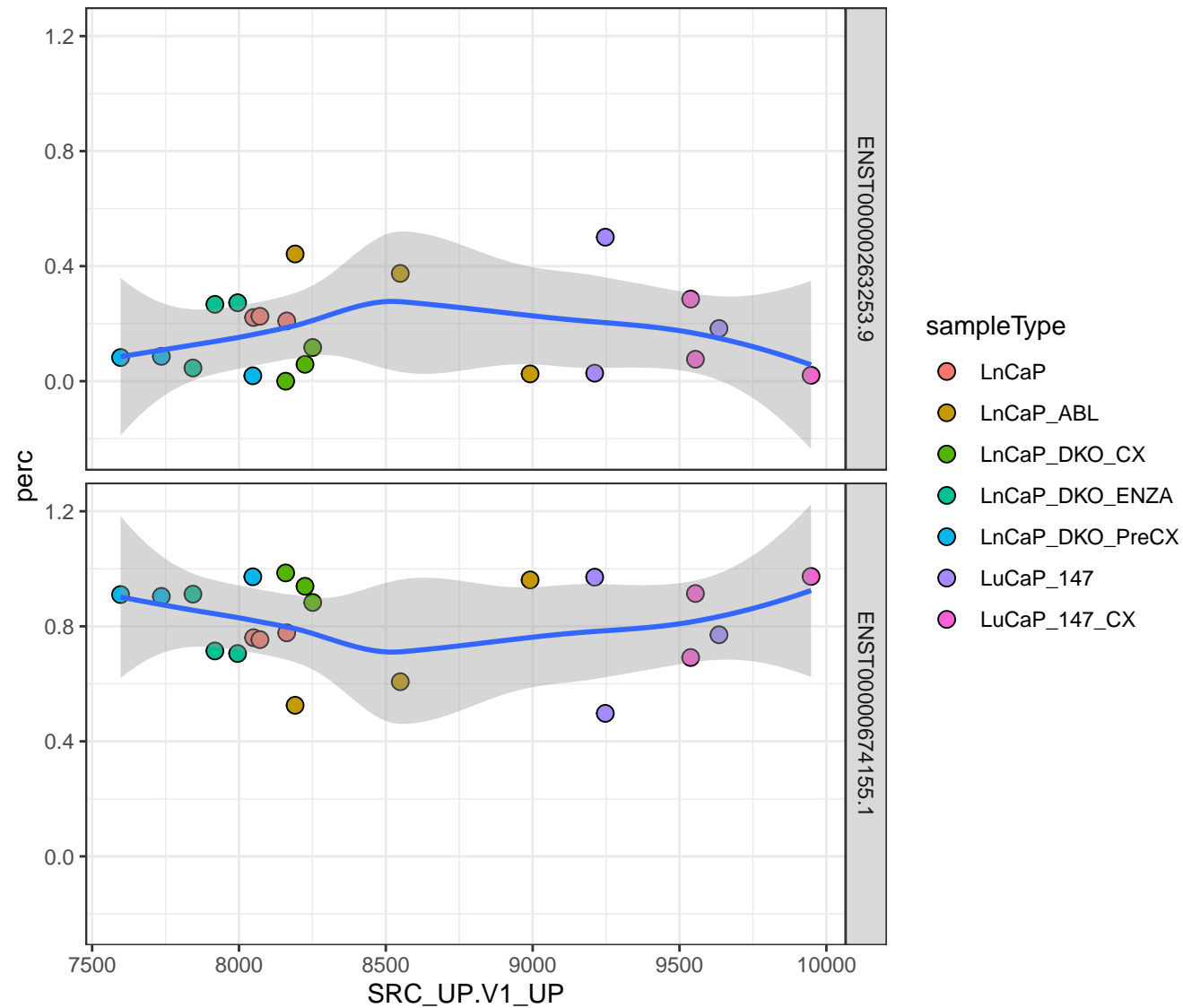
PDX_longitudinal

Gene = EP300

Target = SRC_UP.V1_UP

ENST00000674155.1 (up) = -0.0191891226870778

ENST00000263253.9 (down) = 0.022081665251375



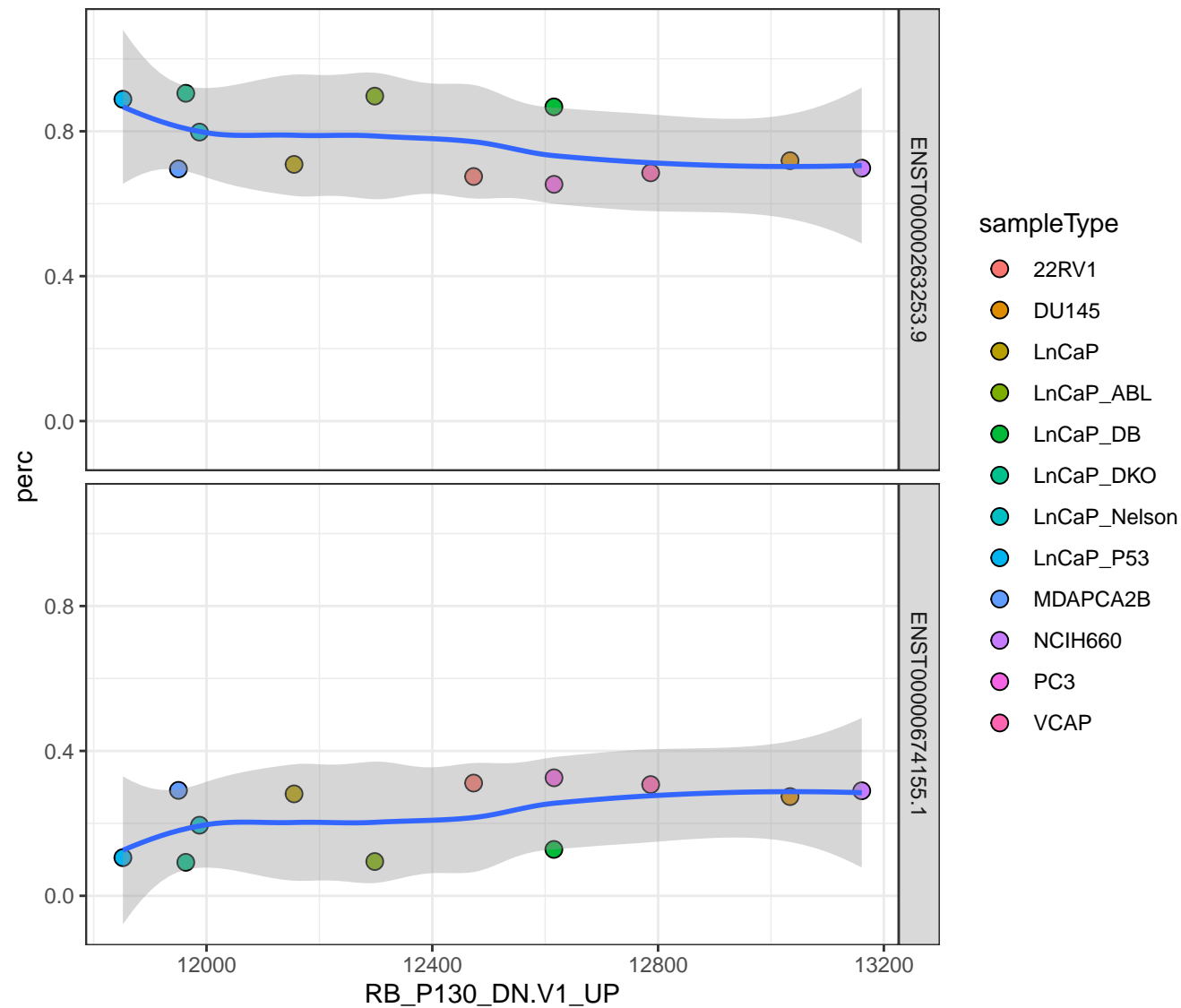
CellLine

Gene = EP300

Target = RB_P130_DN.V1_UP

ENST00000674155.1 (up) = 0.48963020531588

ENST00000263253.9 (down) = -0.48348391167937



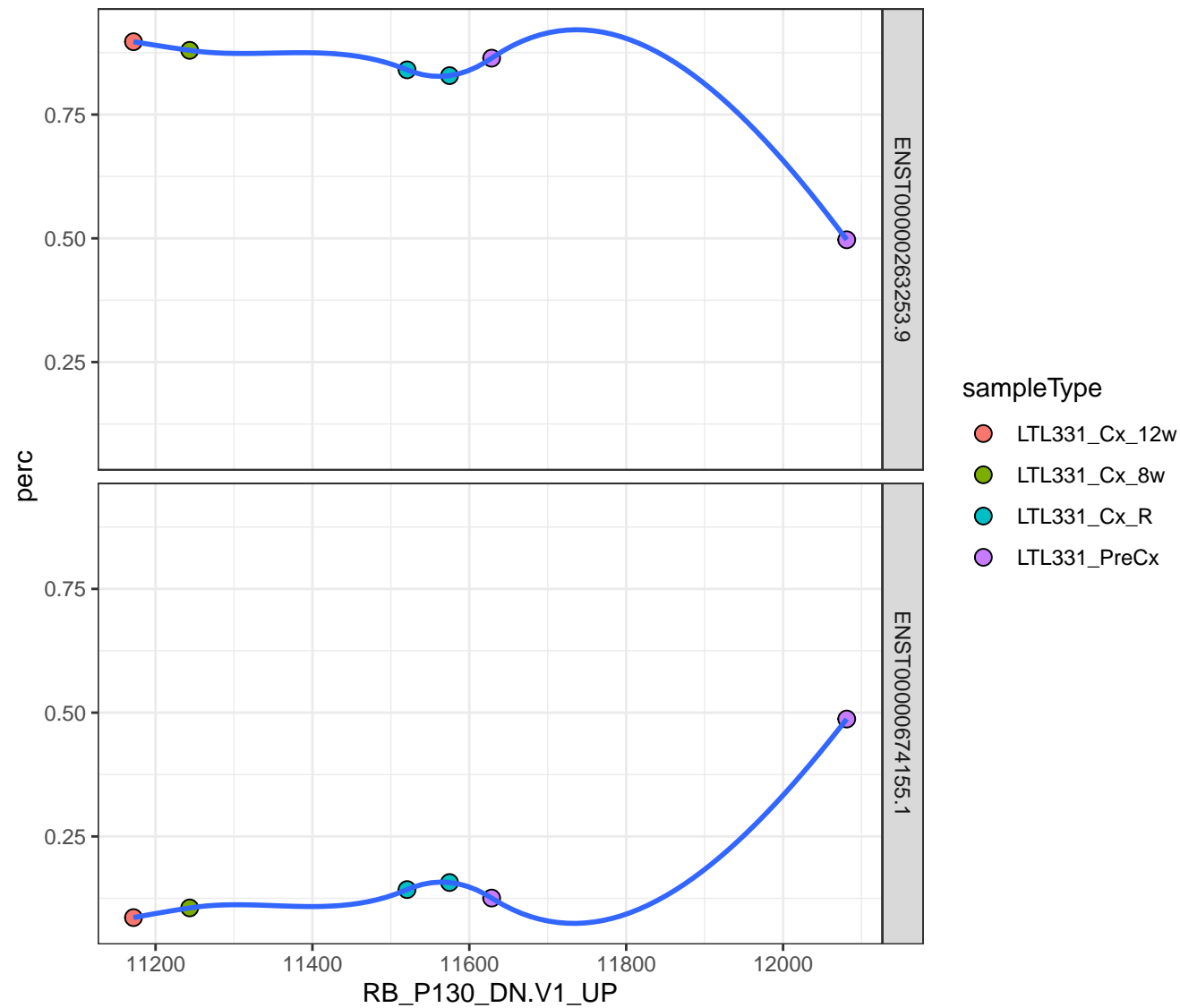
LTL_longitudinal

Gene = EP300

Target = RB_P130_DN.V1_UP

ENST00000674155.1 (up) = 0.892663392429247

ENST00000263253.9 (down) = -0.887962436491458



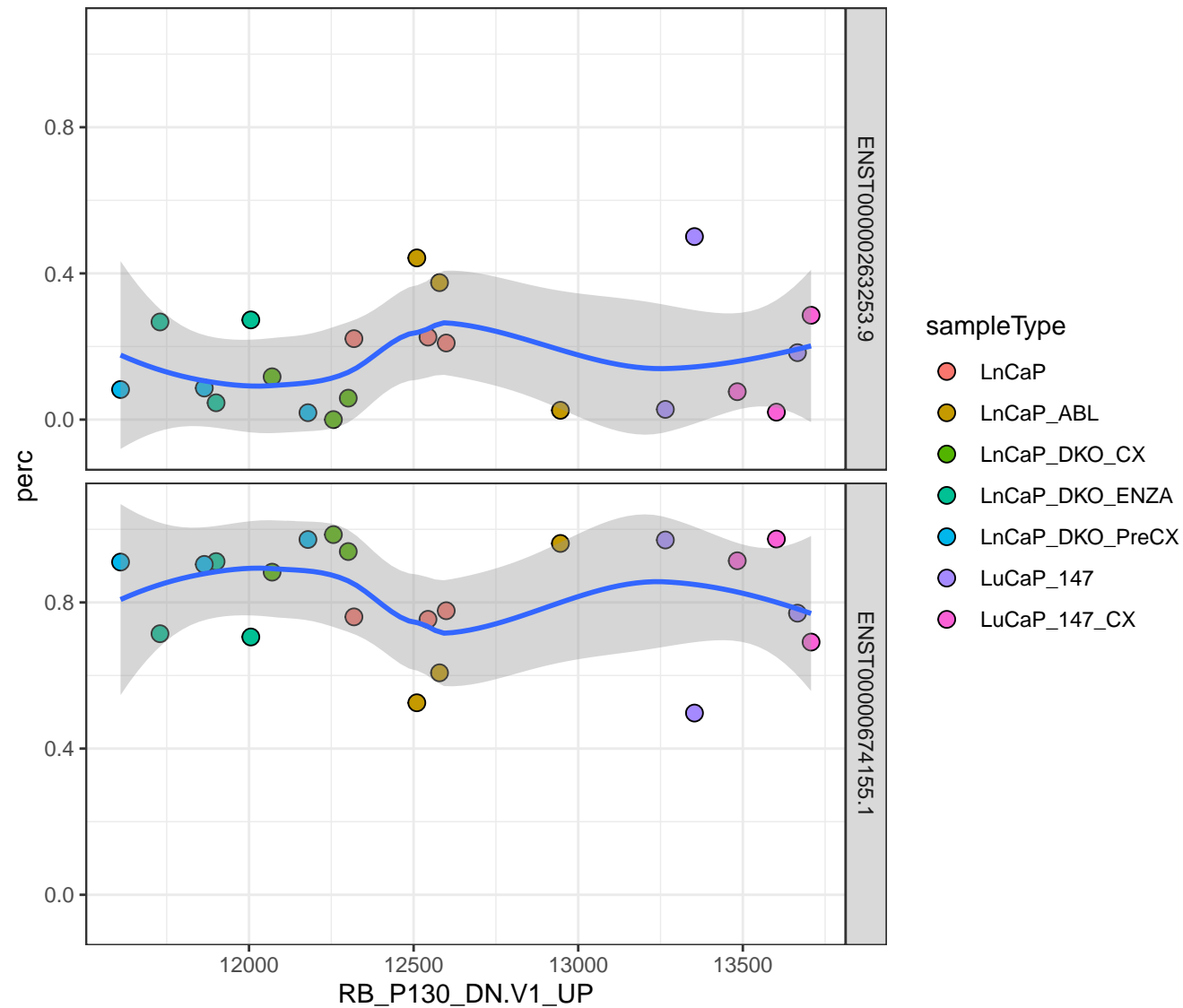
PDX_longitudinal

Gene = EP300

Target = RB_P130_DN.V1_UP

ENST00000674155.1 (up) = -0.120681041892948

ENST00000263253.9 (down) = 0.120140799375119



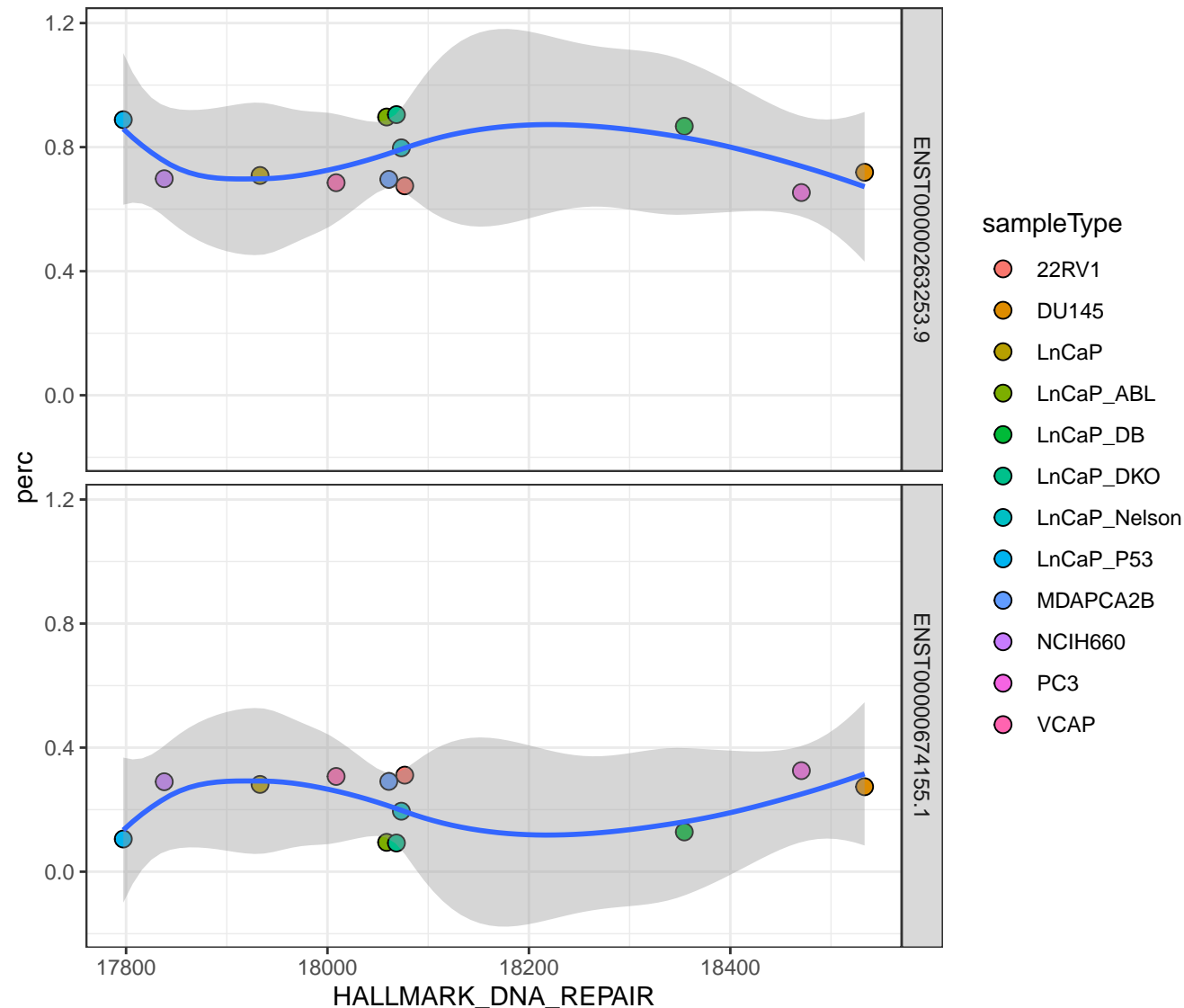
CellLine

Gene = EP300

Target = HALLMARK_DNA_REPAIR

ENST00000674155.1 (up) = 0.174650408153299

ENST00000263253.9 (down) = -0.178136080281216



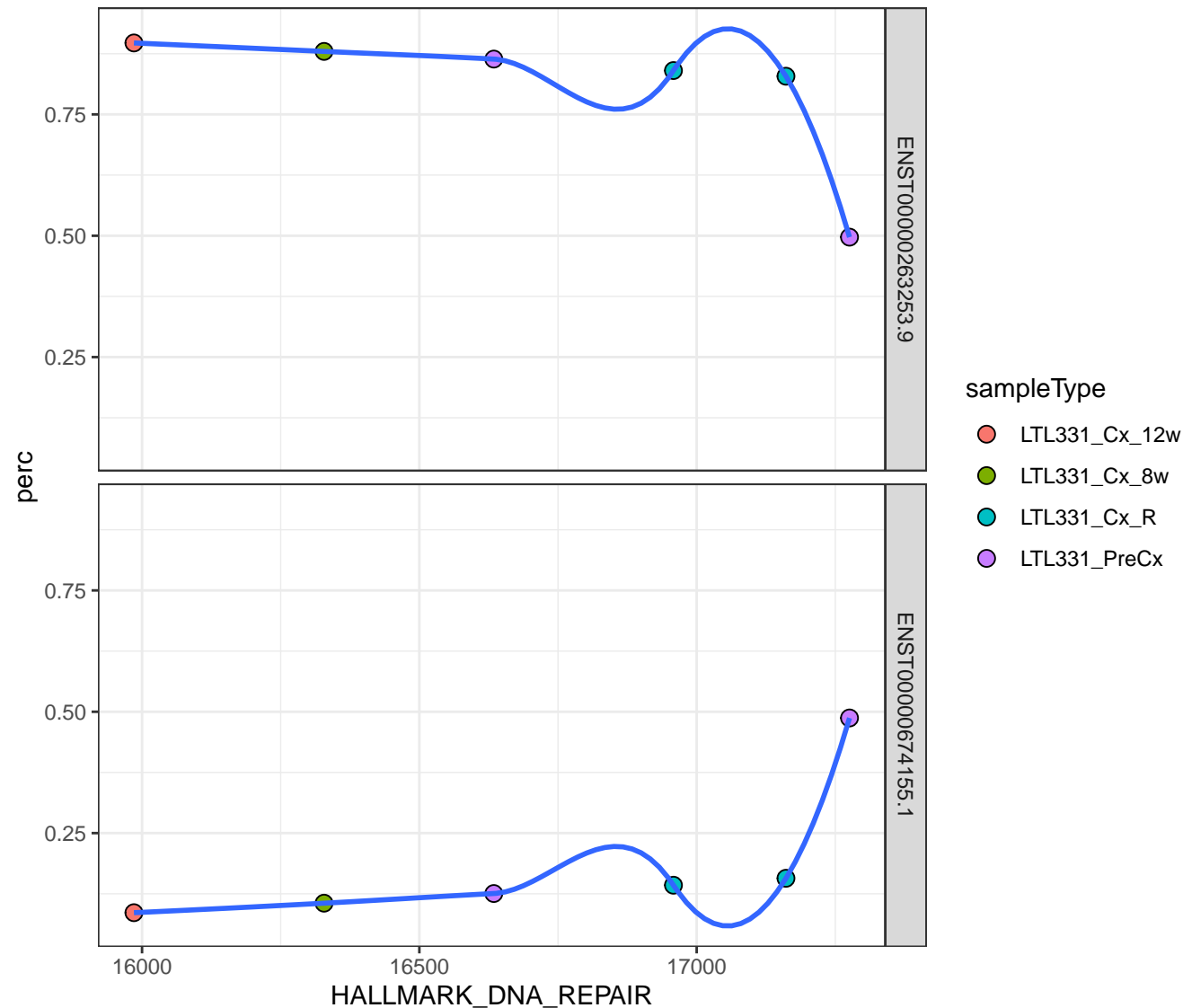
LTL_longitudinal

Gene = EP300

Target = HALLMARK_DNA_REPAIR

ENST00000674155.1 (up) = 0.672416131367312

ENST00000263253.9 (down) = -0.67038808028203



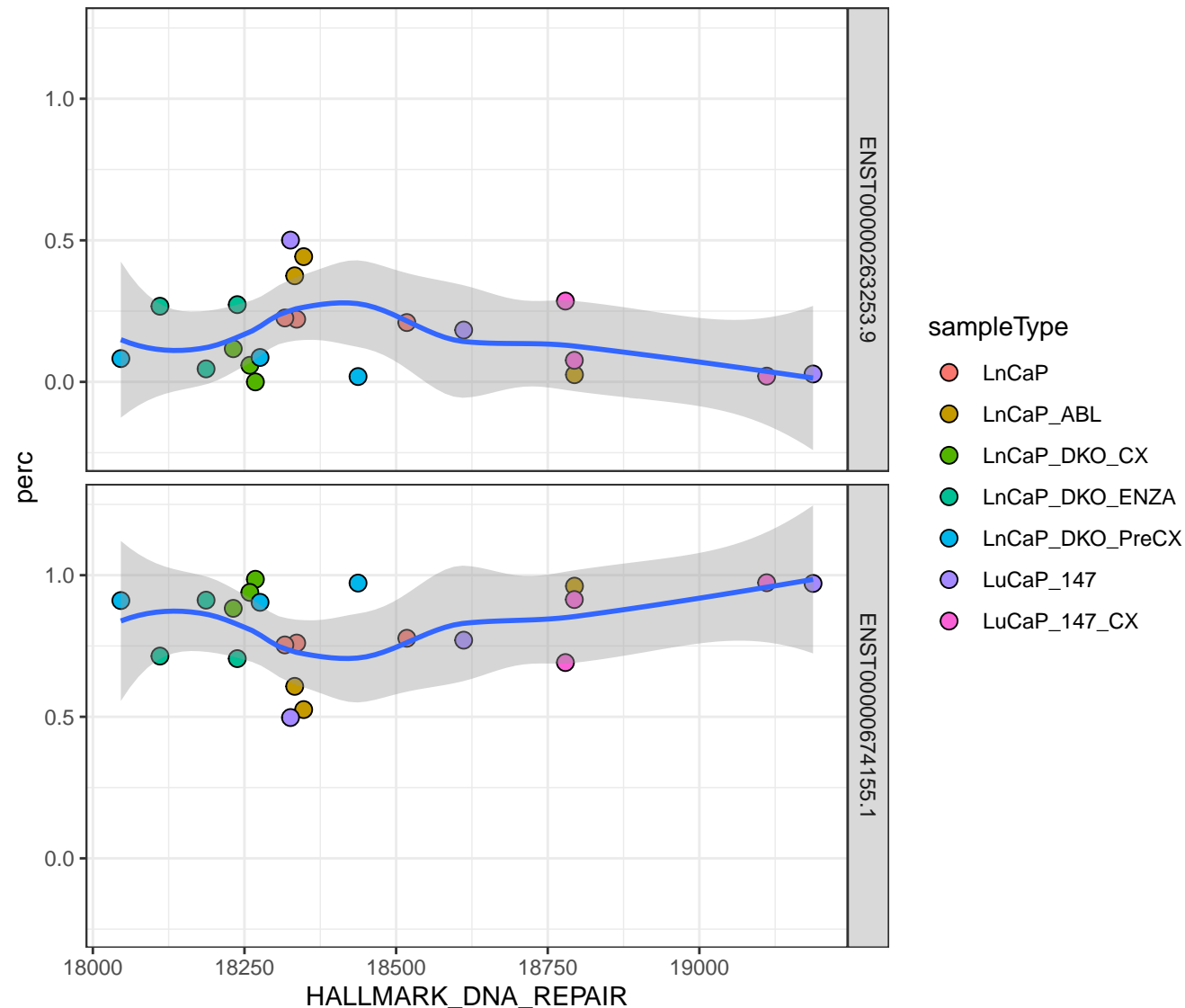
PDX_longitudinal

Gene = EP300

Target = HALLMARK_DNA_REPAIR

ENST00000674155.1 (up) = 0.282397280803224

ENST00000263253.9 (down) = -0.275932348499257



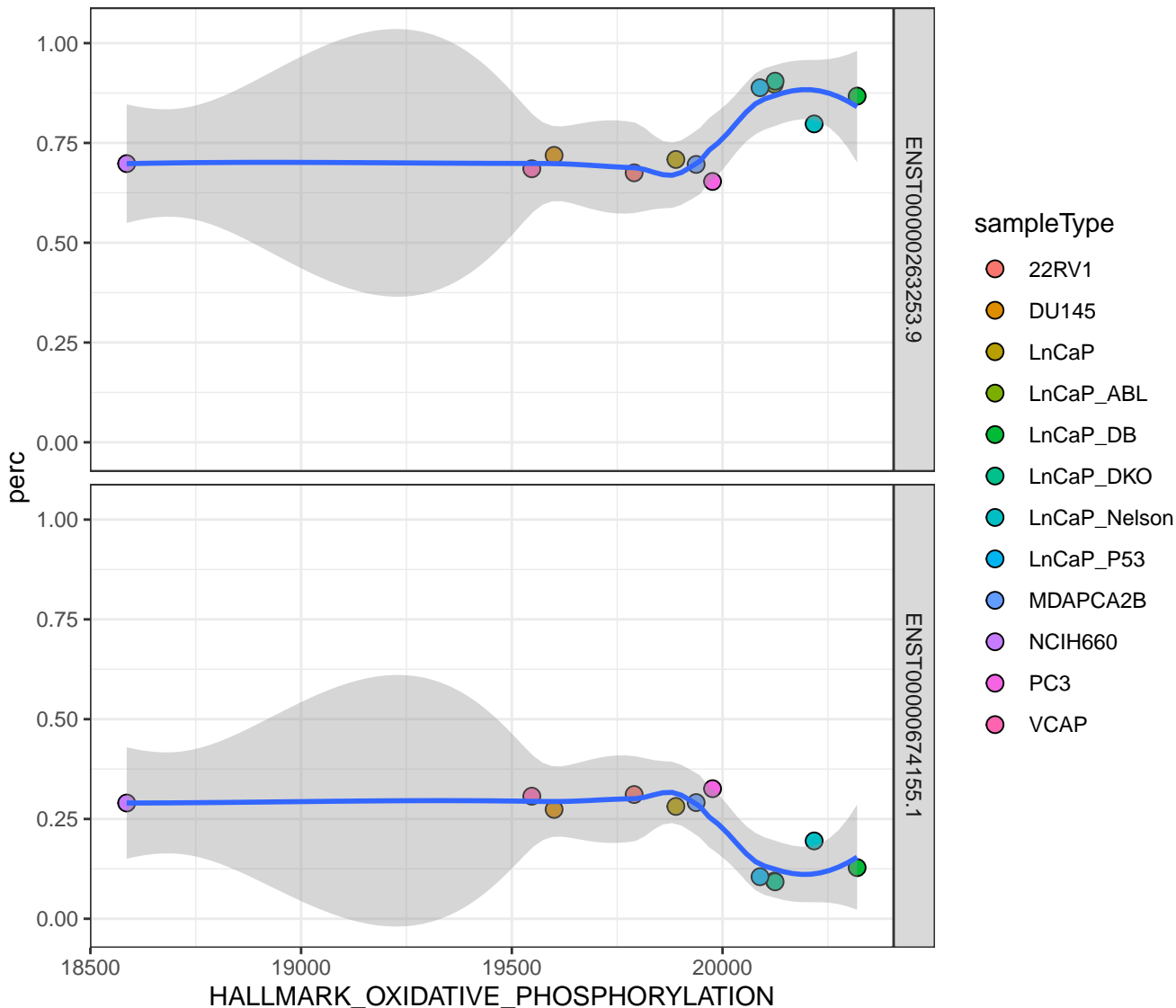
CellLine

Gene = EP300

Target = HALLMARK_OXIDATIVE_PHOSPHORYLATION

ENST00000674155.1 (up) = -0.544979120737338

ENST00000263253.9 (down) = 0.538379010695262



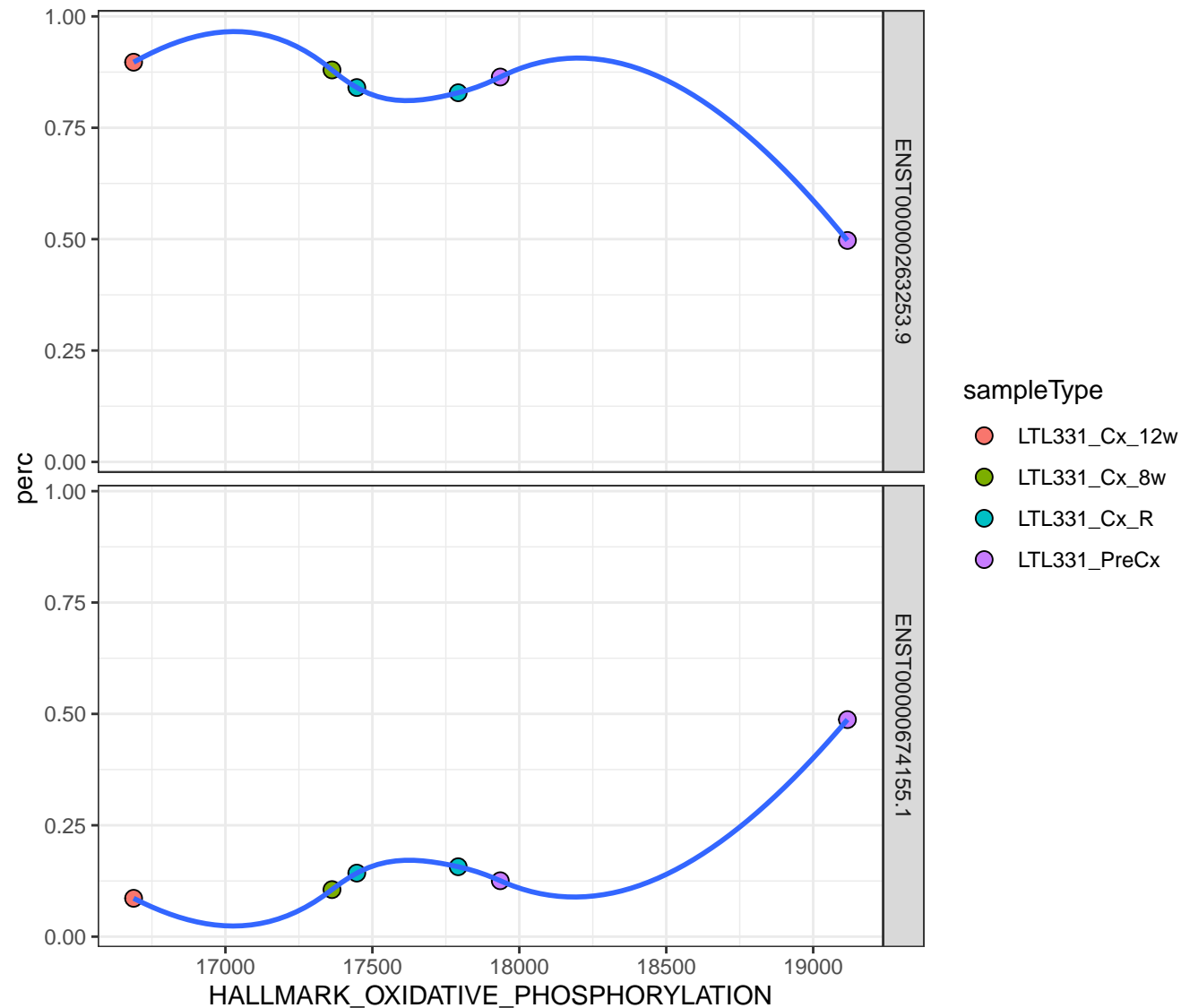
LTL_longitudinal

Gene = EP300

Target = HALLMARK_OXIDATIVE_PHOSPHORYLATION

ENST00000674155.1 (up) = 0.901566034359569

ENST00000263253.9 (down) = -0.895676953071641



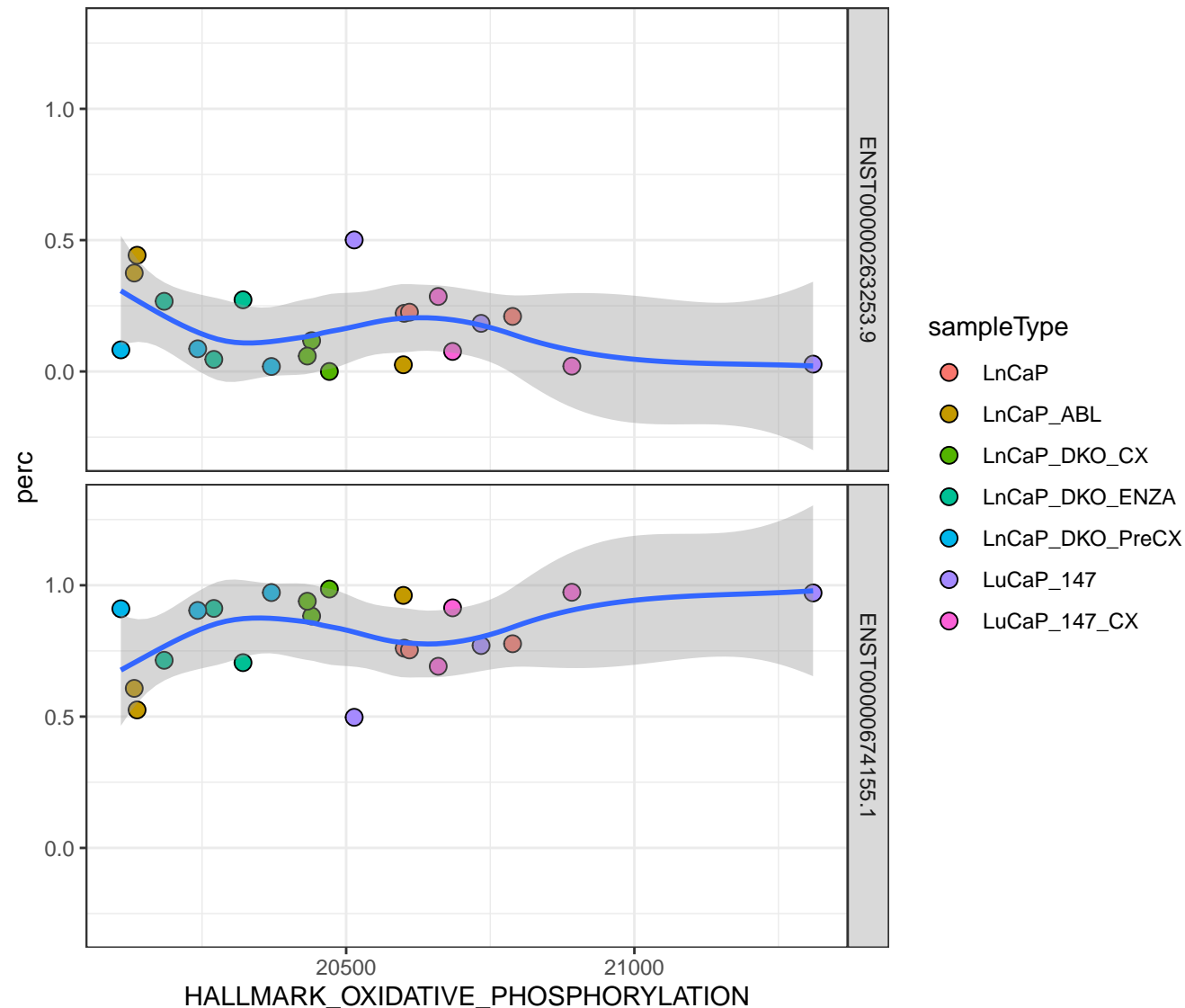
PDX_longitudinal

Gene = EP300

Target = HALLMARK_OXIDATIVE_PHOSPHORYLATION

ENST00000674155.1 (up) = 0.309973164905225

ENST00000263253.9 (down) = -0.299110629037274



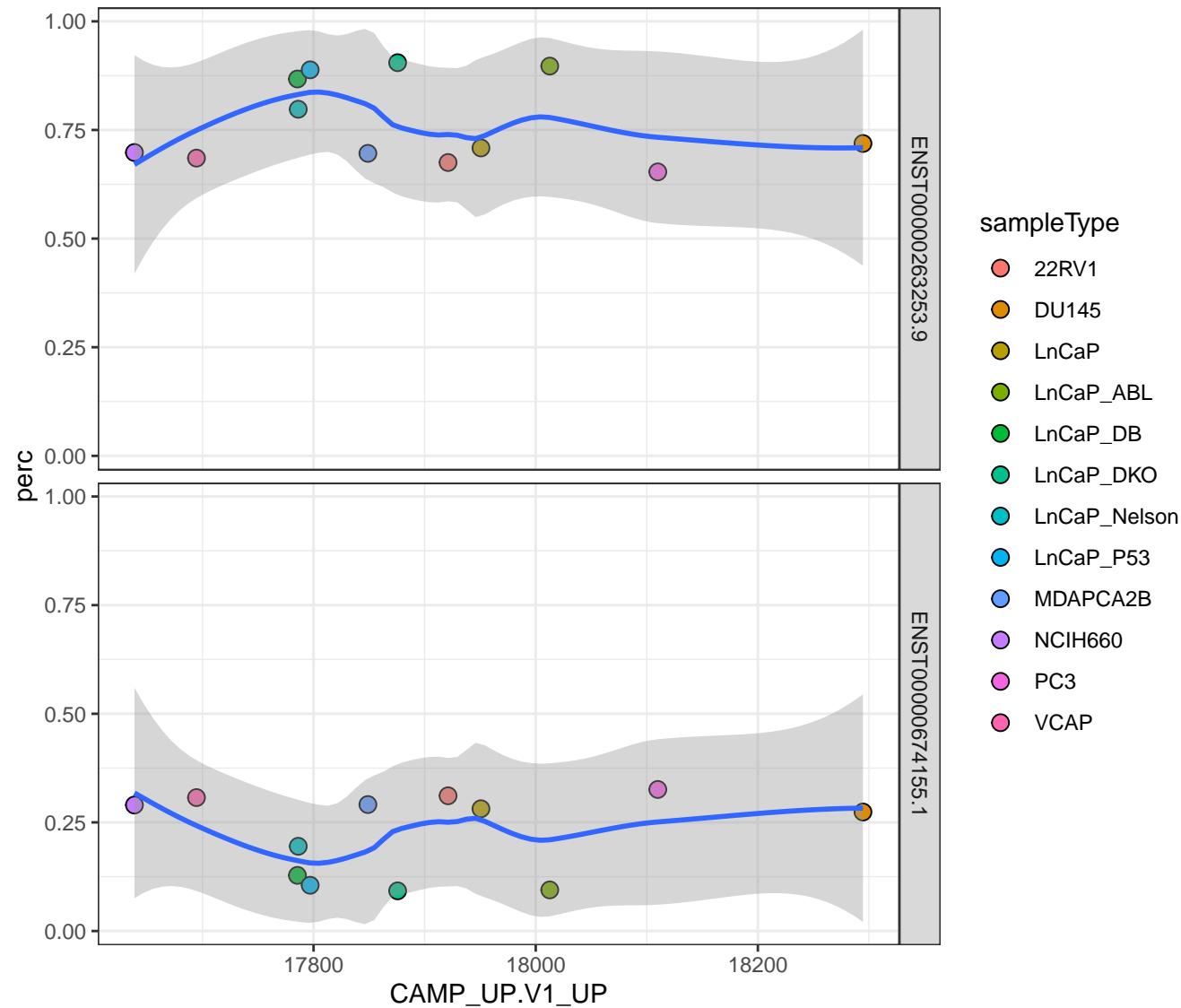
CellLine

Gene = EP300

Target = CAMP_UP.V1_UP

ENST00000674155.1 (up) = 0.12328892753789

ENST00000263253.9 (down) = -0.130653907194808



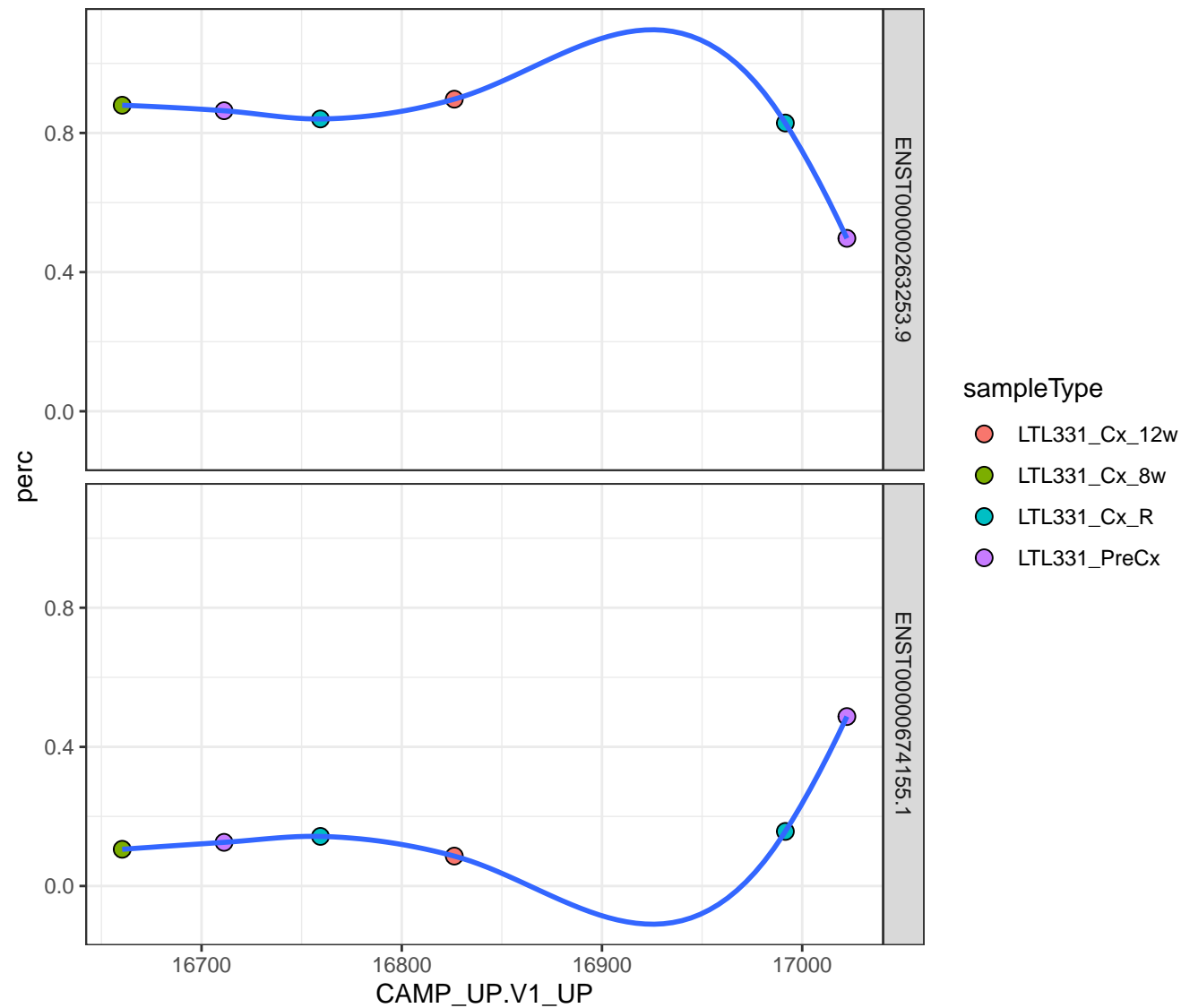
LTL_longitudinal

Gene = EP300

Target = CAMP_UP.V1_UP

ENST00000674155.1 (up) = 0.690225577689066

ENST00000263253.9 (down) = -0.692825737059426



PDX_longitudinal

Gene = EP300

Target = CAMP_UP.V1_UP

ENST00000674155.1 (up) = 0.177661588583952

ENST00000263253.9 (down) = -0.18854284258903

