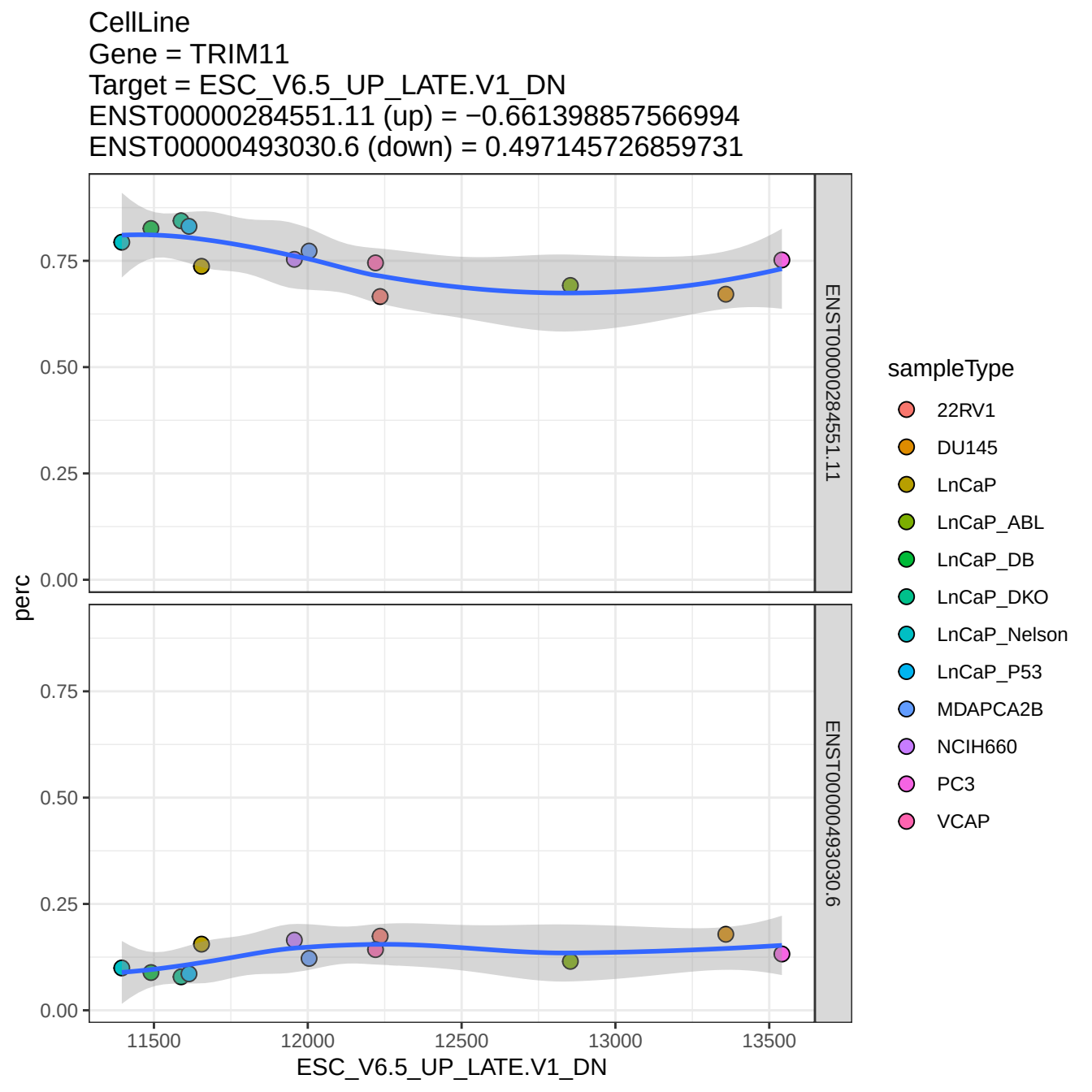
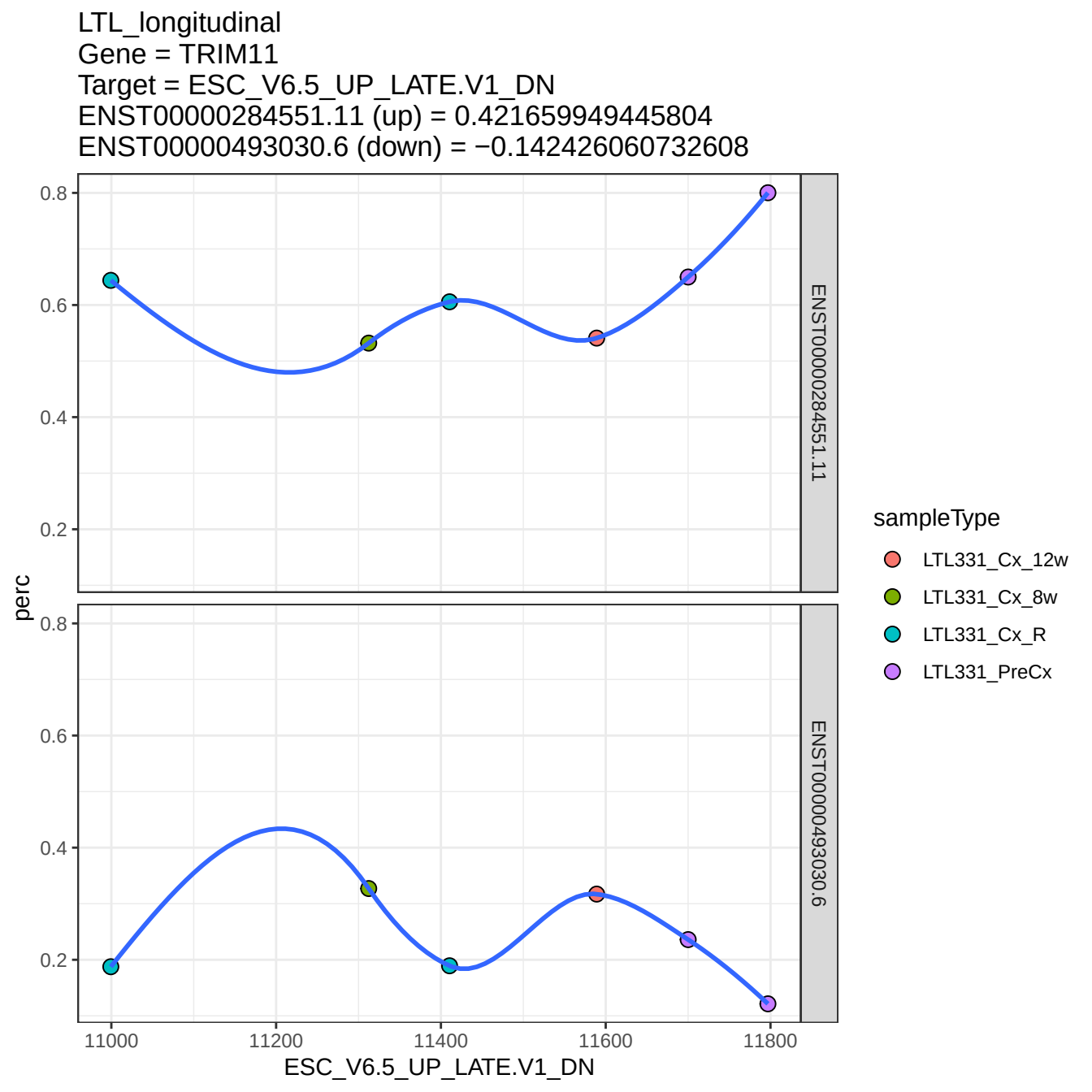
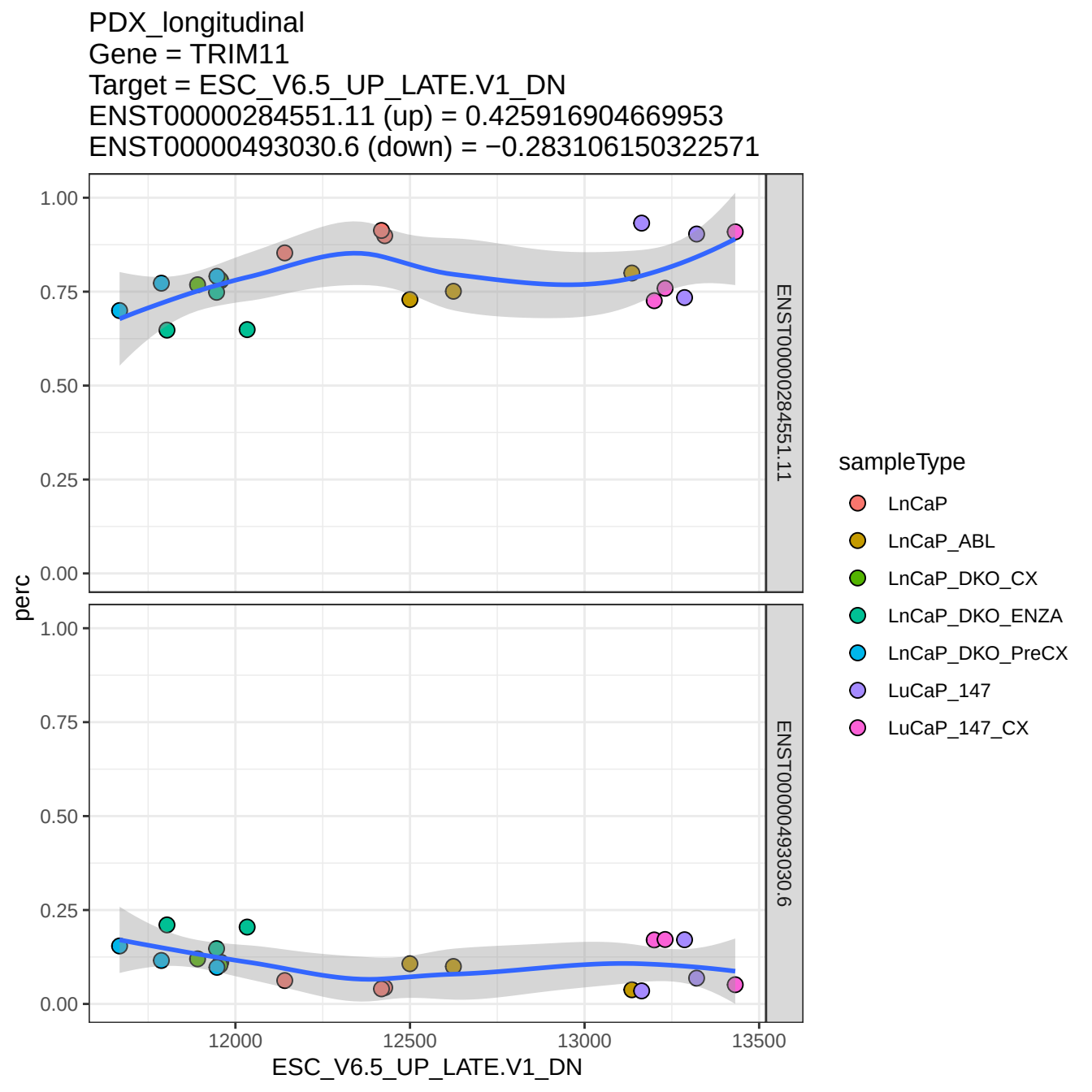
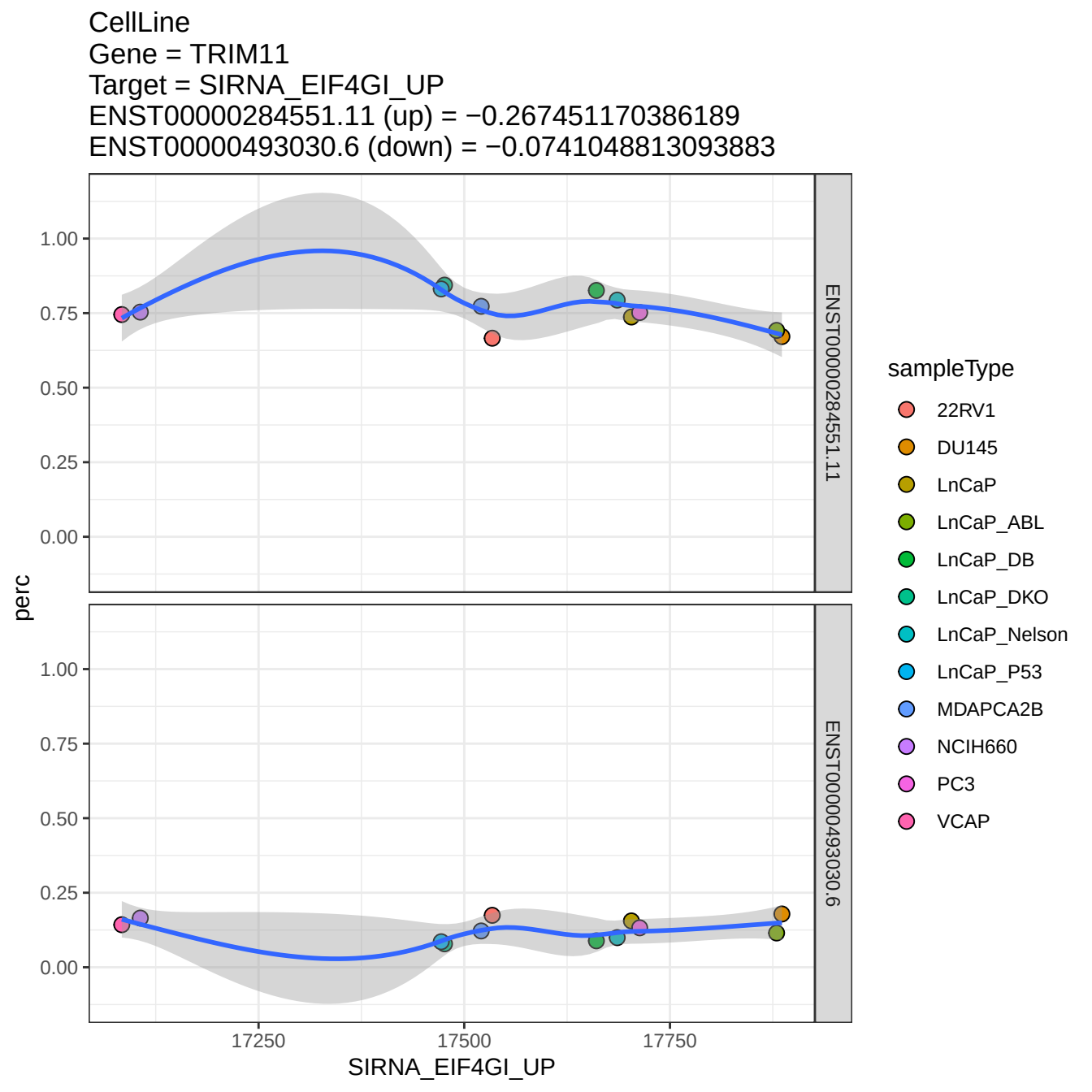
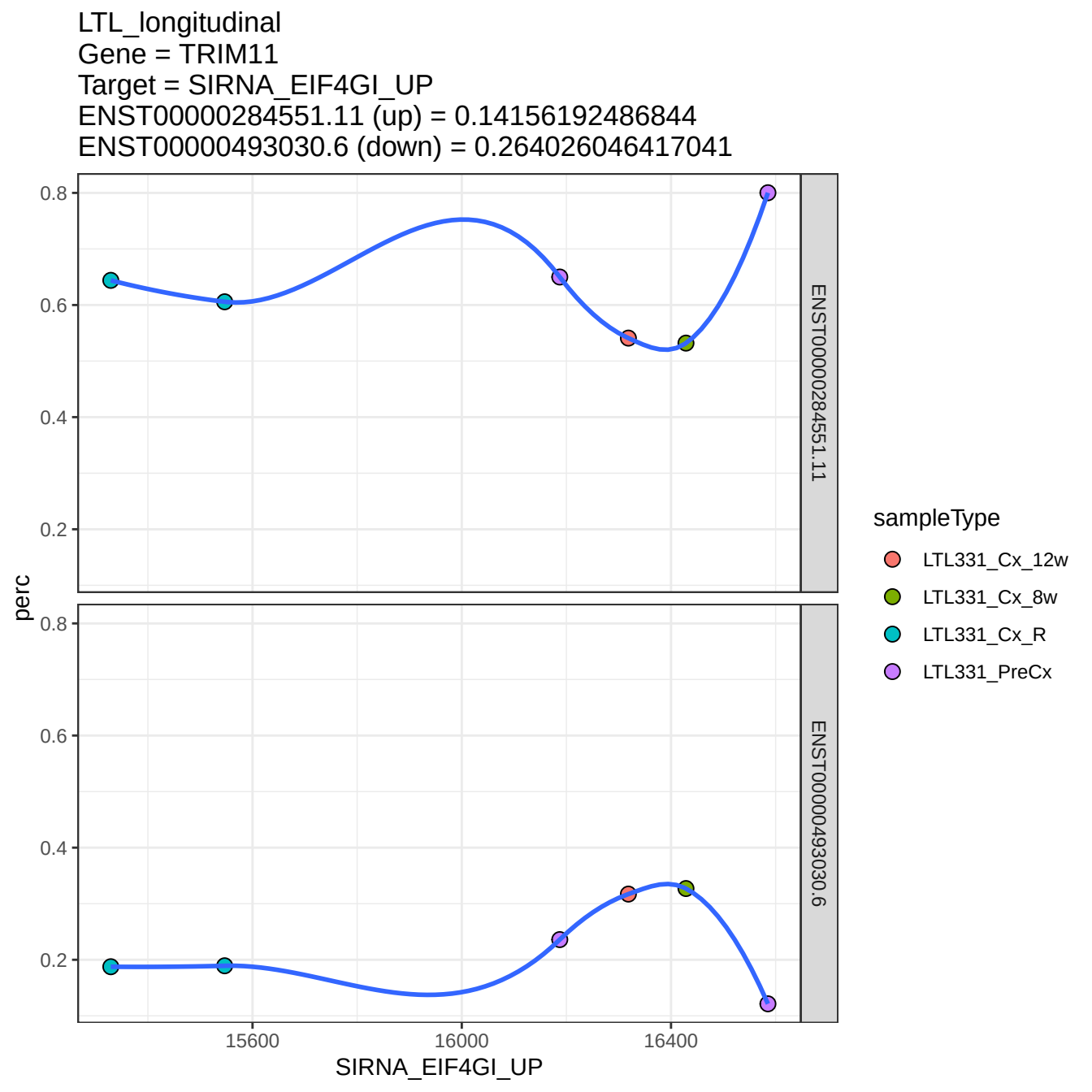
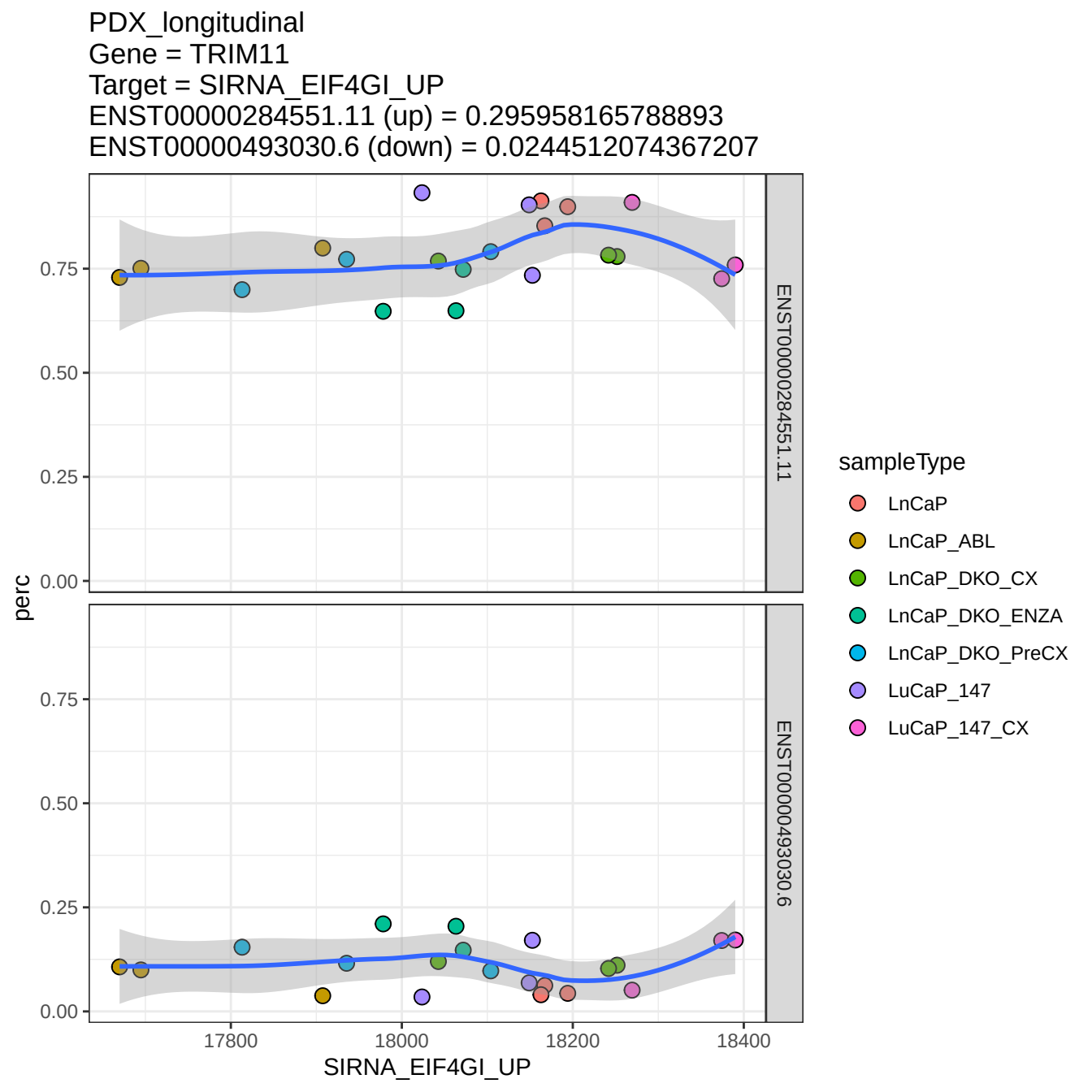

CellLine
Gene = TRIM11
Target = ESC_V6.5_UP_LATE.V1_DN
ENST00000284551.11 (up) = −0.661398857566994
ENST00000493030.6 (down) = 0.497145726859731
perc
0.00
0.25
0.50
0.75
ENST00000284551.11
ENST00000493030.6
11500
12000
12500
13000
13500
ESC_V6.5_UP_LATE.V1_DN
sampleType
22RV1
DU145
LnCaP
LnCaP_ABL
LnCaP_DB
LnCaP_DKO
LnCaP_Nelson
LnCaP_P53
MDAPCA2B
NCIH660
PC3
VCAP

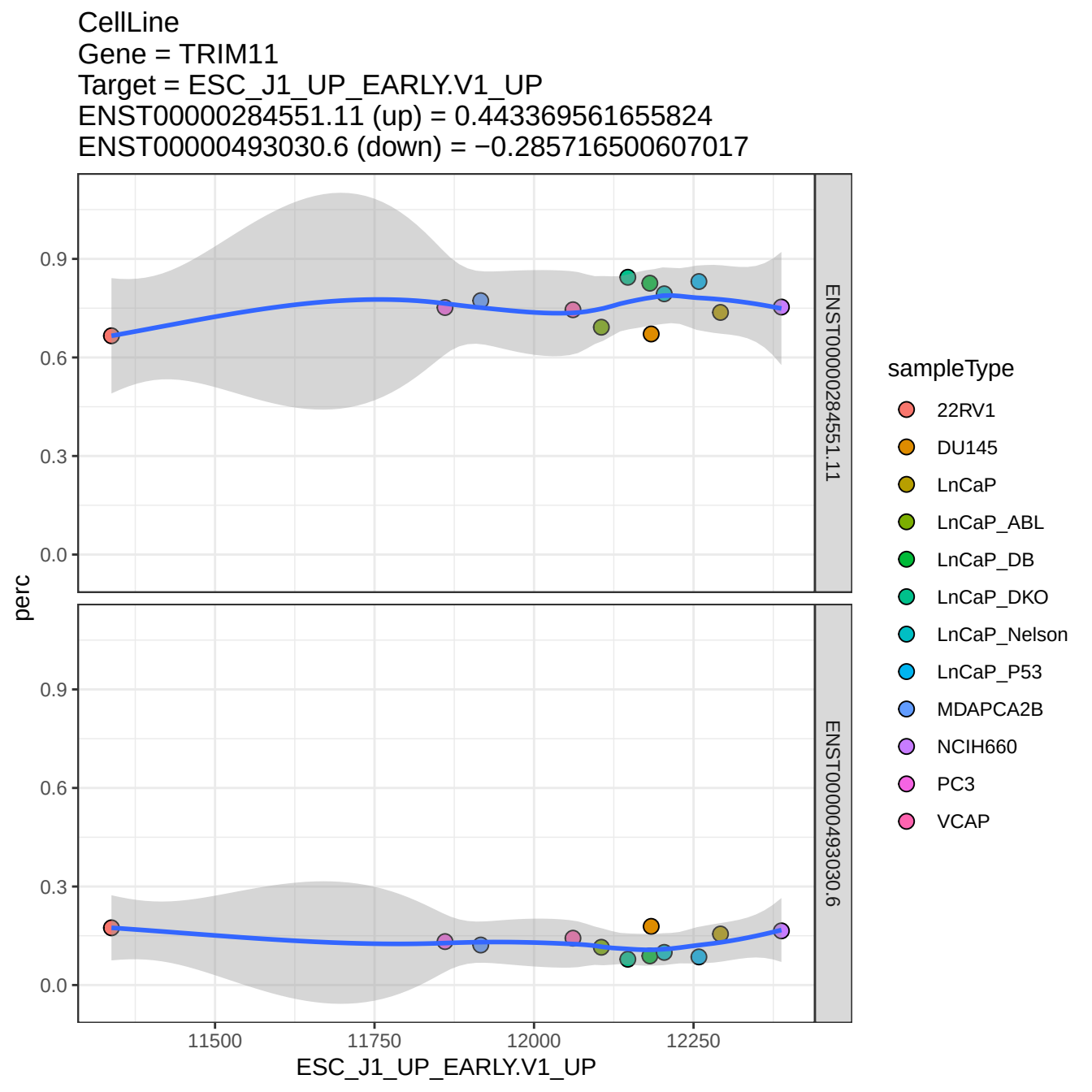

LTL_longitudinal
Gene = TRIM11
Target = ESC_V6.5_UP_LATE.V1_DN
ENST00000284551.11 (up) = 0.421659949445804
ENST00000493030.6 (down) = −0.142426060732608
perc
0.8
0.6
0.4
0.2
ENST00000284551.11
ENST00000493030.6
11000
11200
11400
11600
11800
ESC_V6.5_UP_LATE.V1_DN
sampleType
LTL331_Cx_12w
LTL331_Cx_8w
LTL331_Cx_R
LTL331_PreCx

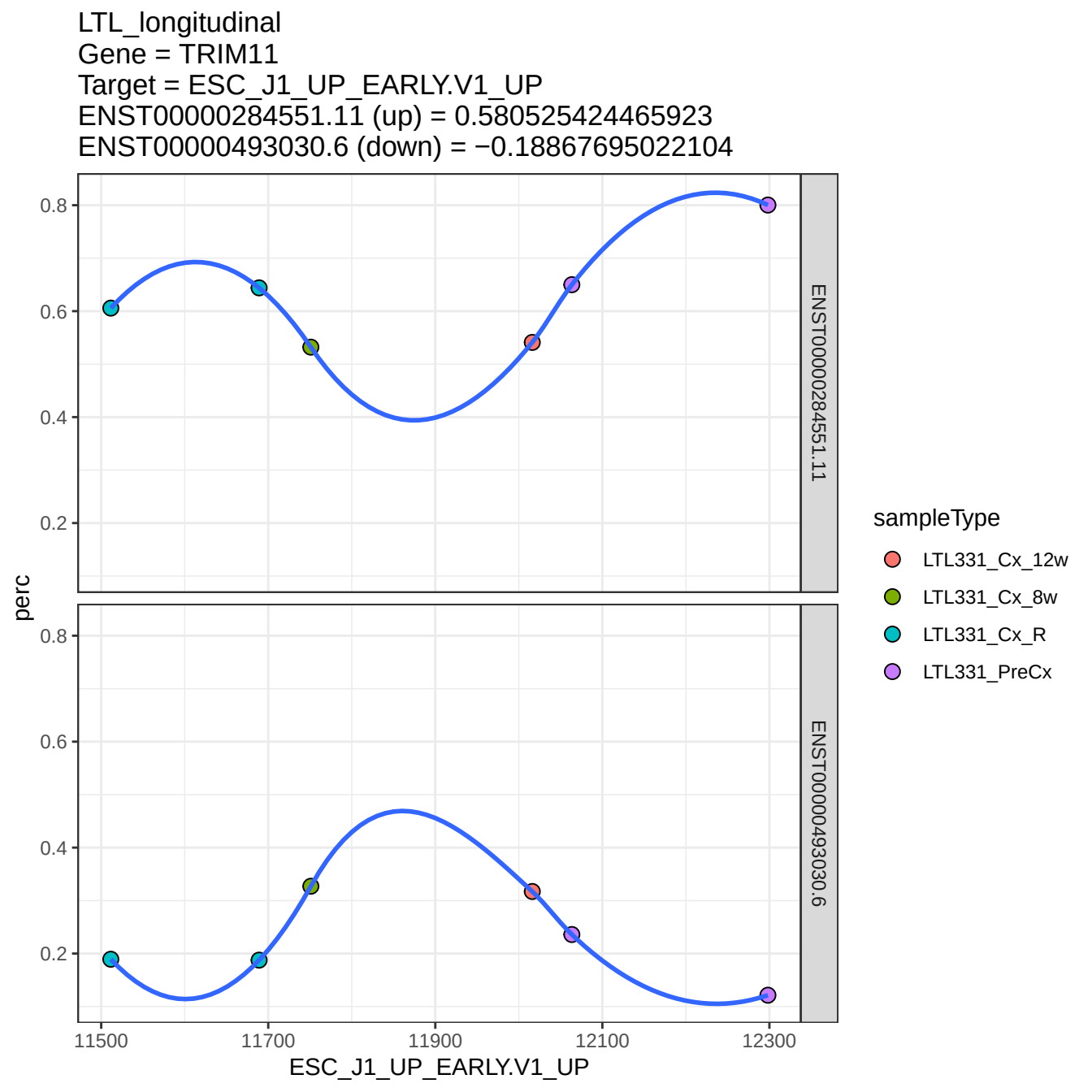

PDX_longitudinal
Gene = TRIM11
Target = ESC_V6.5_UP_LATE.V1_DN
ENST00000284551.11 (up) = 0.425916904669953
ENST00000493030.6 (down) = −0.283106150322571
perc
1.00
0.75
0.50
0.25
0.00
ENST00000284551.11
ENST00000493030.6
12000
12500
13000
13500
ESC_V6.5_UP_LATE.V1_DN
sampleType
LnCaP
LnCaP_ABL
LnCaP_DKO_CX
LnCaP_DKO_ENZA
LnCaP_DKO_PreCX
LuCaP_147
LuCaP_147_CX

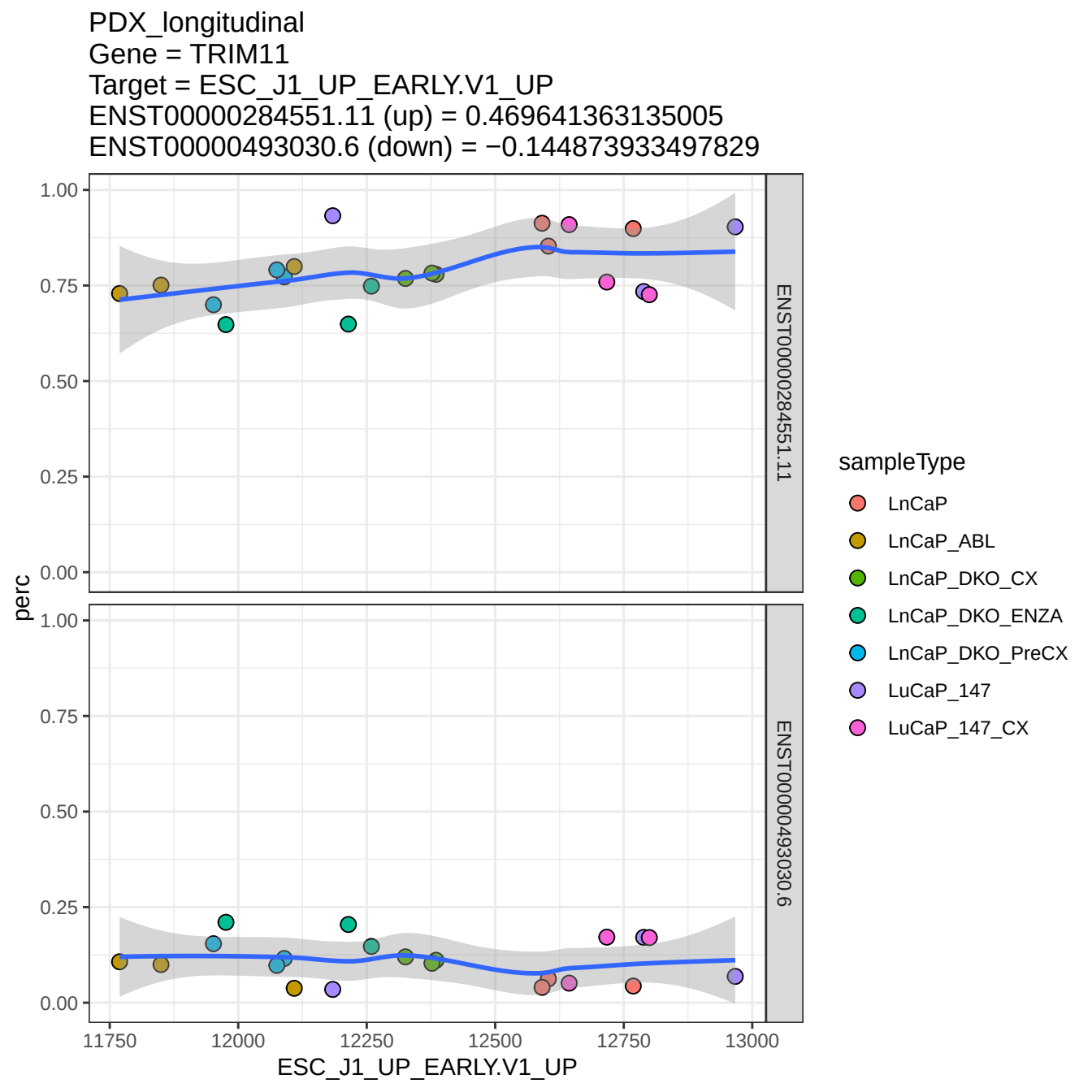

CellLine
Gene = TRIM11
Target = SIRNA_EIF4GI_UP
ENST00000284551.11 (up) = −0.267451170386189
ENST00000493030.6 (down) = −0.0741048813093883
perc
1.00
0.75
0.50
0.25
0.00
ENST00000284551.11
ENST00000493030.6
17250
17500
17750
SIRNA_EIF4GI_UP
sampleType
22RV1
DU145
LnCaP
LnCaP_ABL
LnCaP_DB
LnCaP_DKO
LnCaP_Nelson
LnCaP_P53
MDAPCA2B
NCIH660
PC3
VCAP

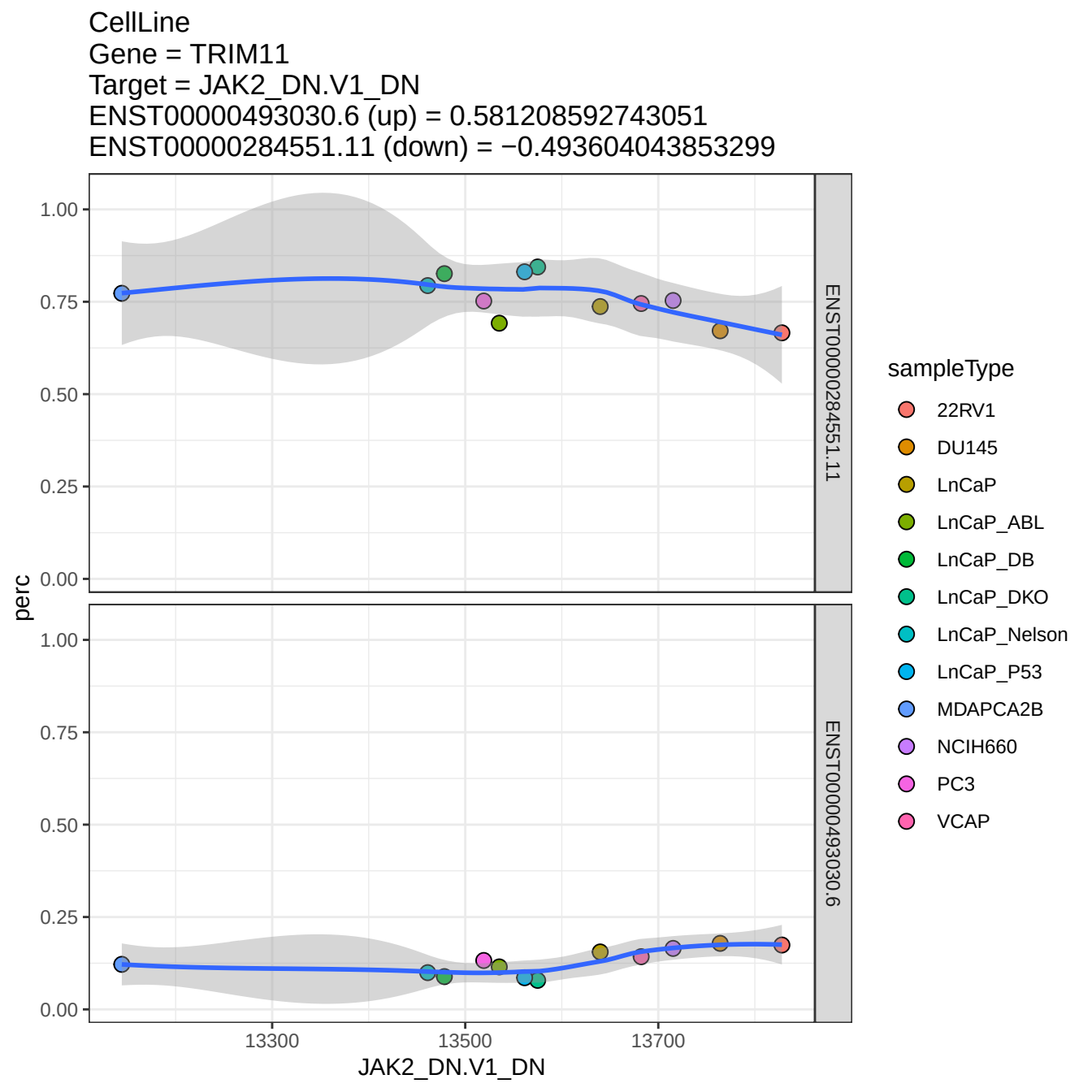

LTL_longitudinal
Gene = TRIM11
Target = SIRNA_EIF4GI_UP
ENST00000284551.11 (up) = 0.14156192486844
ENST00000493030.6 (down) = 0.264026046417041
perc
0.8
0.6
0.4
0.2
ENST00000284551.11
ENST00000493030.6
15600
16000
16400
SIRNA_EIF4GI_UP
sampleType
LTL331_Cx_12w
LTL331_Cx_8w
LTL331_Cx_R
LTL331_PreCx

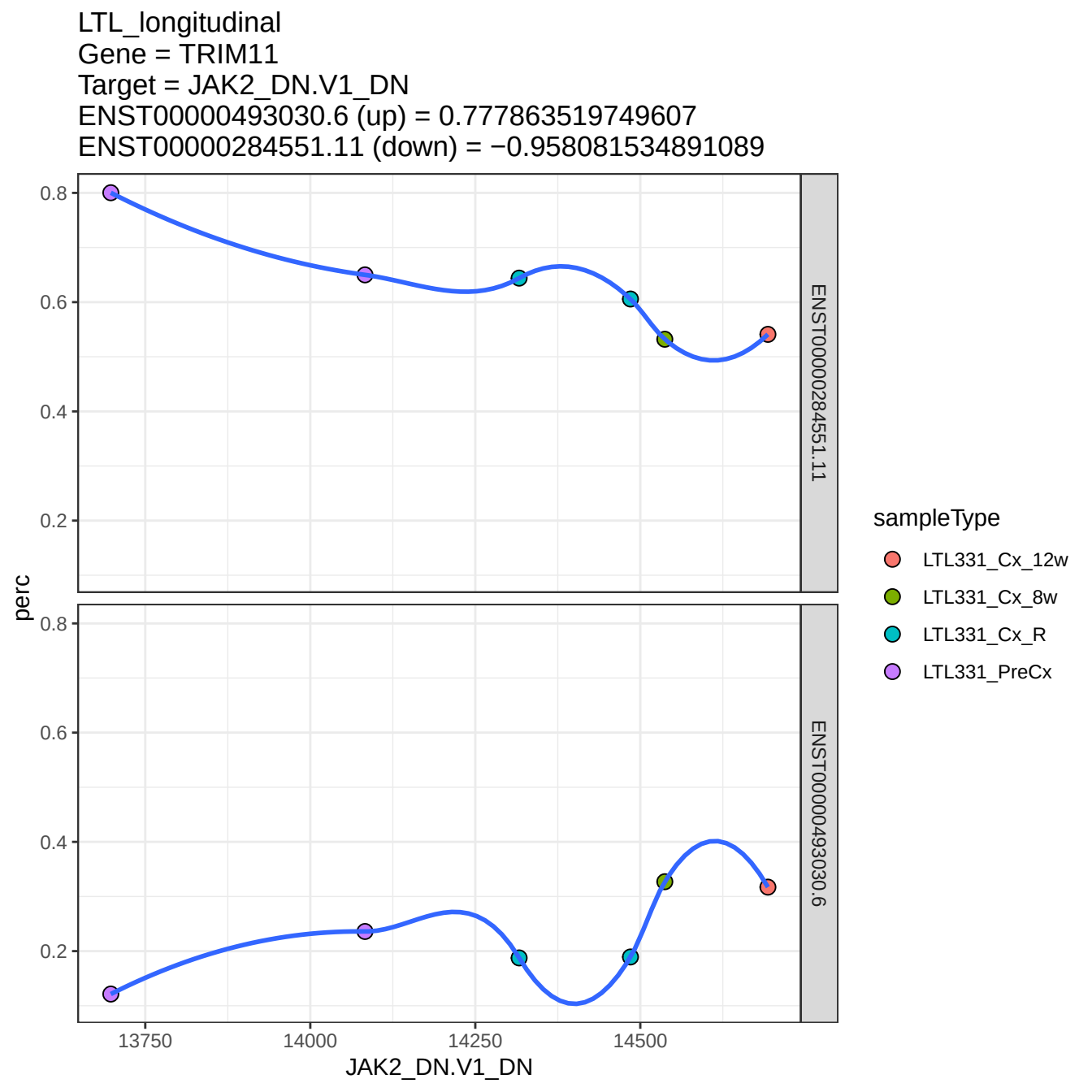

PDX_longitudinal
Gene = TRIM11
Target = SIRNA_EIF4GI_UP
ENST00000284551.11 (up) = 0.295958165788893
ENST00000493030.6 (down) = 0.0244512074367207
perc
0.00
0.25
0.50
0.75
ENST00000284551.11
ENST00000493030.6
17800
18000
18200
18400
SIRNA_EIF4GI_UP
sampleType
LnCaP
LnCaP_ABL
LnCaP_DKO_CX
LnCaP_DKO_ENZA
LnCaP_DKO_PreCX
LuCaP_147
LuCaP_147_CX

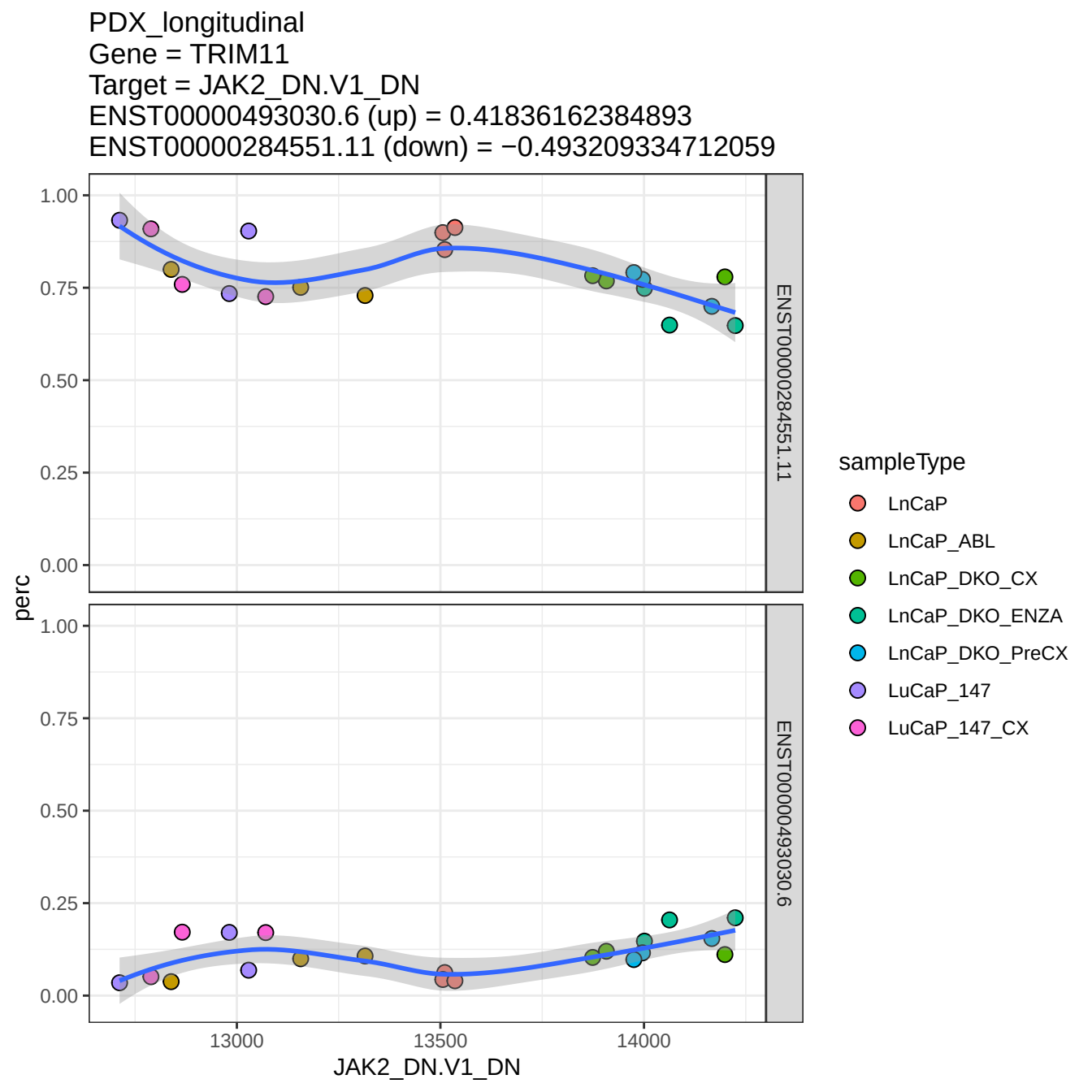

CellLine
Gene = TRIM11
Target = ESC_J1_UP_EARLY.V1_UP
ENST00000284551.11 (up) = 0.443369561655824
ENST00000493030.6 (down) = −0.285716500607017
perc
0.0
0.3
0.6
0.9
ENST00000284551.11
ENST00000493030.6
11500
11750
12000
12250
ESC_J1_UP_EARLY.V1_UP
sampleType
22RV1
DU145
LnCaP
LnCaP_ABL
LnCaP_DB
LnCaP_DKO
LnCaP_Nelson
LnCaP_P53
MDAPCA2B
NCIH660
PC3
VCAP

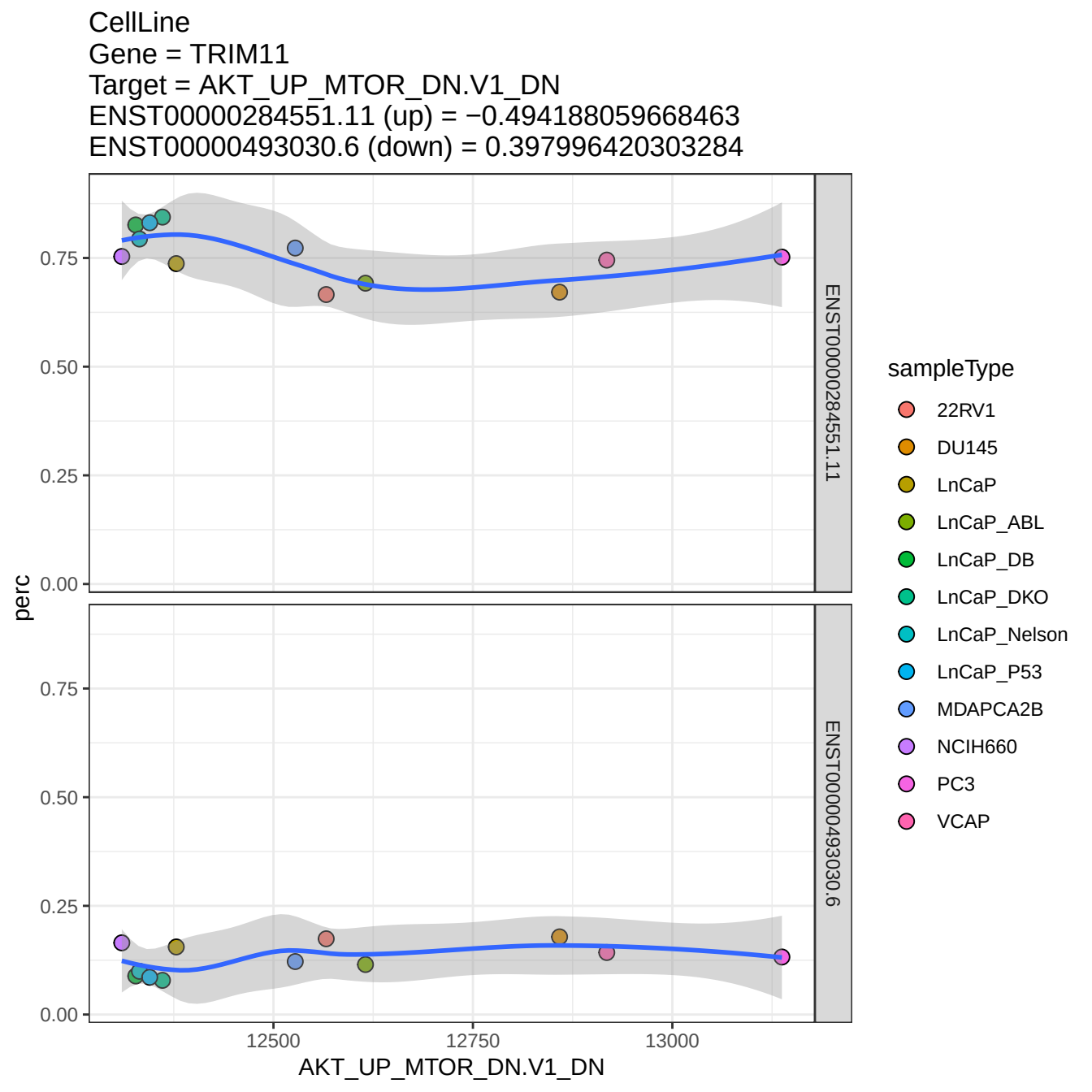

LTL_longitudinal
Gene = TRIM11
Target = ESC_J1_UP_EARLY.V1_UP
ENST00000284551.11 (up) = 0.580525424465923
ENST00000493030.6 (down) = −0.18867695022104
perc
0.8
0.6
0.4
0.2
ENST00000284551.11
ENST00000493030.6
11500
11700
11900
12100
12300
ESC_J1_UP_EARLY.V1_UP
sampleType
LTL331_Cx_12w
LTL331_Cx_8w
LTL331_Cx_R
LTL331_PreCx

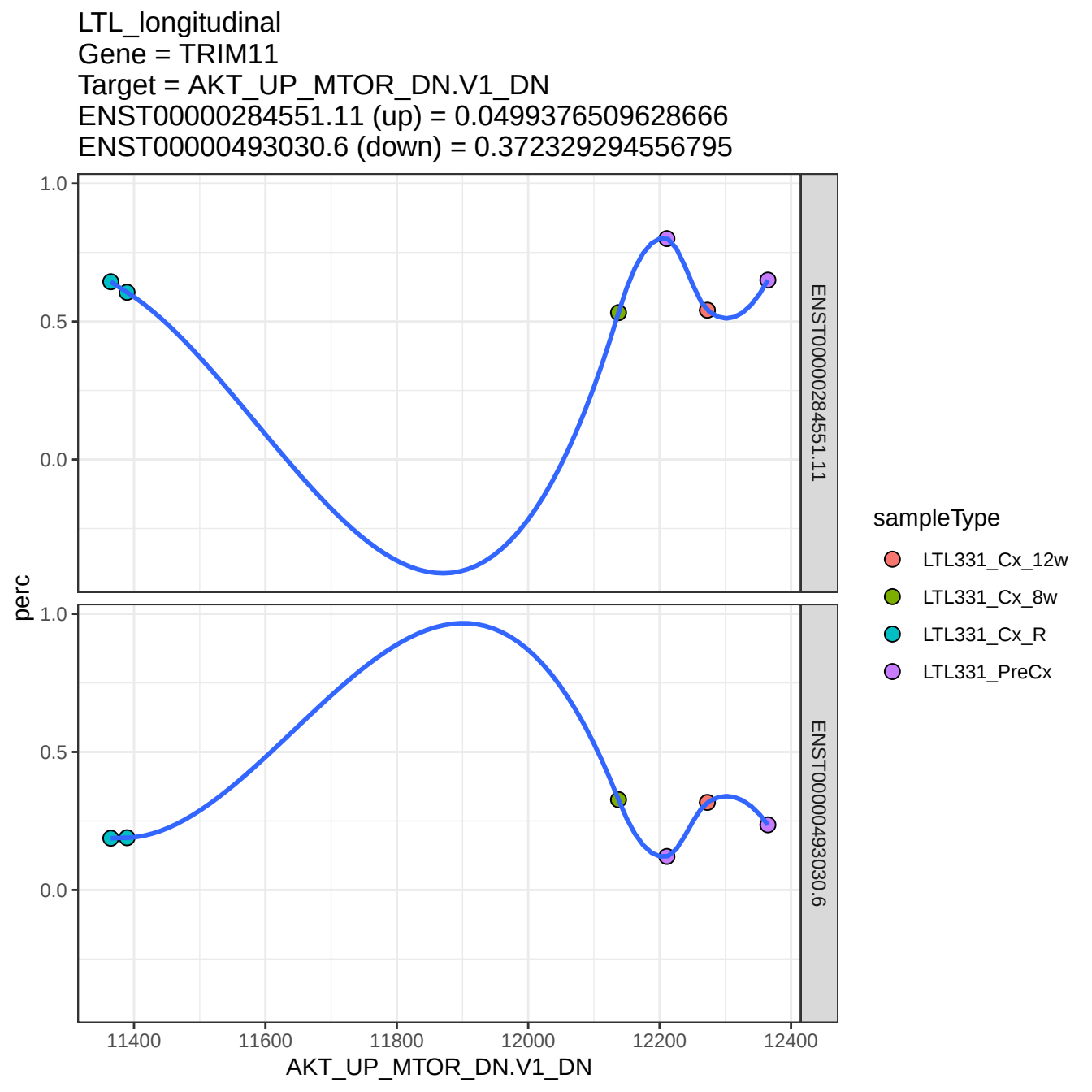

PDX_longitudinal
Gene = TRIM11
Target = ESC_J1_UP_EARLY.V1_UP
ENST00000284551.11 (up) = 0.469641363135005
ENST00000493030.6 (down) = −0.144873933497829
perc
1.00
0.75
0.50
0.25
0.00
ENST00000284551.11
ENST00000493030.6
11750
12000
12250
12500
12750
13000
ESC_J1_UP_EARLY.V1_UP
sampleType
LnCaP
LnCaP_ABL
LnCaP_DKO_CX
LnCaP_DKO_ENZA
LnCaP_DKO_PreCX
LuCaP_147
LuCaP_147_CX

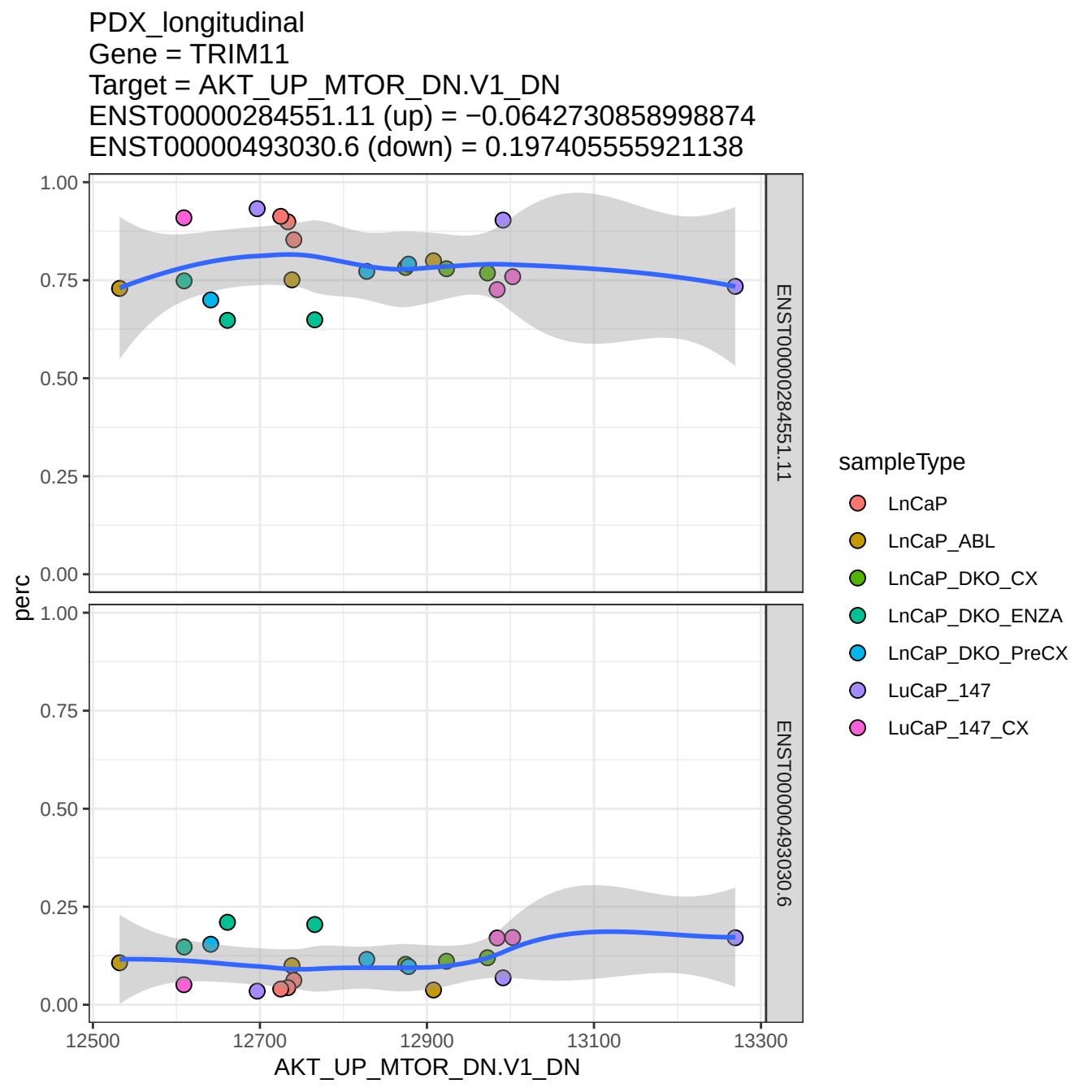

CellLine
Gene = TRIM11
Target = JAK2_DN.V1_DN
ENST00000493030.6 (up) = 0.581208592743051
ENST00000284551.11 (down) = −0.493604043853299
perc
1.00
0.75
0.50
0.25
0.00
ENST00000284551.11
ENST00000493030.6
13300
13500
13700
JAK2_DN.V1_DN
sampleType
22RV1
DU145
LnCaP
LnCaP_ABL
LnCaP_DB
LnCaP_DKO
LnCaP_Nelson
LnCaP_P53
MDAPCA2B
NCIH660
PC3
VCAP

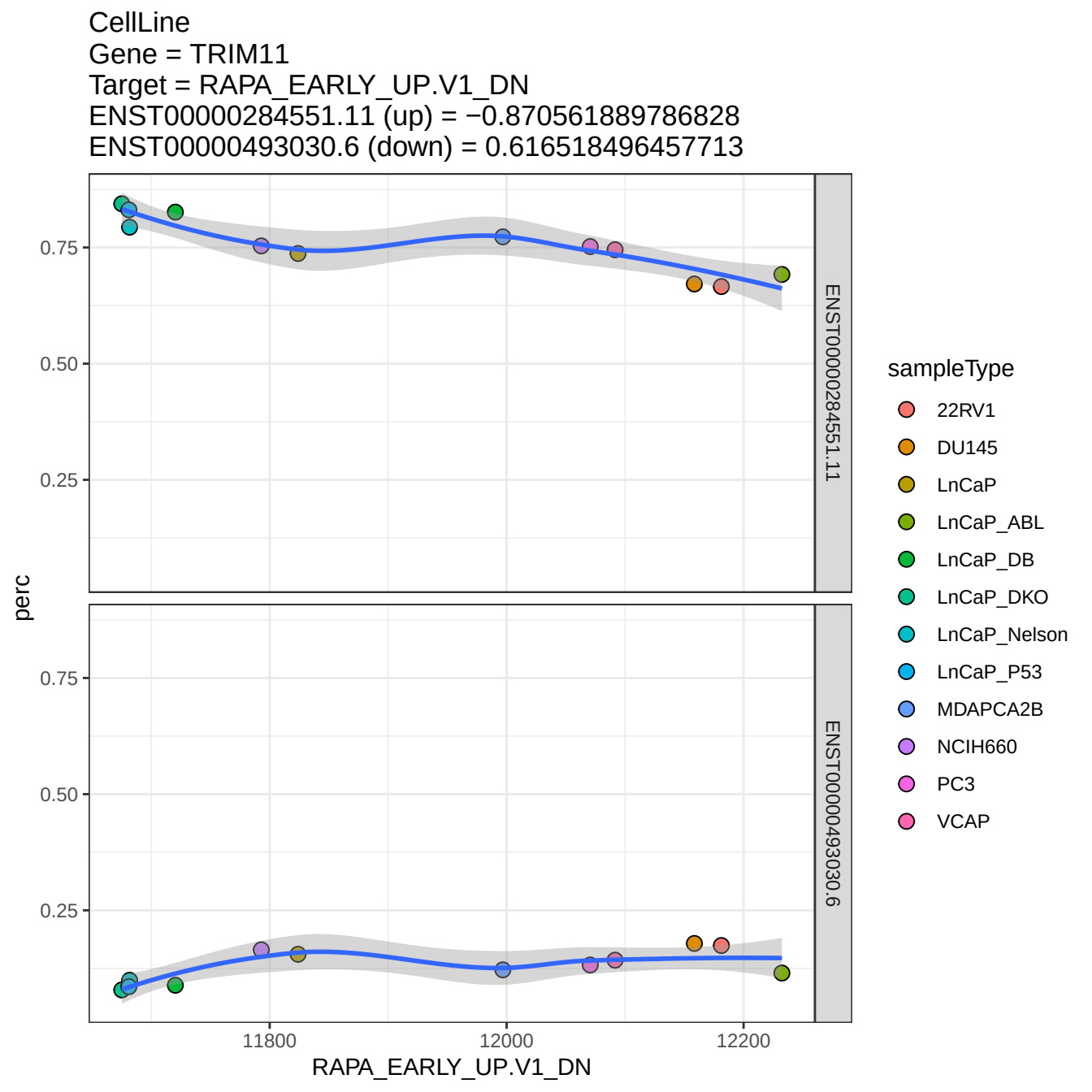

LTL_longitudinal
Gene = TRIM11
Target = JAK2_DN.V1_DN
ENST00000493030.6 (up) = 0.777863519749607
ENST00000284551.11 (down) = −0.958081534891089
perc
0.8
0.6
0.4
0.2
ENST00000284551.11
ENST00000493030.6
13750
14000
14250
14500
JAK2_DN.V1_DN
sampleType
LTL331_Cx_12w
LTL331_Cx_8w
LTL331_Cx_R
LTL331_PreCx

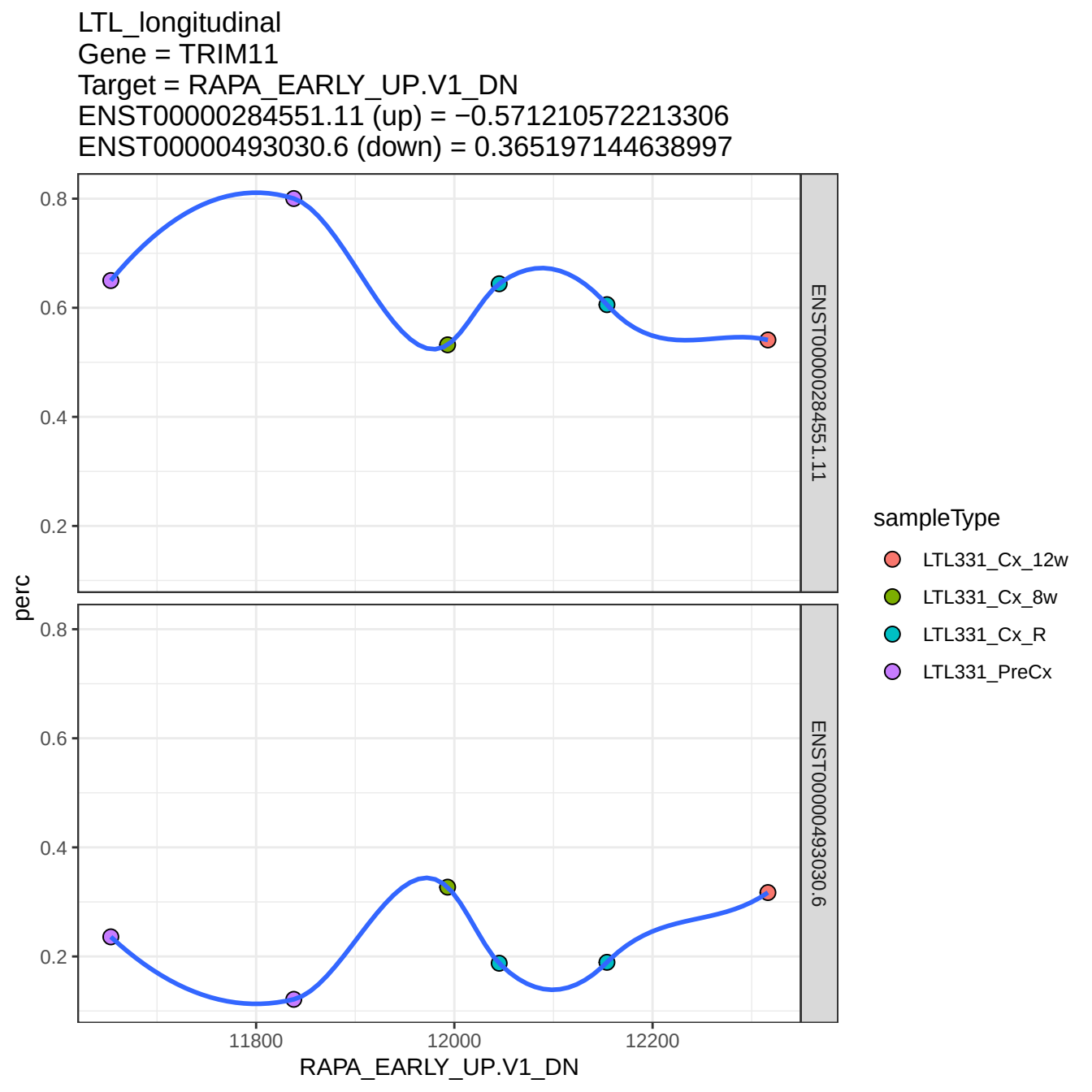

PDX_longitudinal
Gene = TRIM11
Target = JAK2_DN.V1_DN
ENST00000493030.6 (up) = 0.41836162384893
ENST00000284551.11 (down) = −0.493209334712059
perc
1.00
0.75
0.50
0.25
0.00
ENST00000284551.11
ENST00000493030.6
13000
13500
14000
JAK2_DN.V1_DN
sampleType
LnCaP
LnCaP_ABL
LnCaP_DKO_CX
LnCaP_DKO_ENZA
LnCaP_DKO_PreCX
LuCaP_147
LuCaP_147_CX


CellLine
Gene = TRIM11
Target = AKT_UP_MTOR_DN.V1_DN
ENST00000284551.11 (up) = −0.494188059668463
ENST00000493030.6 (down) = 0.397996420303284
perc
0.00
0.25
0.50
0.75
ENST00000284551.11
ENST00000493030.6
12500
12750
13000
AKT_UP_MTOR_DN.V1_DN
sampleType
22RV1
DU145
LnCaP
LnCaP_ABL
LnCaP_DB
LnCaP_DKO
LnCaP_Nelson
LnCaP_P53
MDAPCA2B
NCIH660
PC3
VCAP

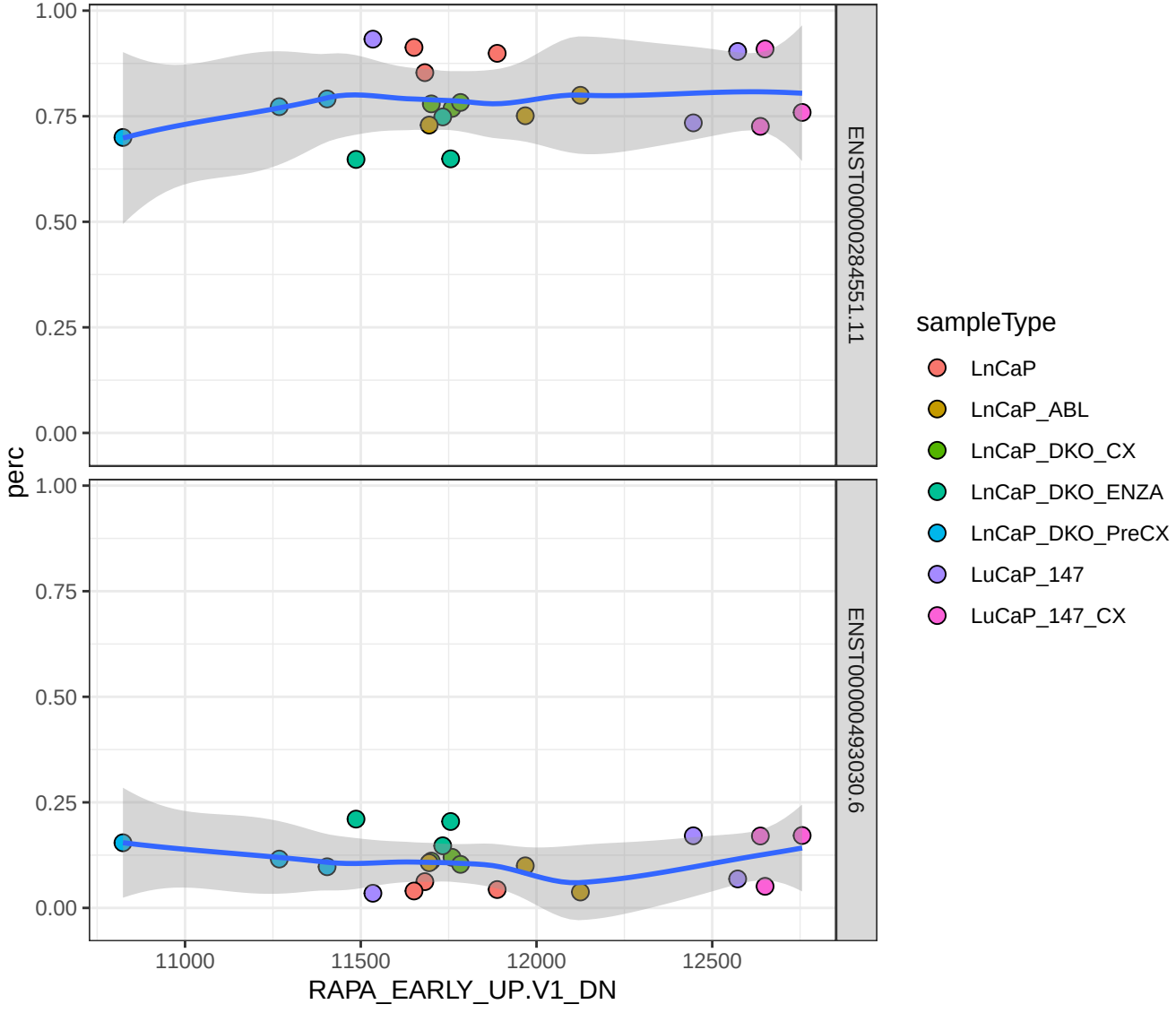

LTL_longitudinal
Gene = TRIM11
Target = AKT_UP_MTOR_DN.V1_DN
ENST00000284551.11 (up) = 0.0499376509628666
ENST00000493030.6 (down) = 0.372329294556795
perc
1.0
0.5
0.0
ENST00000284551.11
ENST00000493030.6
11400
11600
11800
12000
12200
12400
AKT_UP_MTOR_DN.V1_DN
sampleType
LTL331_Cx_12w
LTL331_Cx_8w
LTL331_Cx_R
LTL331_PreCx

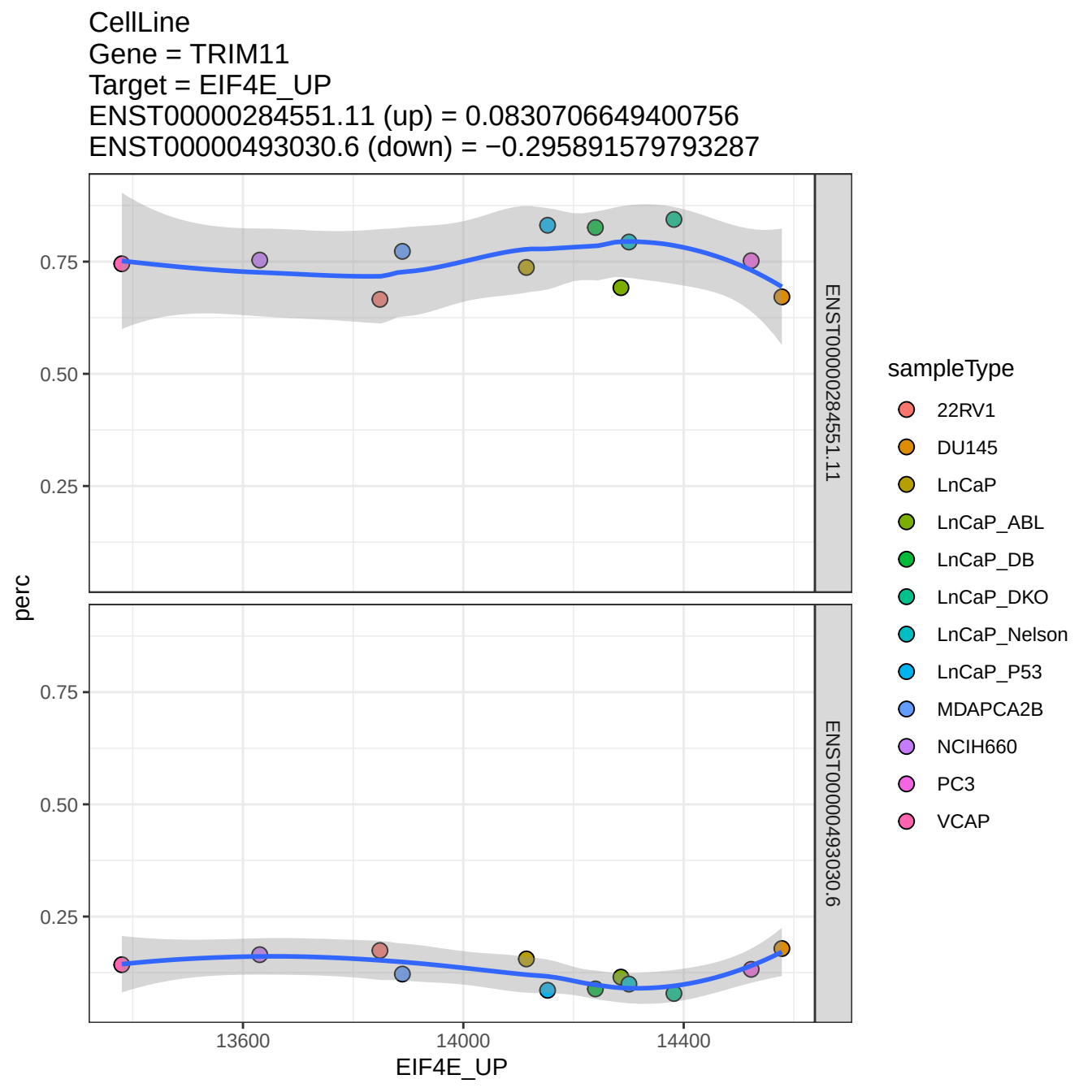

PDX_longitudinal
Gene = TRIM11
Target = AKT_UP_MTOR_DN.V1_DN
ENST00000284551.11 (up) = −0.0642730858998874
ENST00000493030.6 (down) = 0.197405555921138
perc
1.00
0.75
0.50
0.25
0.00
ENST00000284551.11
ENST00000493030.6
12500
12700
12900
13100
13300
AKT_UP_MTOR_DN.V1_DN
sampleType
LnCaP
LnCaP_ABL
LnCaP_DKO_CX
LnCaP_DKO_ENZA
LnCaP_DKO_PreCX
LuCaP_147
LuCaP_147_CX

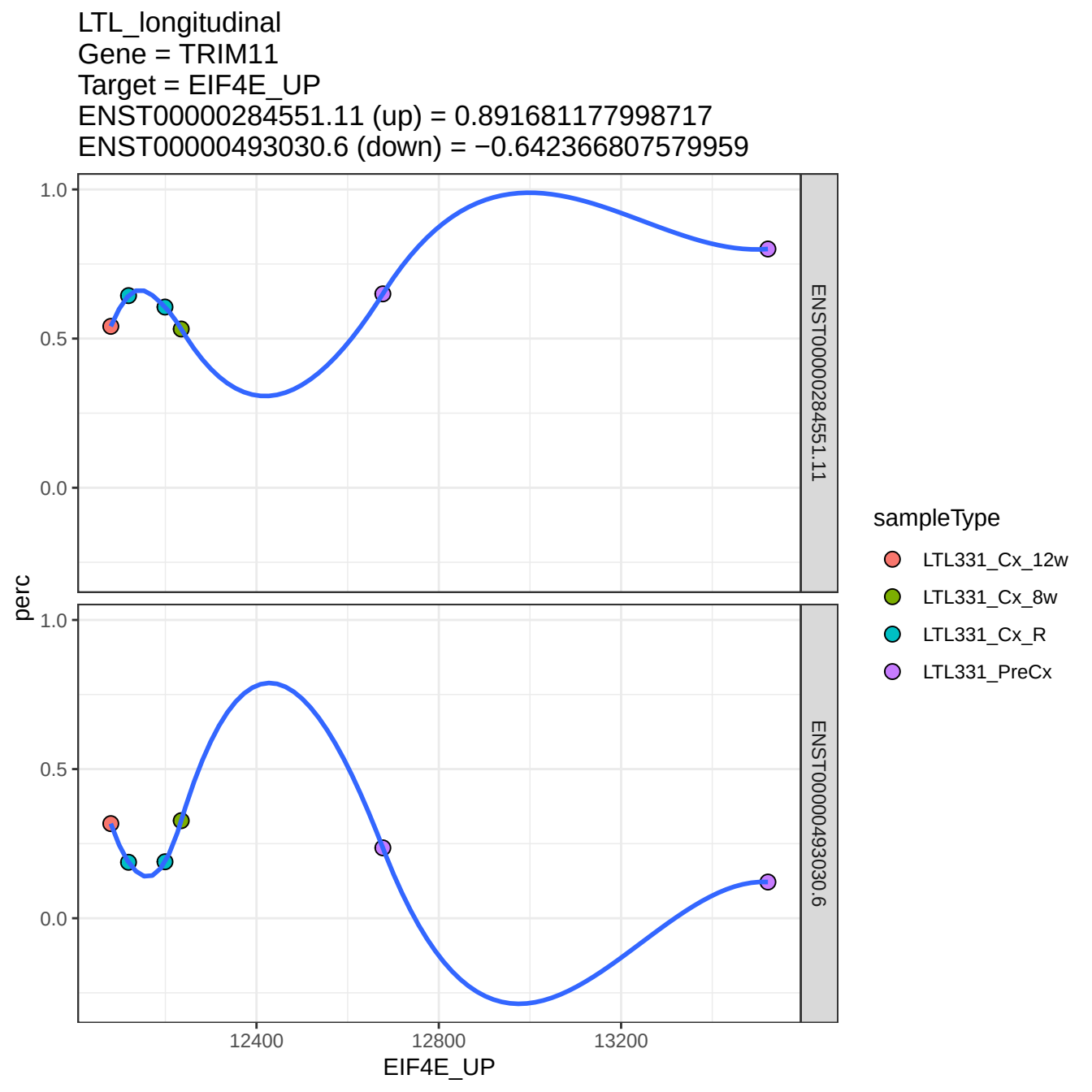

CellLine
Gene = TRIM11
Target = RAPA_EARLY_UP.V1_DN
ENST00000284551.11 (up) = −0.870561889786828
ENST00000493030.6 (down) = 0.616518496457713
perc
0.75
0.50
0.25
ENST00000284551.11
ENST00000493030.6
11800
12000
12200
RAPA_EARLY_UP.V1_DN
sampleType
22RV1
DU145
LnCaP
LnCaP_ABL
LnCaP_DB
LnCaP_DKO
LnCaP_Nelson
LnCaP_P53
MDAPCA2B
NCIH660
PC3
VCAP


LTL_longitudinal
Gene = TRIM11
Target = RAPA_EARLY_UP.V1_DN
ENST00000284551.11 (up) = −0.571210572213306
ENST00000493030.6 (down) = 0.365197144638997
perc
0.8
0.6
0.4
0.2
ENST00000284551.11
ENST00000493030.6
11800
12000
12200
RAPA_EARLY_UP.V1_DN
sampleType
LTL331_Cx_12w
LTL331_Cx_8w
LTL331_Cx_R
LTL331_PreCx

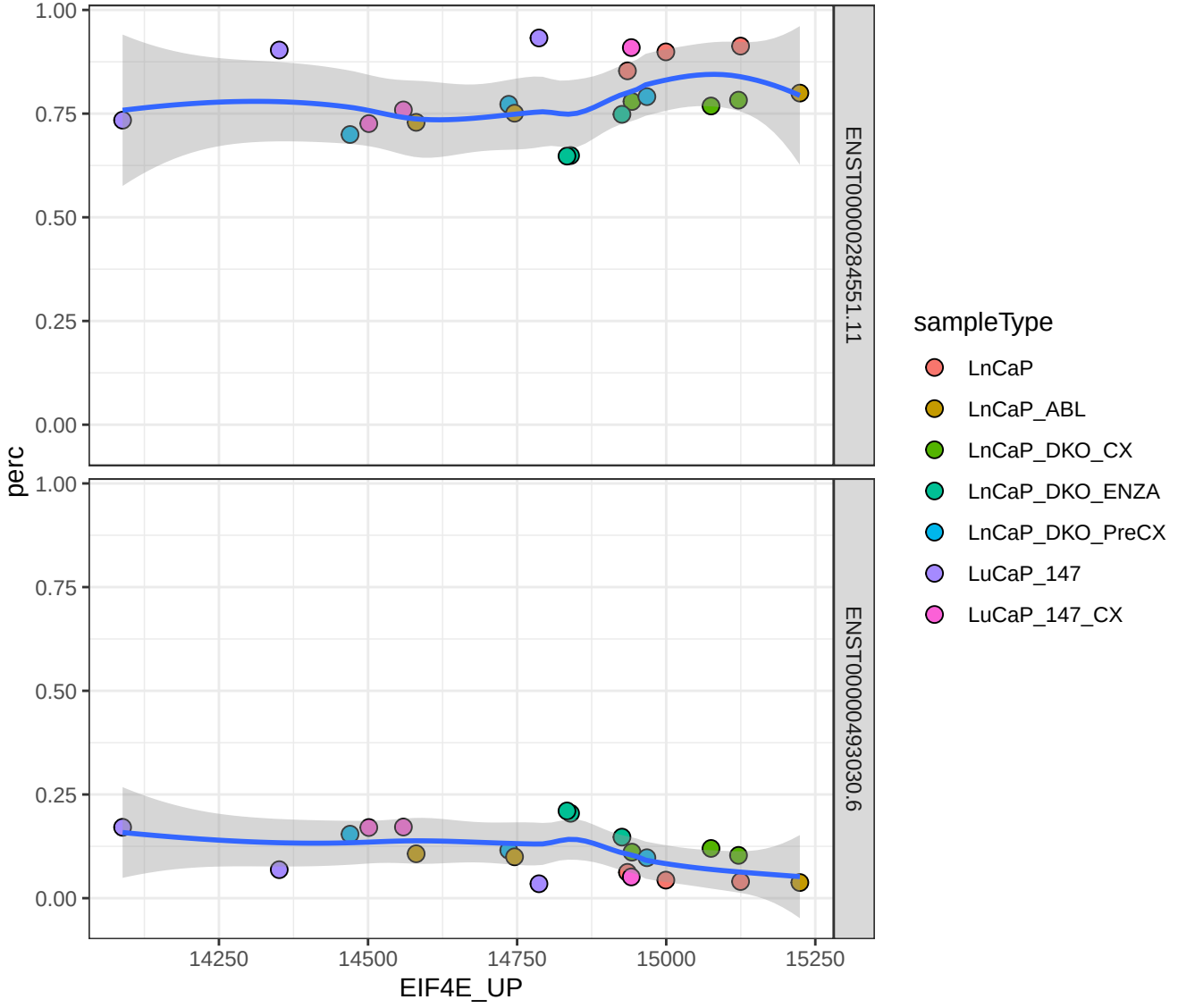
PDX_longitudinal
Gene = TRIM11
Target = RAPA_EARLY_UP.V1_DN
ENST00000284551.11 (up) = 0.206590200796484
ENST00000493030.6 (down) = −0.0156342800035312

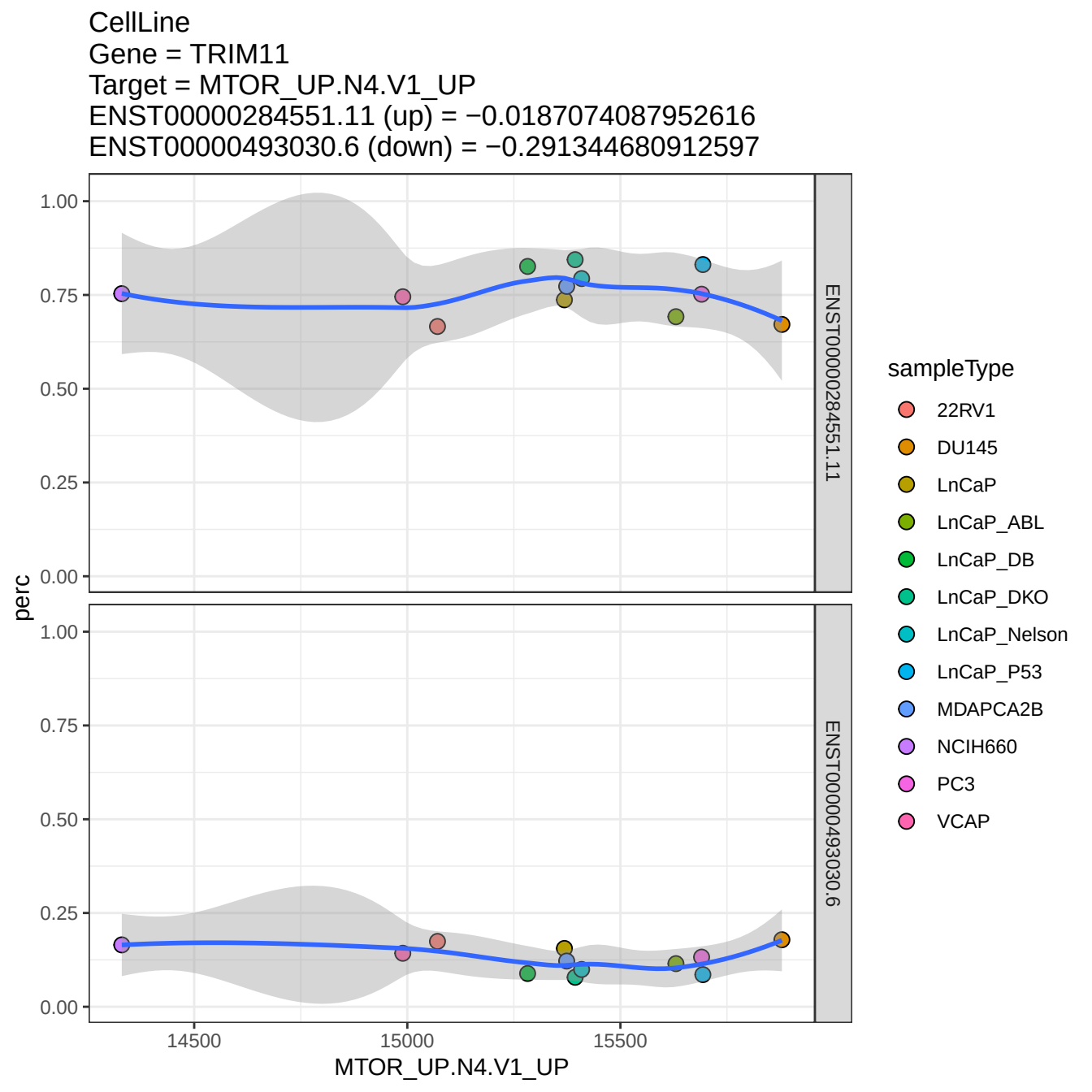


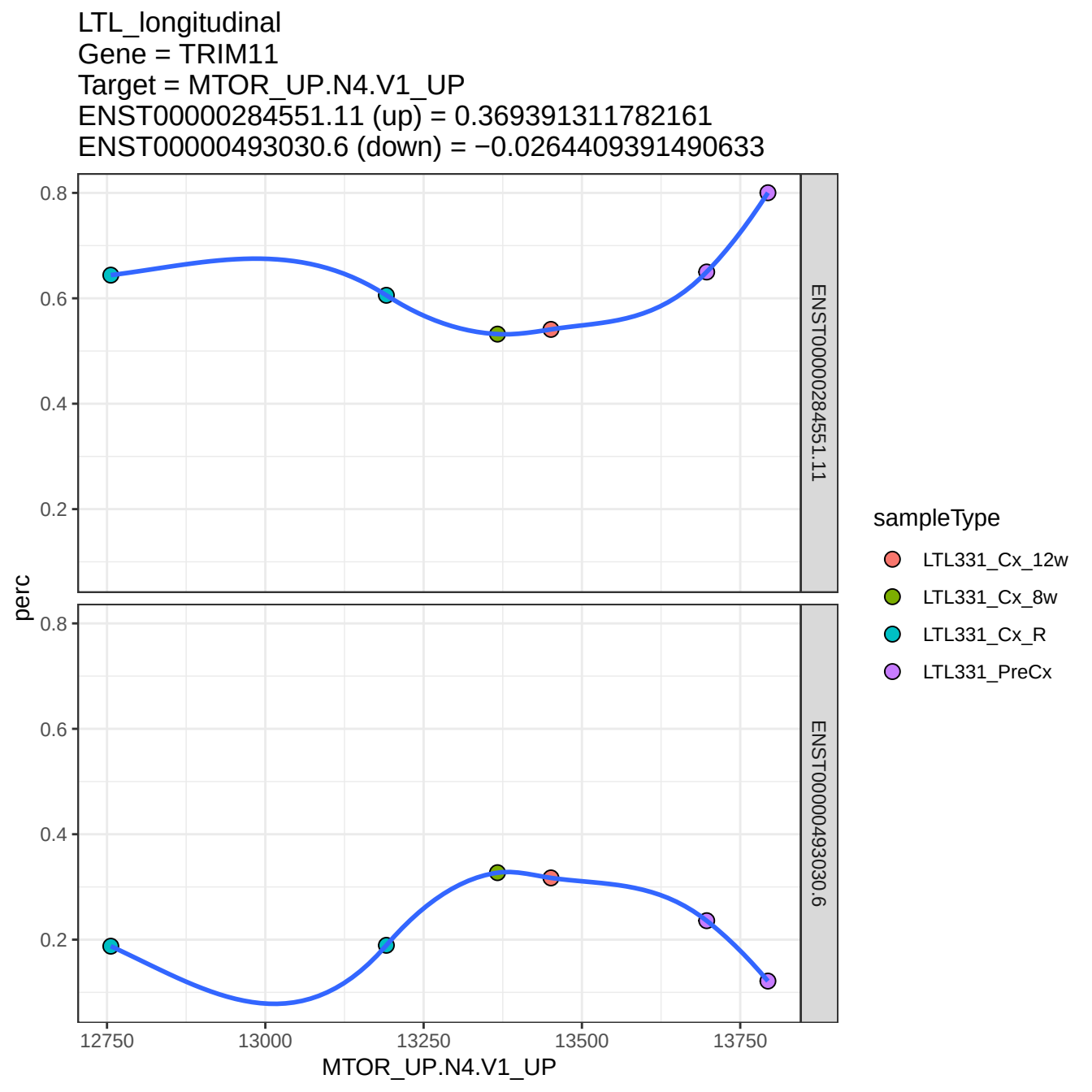

CellLine
Gene = TRIM11
Target = EIF4E_UP
ENST00000284551.11 (up) = 0.0830706649400756
ENST00000493030.6 (down) = −0.295891579793287
perc
0.75
0.50
0.25
ENST00000284551.11
ENST00000493030.6
13600
14000
14400
EIF4E_UP
sampleType
22RV1
DU145
LnCaP
LnCaP_ABL
LnCaP_DB
LnCaP_DKO
LnCaP_Nelson
LnCaP_P53
MDAPCA2B
NCIH660
PC3
VCAP

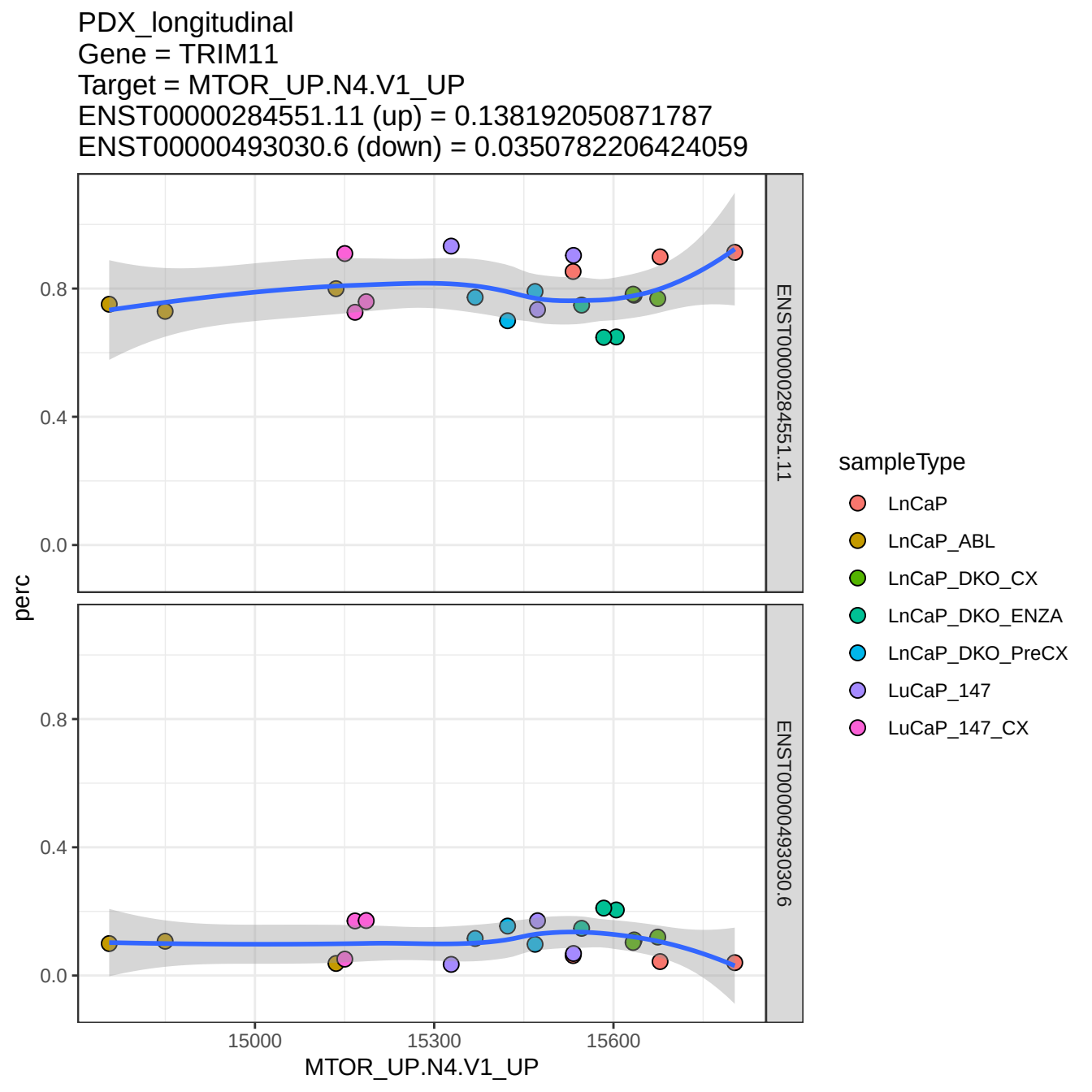

LTL_longitudinal
Gene = TRIM11
Target = EIF4E_UP
ENST00000284551.11 (up) = 0.891681177998717
ENST00000493030.6 (down) = −0.642366807579959
1.0
0.5
0.0
perc
ENST00000284551.11
ENST00000493030.6
12400
12800
13200
EIF4E_UP
sampleType
LTL331_Cx_12w
LTL331_Cx_8w
LTL331_Cx_R
LTL331_PreCx

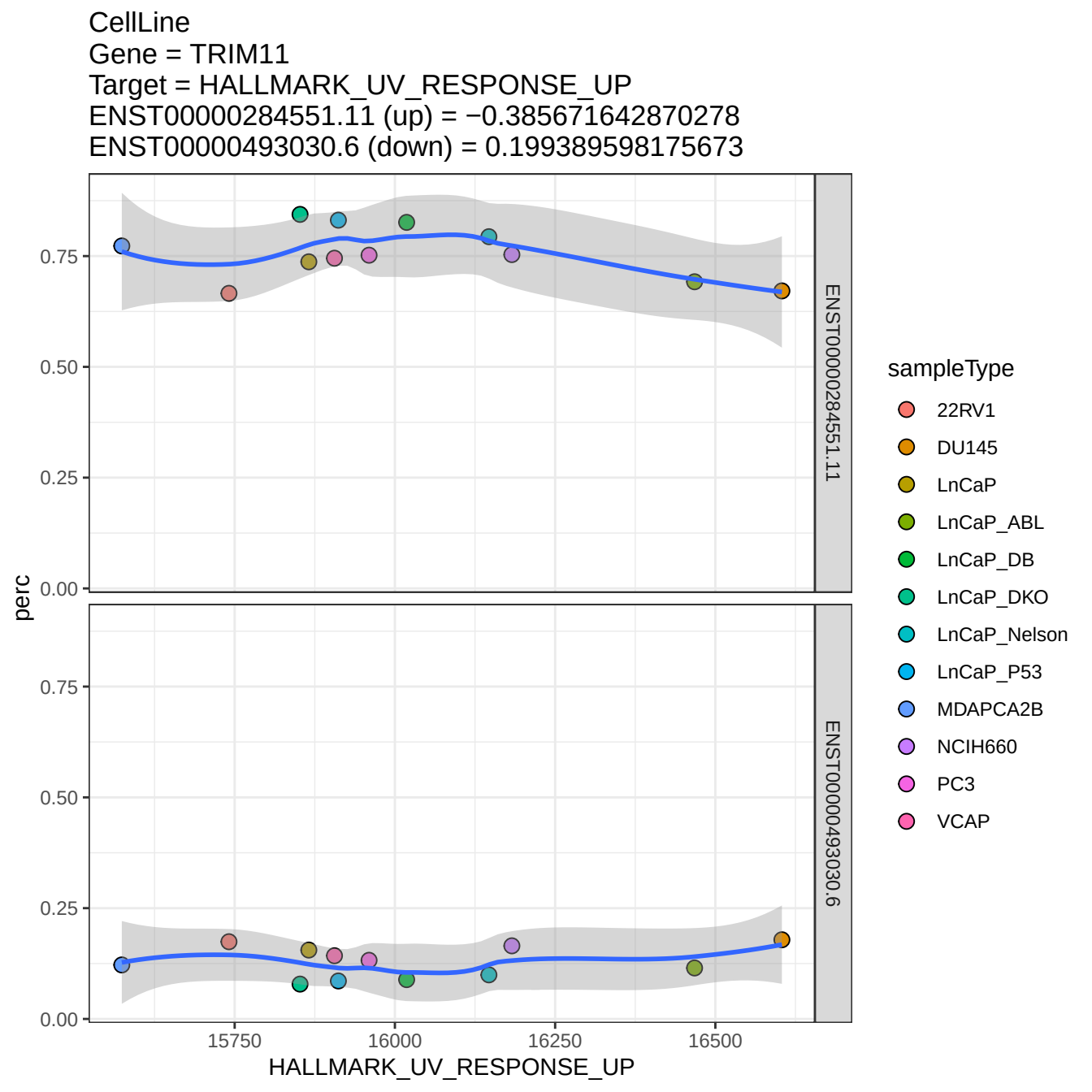
PDX_longitudinal
Gene = TRIM11
Target = EIF4E_UP
ENST00000284551.11 (up) = 0.260516463599295
ENST00000493030.6 (down) = −0.446445702301439

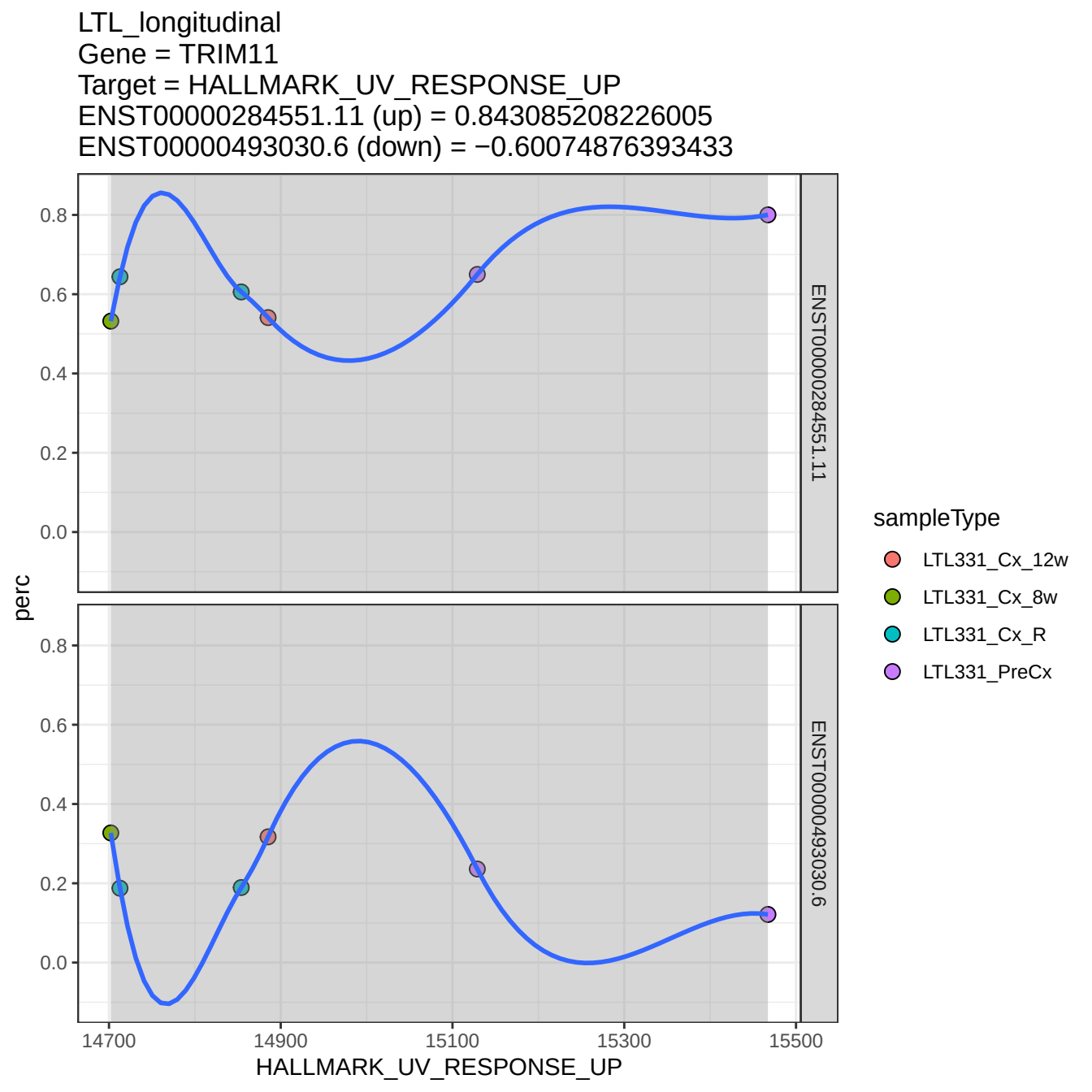



CellLine
Gene = TRIM11
Target = MTOR_UP.N4.V1_UP
ENST00000284551.11 (up) = −0.0187074087952616
ENST00000493030.6 (down) = −0.291344680912597
perc
1.00
0.75
0.50
0.25
0.00
ENST00000284551.11
ENST00000493030.6
14500
15000
15500
MTOR_UP.N4.V1_UP
sampleType
22RV1
DU145
LnCaP
LnCaP_ABL
LnCaP_DB
LnCaP_DKO
LnCaP_Nelson
LnCaP_P53
MDAPCA2B
NCIH660
PC3
VCAP

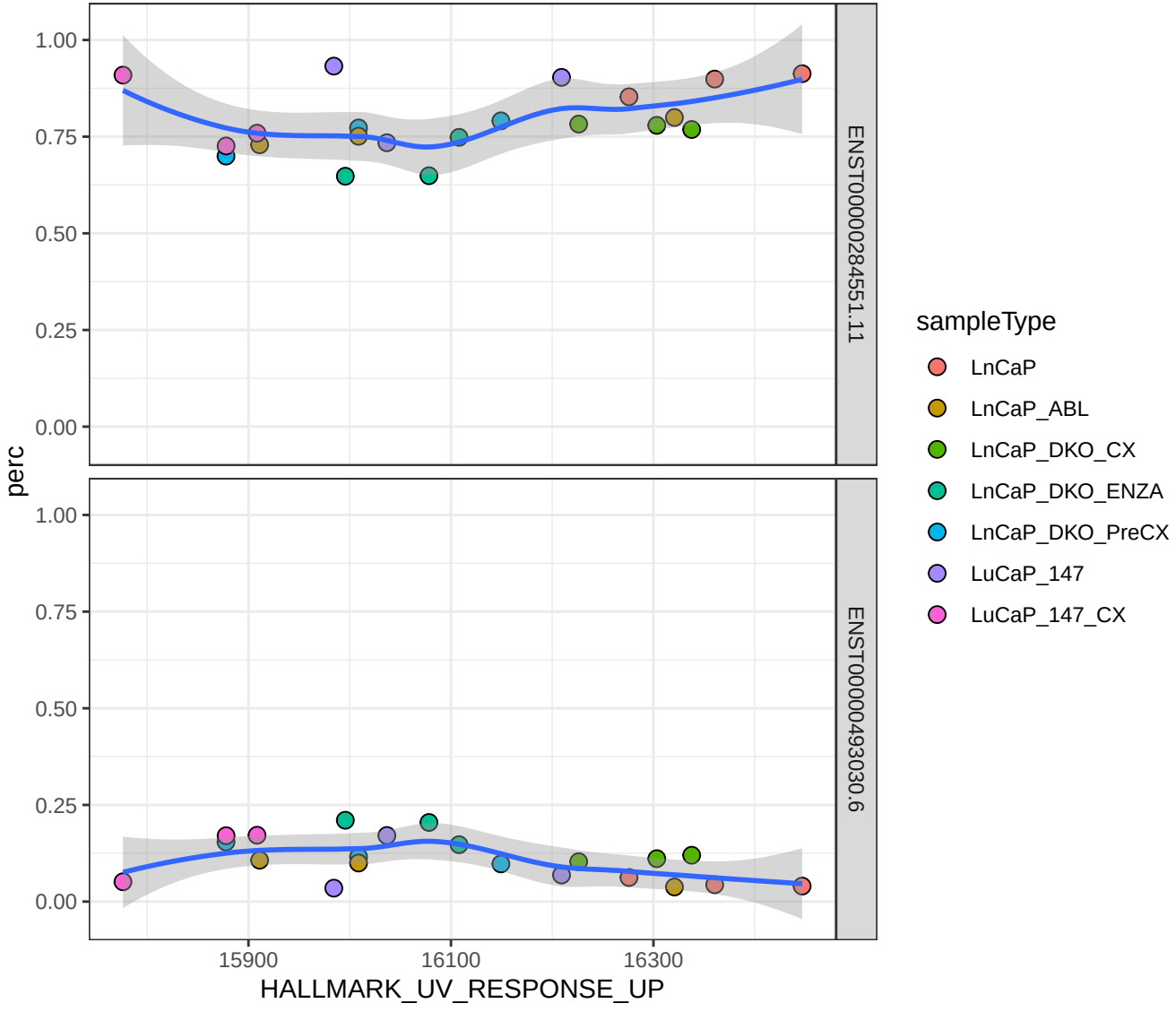

LTL_longitudinal
Gene = TRIM11
Target = MTOR_UP.N4.V1_UP
ENST00000284551.11 (up) = 0.369391311782161
ENST00000493030.6 (down) = −0.0264409391490633
perc
0.8
0.6
0.4
0.2
ENST00000284551.11
ENST00000493030.6
12750
13000
13250
13500
13750
MTOR_UP.N4.V1_UP
sampleType
LTL331_Cx_12w
LTL331_Cx_8w
LTL331_Cx_R
LTL331_PreCx


PDX_longitudinal
Gene = TRIM11
Target = MTOR_UP.N4.V1_UP
ENST00000284551.11 (up) = 0.138192050871787
ENST00000493030.6 (down) = 0.0350782206424059
perc
0.0
0.4
0.8
ENST00000284551.11
ENST00000493030.6
15000
15300
15600
MTOR_UP.N4.V1_UP
sampleType
LnCaP
LnCaP_ABL
LnCaP_DKO_CX
LnCaP_DKO_ENZA
LnCaP_DKO_PreCX
LuCaP_147
LuCaP_147_CX

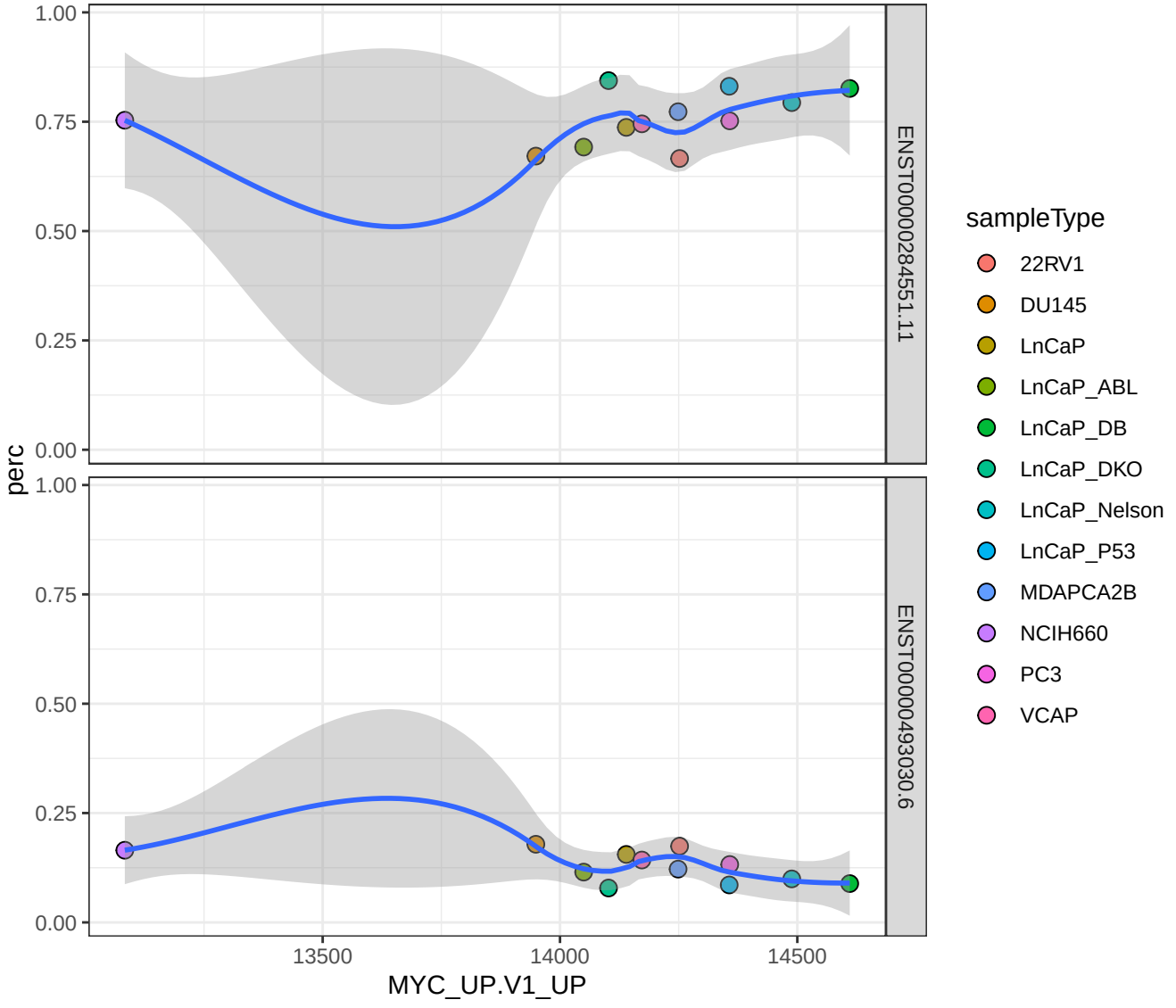

CellLine
Gene = TRIM11
Target = HALLMARK_UV_RESPONSE_UP
ENST00000284551.11 (up) = −0.385671642870278
ENST00000493030.6 (down) = 0.199389598175673
perc
0.00
0.25
0.50
0.75
ENST00000284551.11
ENST00000493030.6
15750
16000
16250
16500
HALLMARK_UV_RESPONSE_UP
sampleType
22RV1
DU145
LnCaP
LnCaP_ABL
LnCaP_DB
LnCaP_DKO
LnCaP_Nelson
LnCaP_P53
MDAPCA2B
NCIH660
PC3
VCAP

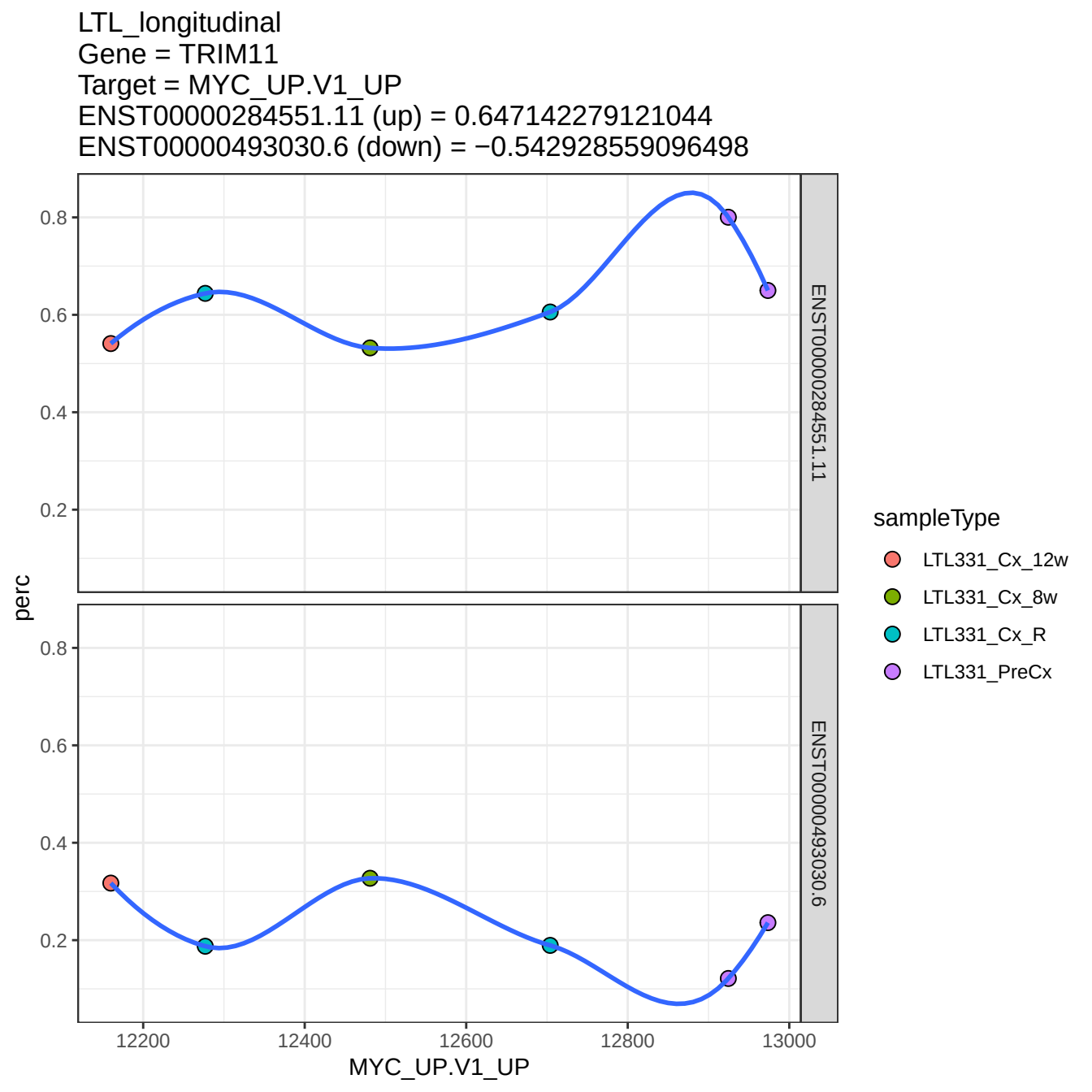

LTL_longitudinal
Gene = TRIM11
Target = HALLMARK_UV_RESPONSE_UP
ENST00000284551.11 (up) = 0.843085208226005
ENST00000493030.6 (down) = −0.60074876393433
perc
0.0
0.2
0.4
0.6
0.8
ENST00000284551.11
ENST00000493030.6
14700
14900
15100
15300
15500
HALLMARK_UV_RESPONSE_UP
sampleType
LTL331_Cx_12w
LTL331_Cx_8w
LTL331_Cx_R
LTL331_PreCx

PDX_longitudinal
Gene = TRIM11
Target = HALLMARK_UV_RESPONSE_UP
ENST00000284551.11 (up) = 0.348591548707166
ENST00000493030.6 (down) = −0.436706843337271

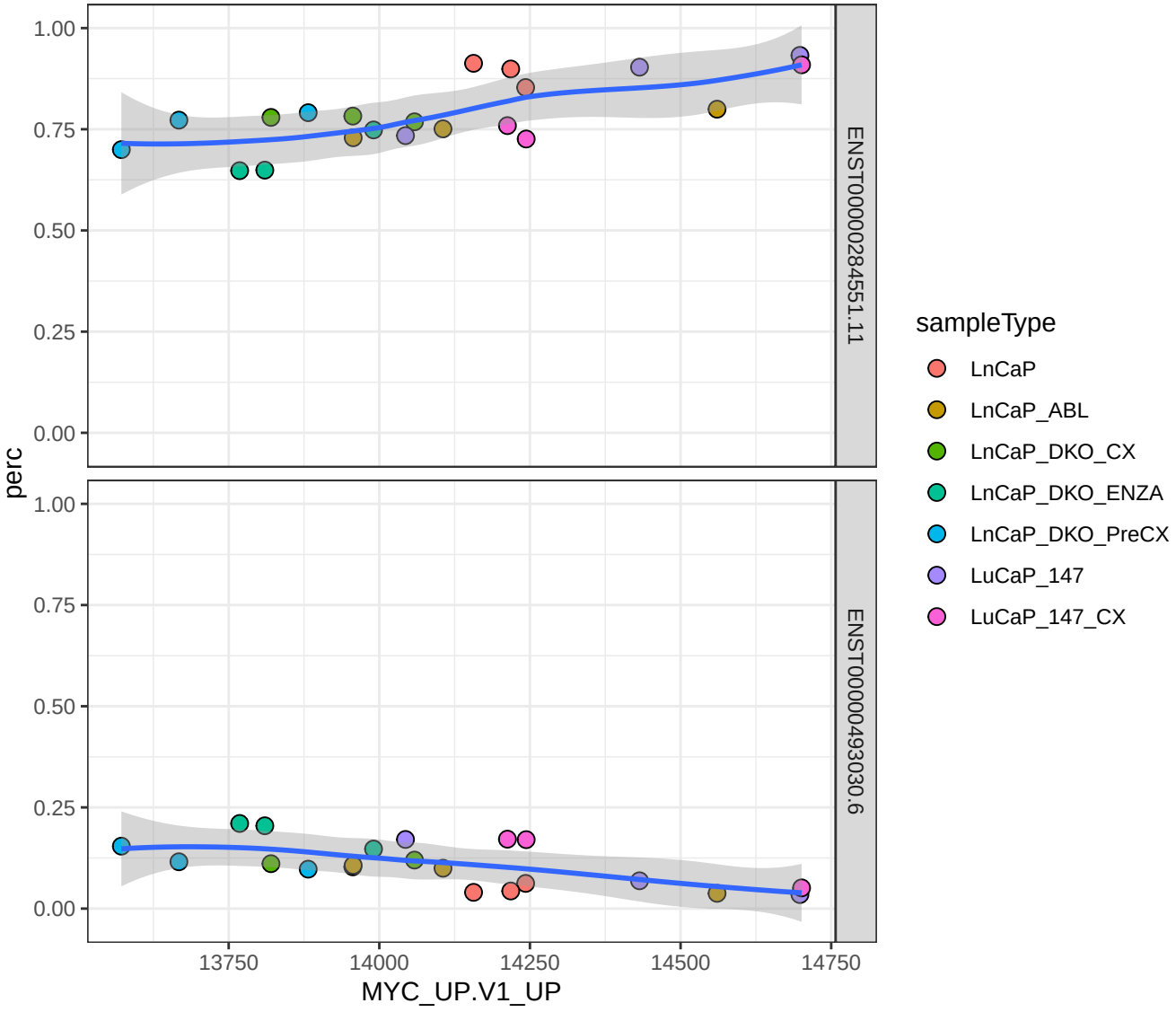


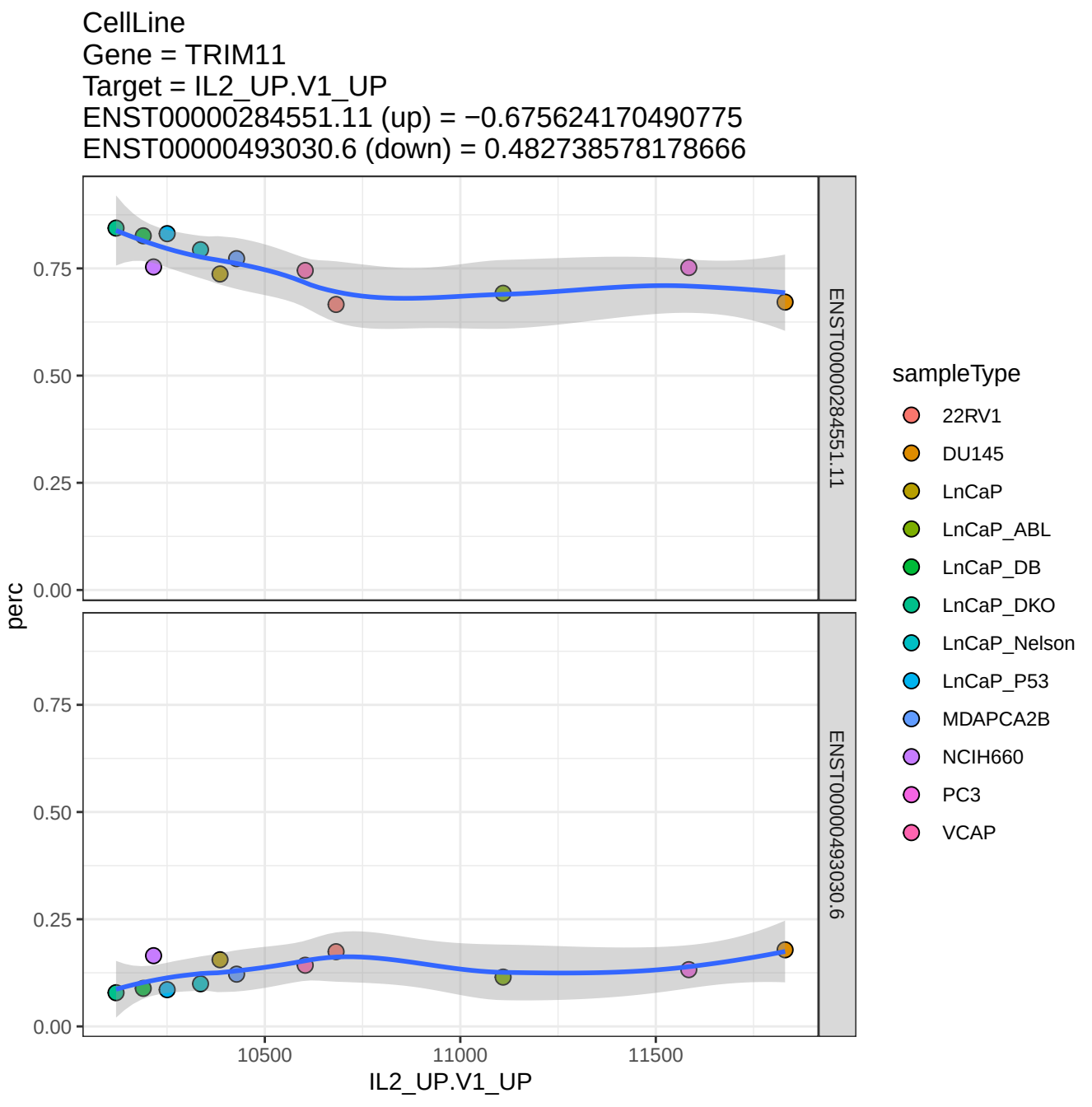
CellLine
Gene = TRIM11
Target = MYC_UP.V1_UP
ENST00000284551.11 (up) = 0.291156520742859
ENST00000493030.6 (down) = −0.516136417783064

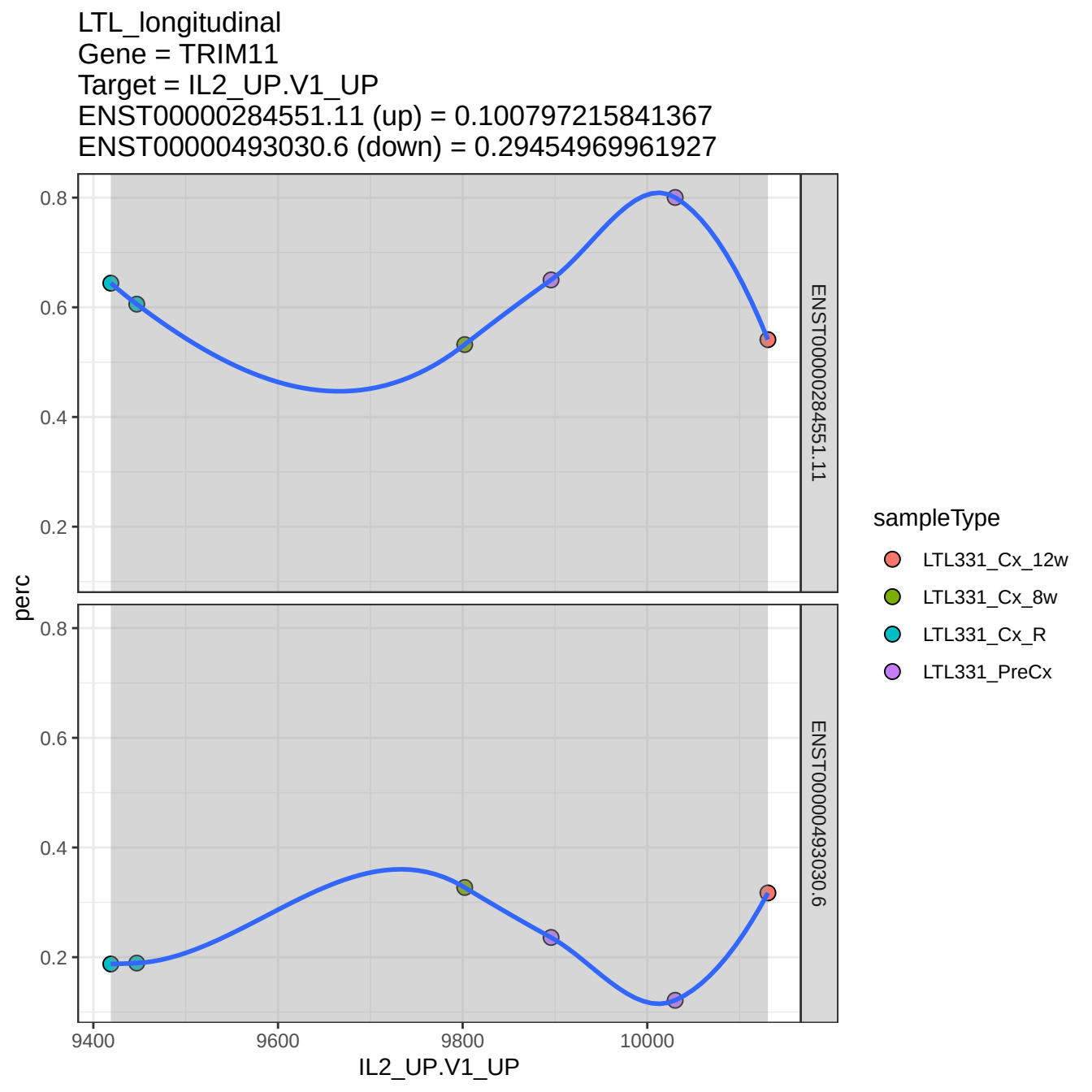


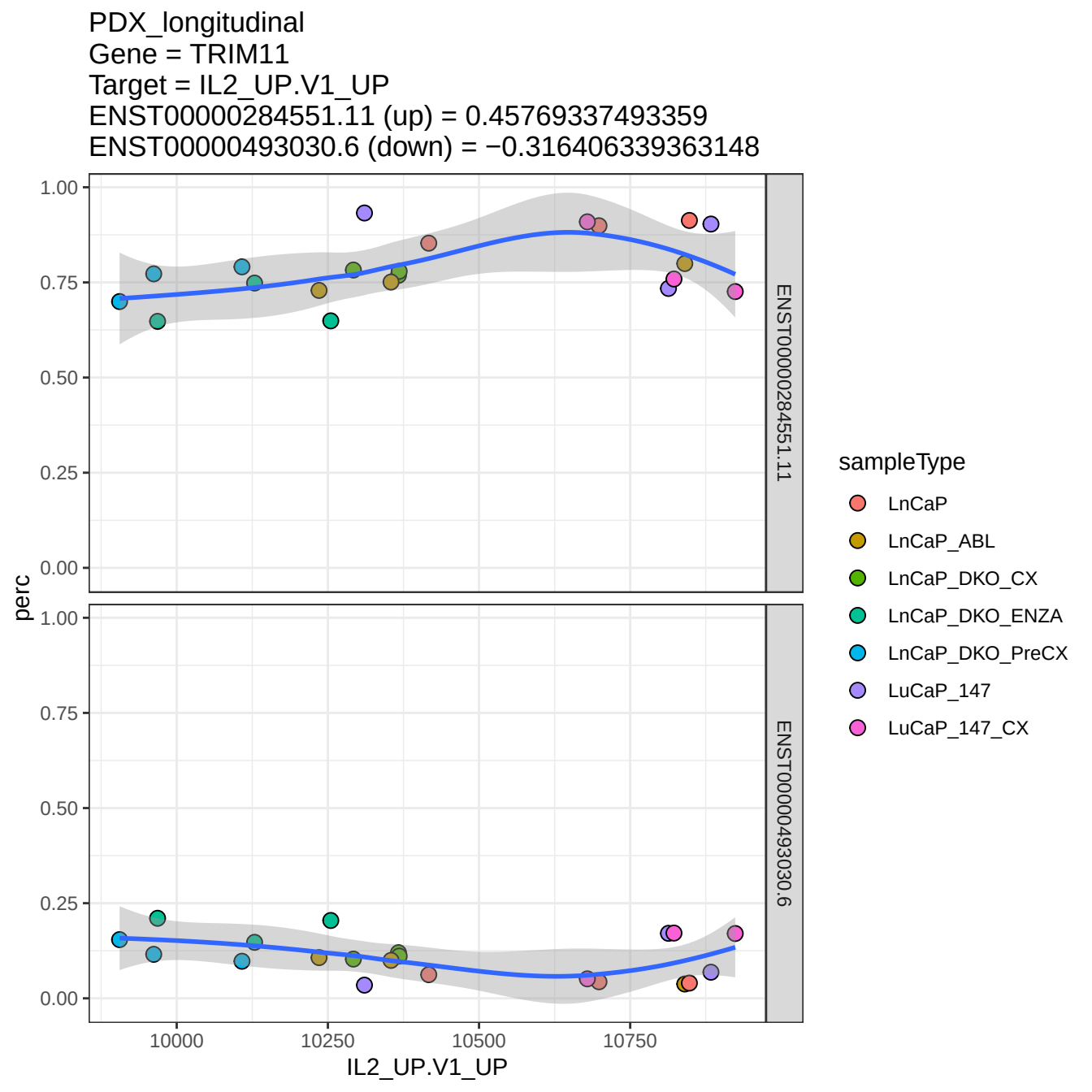

LTL_longitudinal
Gene = TRIM11
Target = MYC_UP.V1_UP
ENST00000284551.11 (up) = 0.647142279121044
ENST00000493030.6 (down) = −0.542928559096498
perc
0.8
0.6
0.4
0.2
ENST00000284551.11
ENST00000493030.6
12200
12400
12600
12800
13000
MYC_UP.V1_UP
sampleType
LTL331_Cx_12w
LTL331_Cx_8w
LTL331_Cx_R
LTL331_PreCx

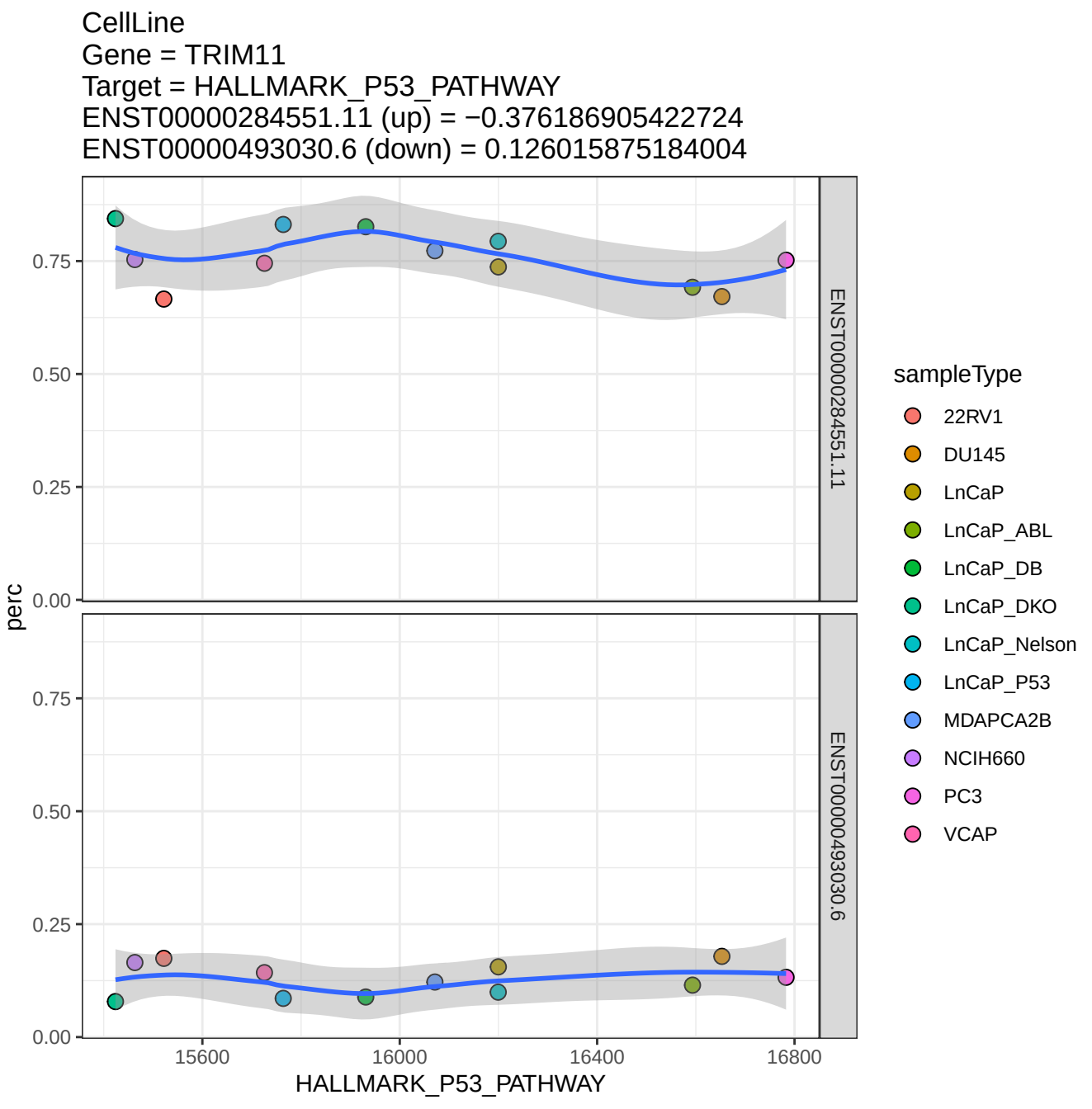
PDX_longitudinal
Gene = TRIM11
Target = MYC_UP.V1_UP
ENST00000284551.11 (up) = 0.718051576803842
ENST00000493030.6 (down) = −0.636917809156402

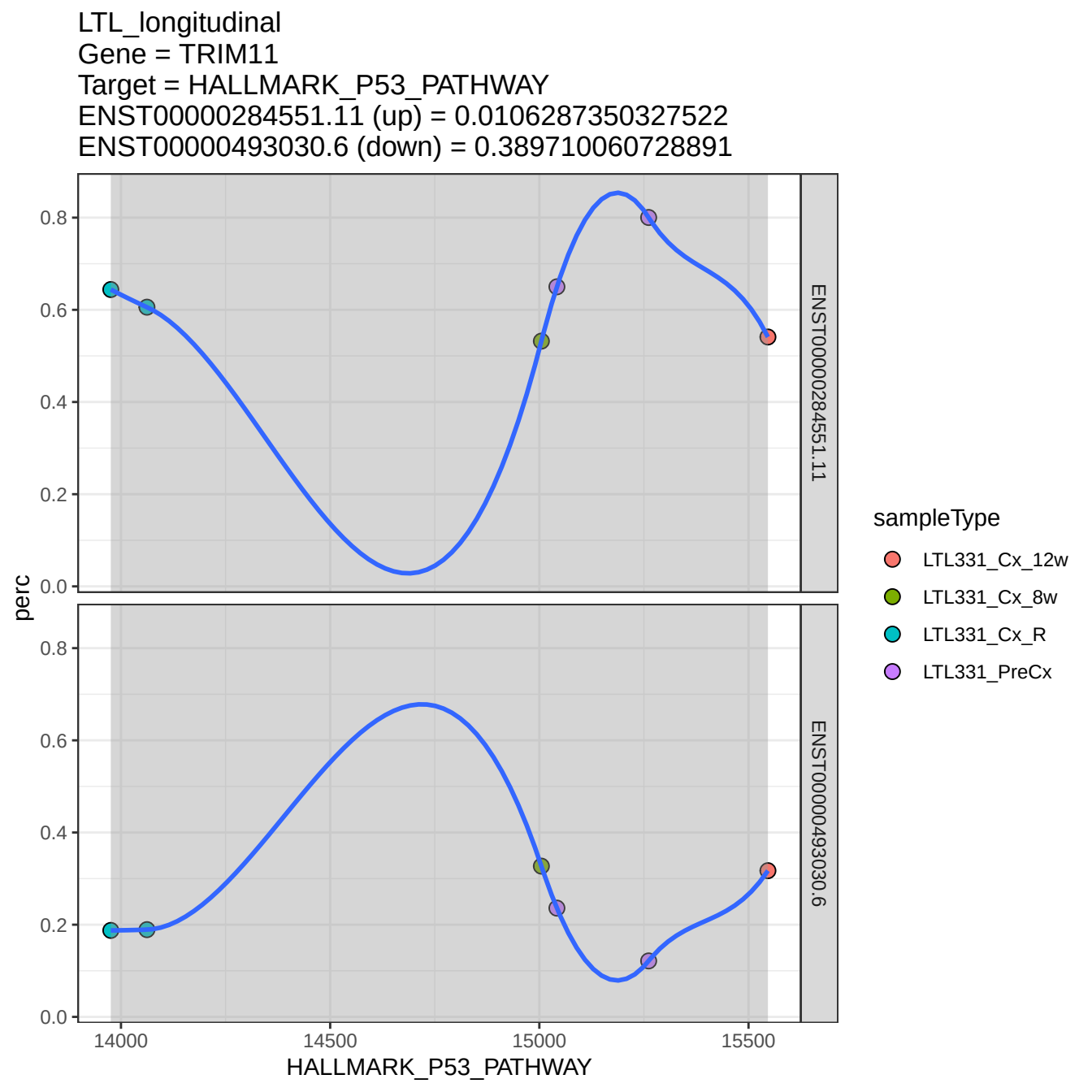



CellLine
Gene = TRIM11
Target = IL2_UP.V1_UP
ENST00000284551.11 (up) = −0.675624170490775
ENST00000493030.6 (down) = 0.482738578178666
perc
0.00
0.25
0.50
0.75
ENST00000284551.11
ENST00000493030.6
10500
11000
11500
IL2_UP.V1_UP
sampleType
22RV1
DU145
LnCaP
LnCaP_ABL
LnCaP_DB
LnCaP_DKO
LnCaP_Nelson
LnCaP_P53
MDAPCA2B
NCIH660
PC3
VCAP

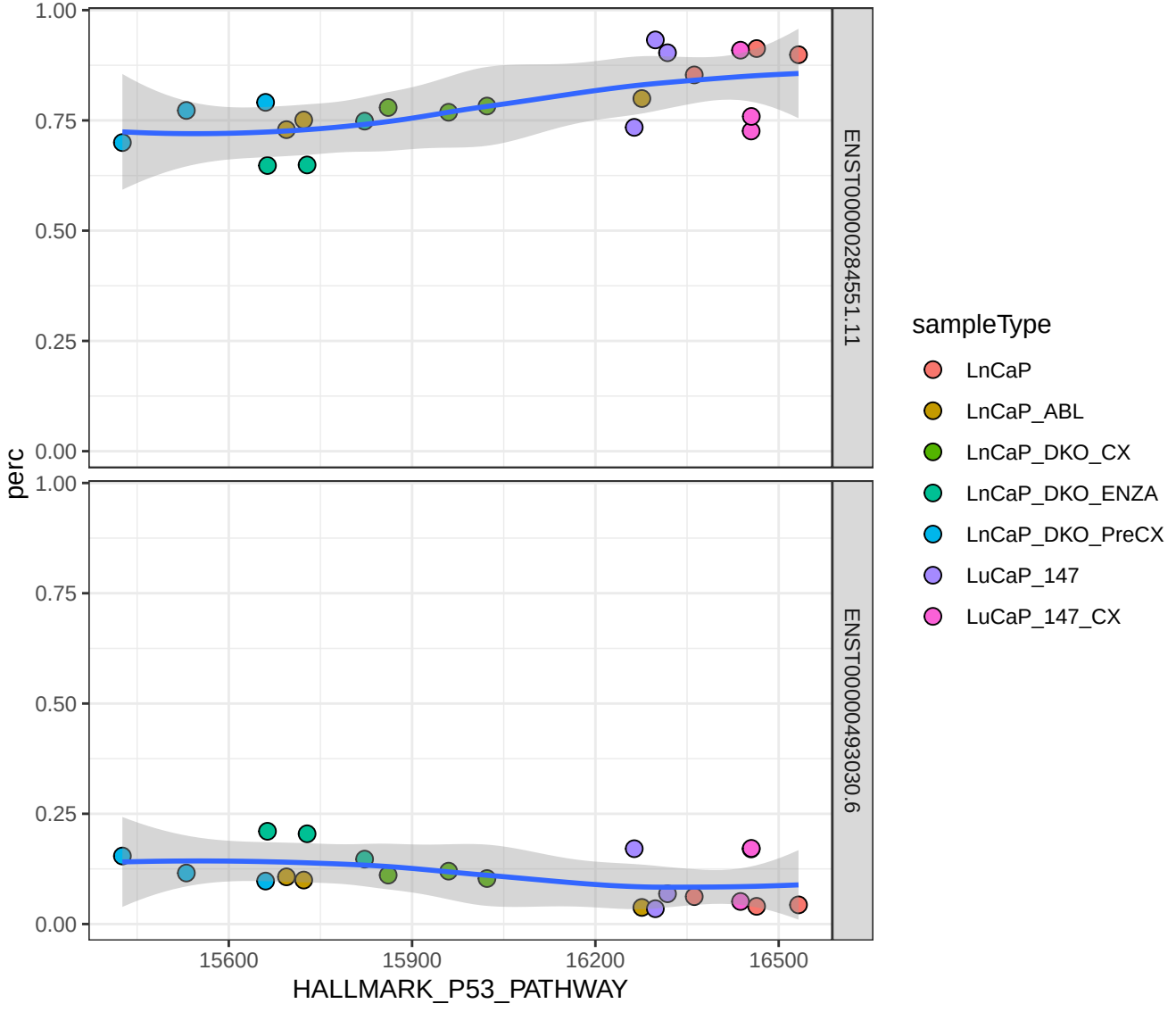

LTL_longitudinal
Gene = TRIM11
Target = IL2_UP.V1_UP
ENST00000284551.11 (up) = 0.100797215841367
ENST00000493030.6 (down) = 0.29454969961927
perc
0.8
0.6
0.4
0.2
ENST00000284551.11
ENST00000493030.6
9400
9600
9800
10000
IL2_UP.V1_UP
sampleType
LTL331_Cx_12w
LTL331_Cx_8w
LTL331_Cx_R
LTL331_PreCx

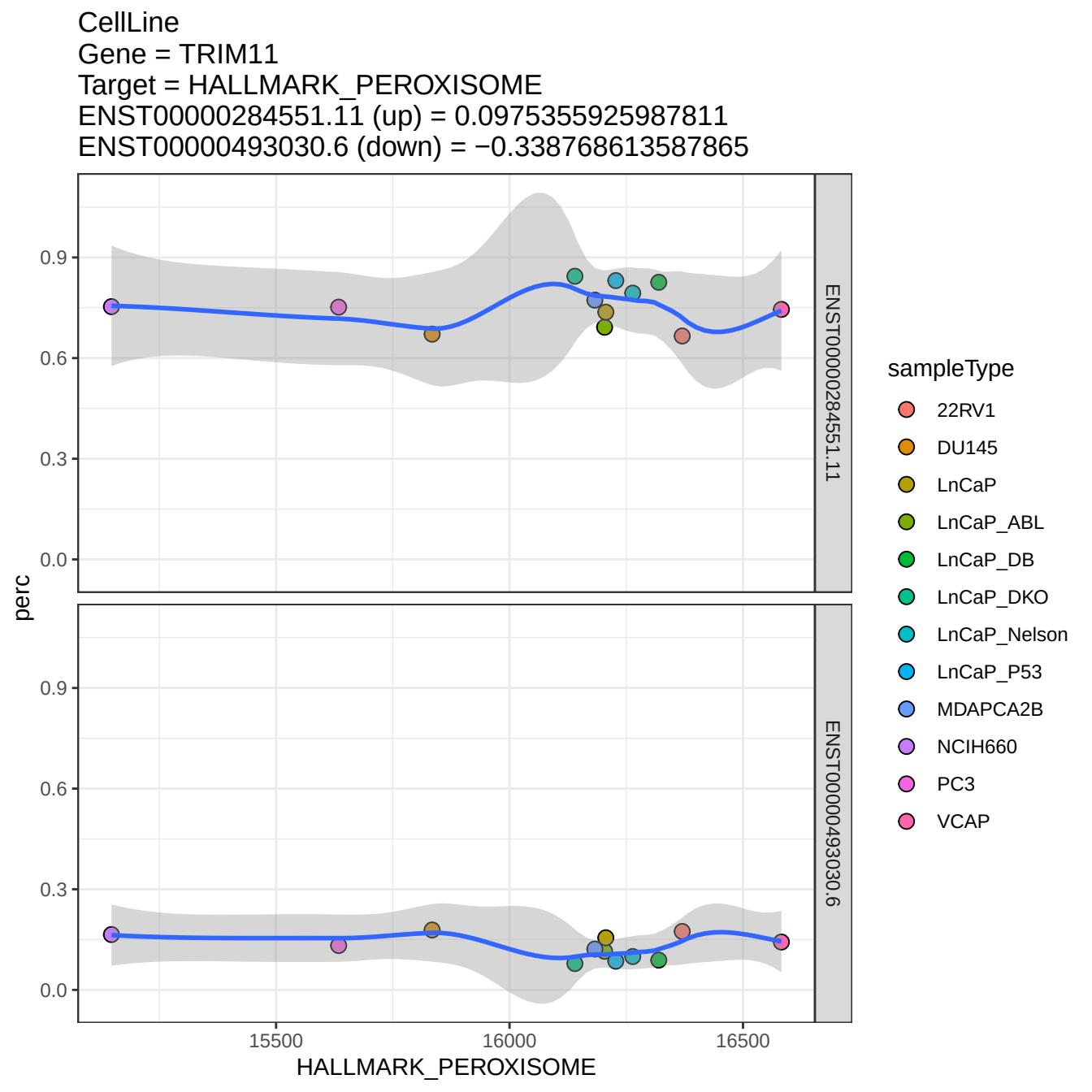

PDX_longitudinal
Gene = TRIM11
Target = IL2_UP.V1_UP
ENST00000284551.11 (up) = 0.45769337493359
ENST00000493030.6 (down) = −0.316406339363148
perc
1.00
0.75
0.50
0.25
0.00
ENST00000284551.11
ENST00000493030.6
10000
10250
10500
10750
IL2_UP.V1_UP
sampleType
LnCaP
LnCaP_ABL
LnCaP_DKO_CX
LnCaP_DKO_ENZA
LnCaP_DKO_PreCX
LuCaP_147
LuCaP_147_CX

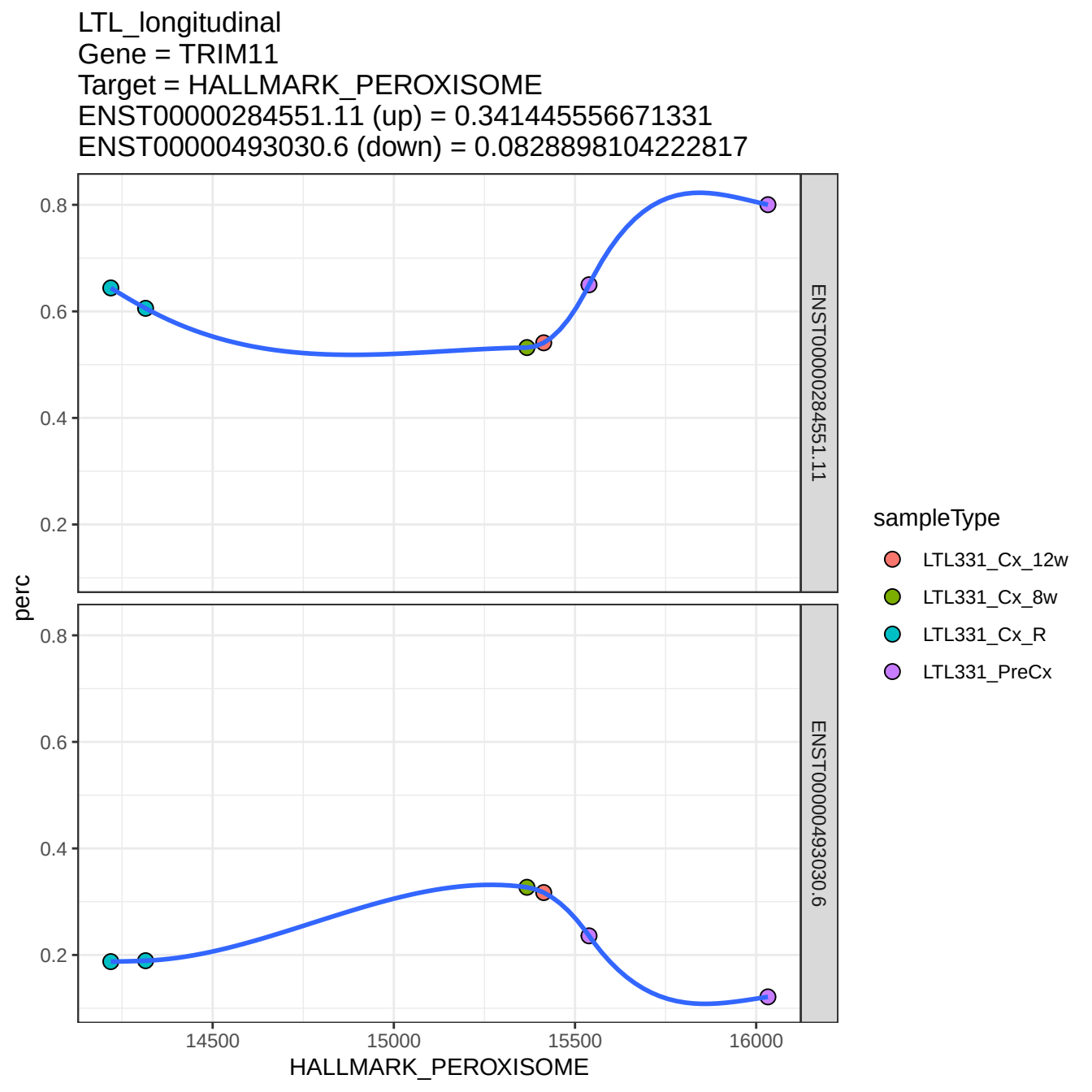

CellLine
Gene = TRIM11
Target = HALLMARK_P53_PATHWAY
ENST00000284551.11 (up) = −0.376186905422724
ENST00000493030.6 (down) = 0.126015875184004
perc
0.00
0.25
0.50
0.75
ENST00000284551.11
ENST00000493030.6
15600
16000
16400
16800
HALLMARK_P53_PATHWAY
sampleType
22RV1
DU145
LnCaP
LnCaP_ABL
LnCaP_DB
LnCaP_DKO
LnCaP_Nelson
LnCaP_P53
MDAPCA2B
NCIH660
PC3
VCAP


LTL_longitudinal
Gene = TRIM11
Target = HALLMARK_P53_PATHWAY
ENST00000284551.11 (up) = 0.0106287350327522
ENST00000493030.6 (down) = 0.389710060728891
perc
0.0
0.2
0.4
0.6
0.8
ENST00000284551.11
ENST00000493030.6
14000
14500
15000
15500
HALLMARK_P53_PATHWAY
sampleType
LTL331_Cx_12w
LTL331_Cx_8w
LTL331_Cx_R
LTL331_PreCx

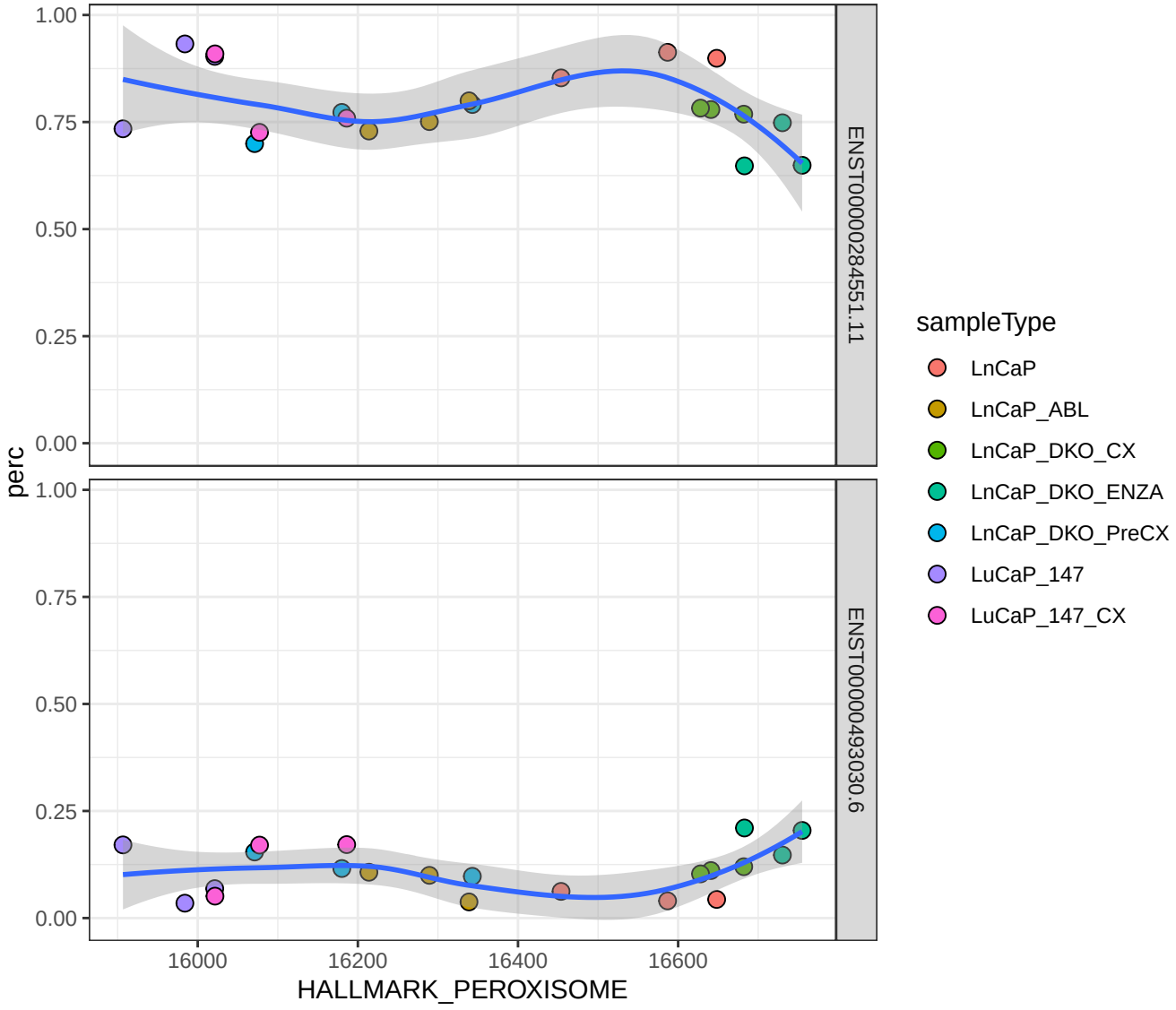
PDX_longitudinal
Gene = TRIM11
Target = HALLMARK_P53_PATHWAY
ENST00000284551.11 (up) = 0.656535863402131
ENST00000493030.6 (down) = −0.455071154611125

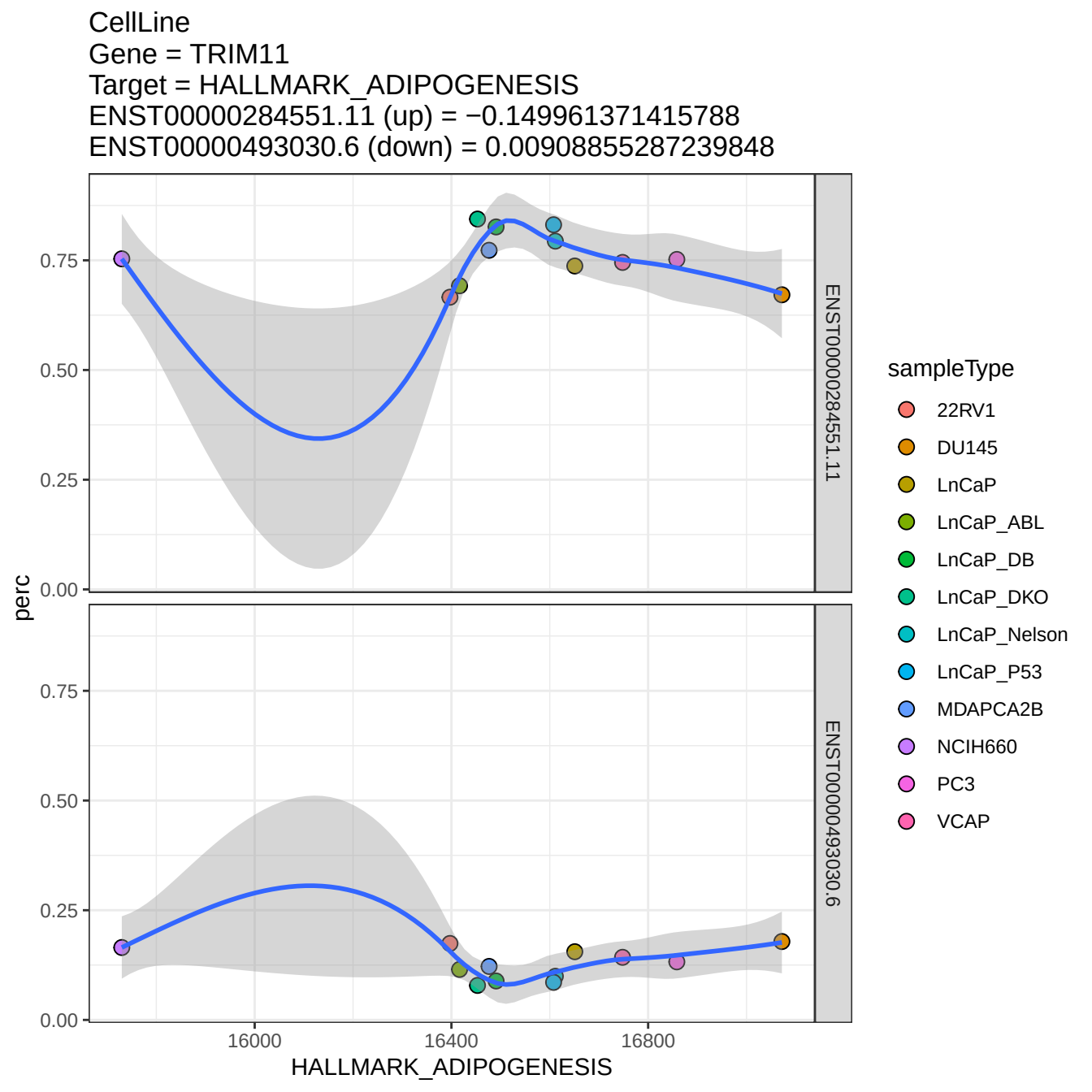


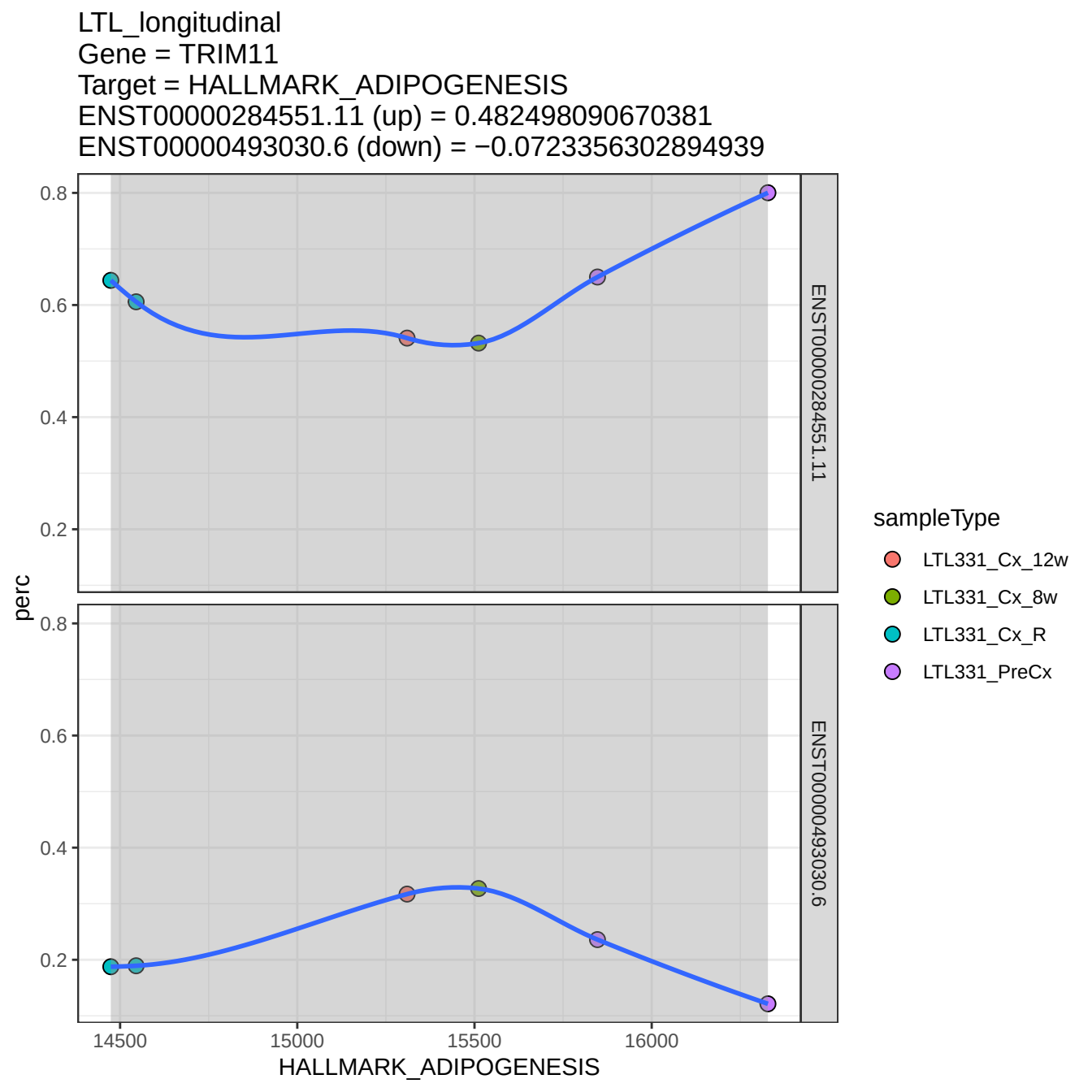

CellLine
Gene = TRIM11
Target = HALLMARK_PEROXISOME
ENST00000284551.11 (up) = 0.0975355925987811
ENST00000493030.6 (down) = −0.338768613587865
perc
0.0
0.3
0.6
0.9
ENST00000284551.11
ENST00000493030.6
15500
16000
16500
HALLMARK_PEROXISOME
sampleType
22RV1
DU145
LnCaP
LnCaP_ABL
LnCaP_DB
LnCaP_DKO
LnCaP_Nelson
LnCaP_P53
MDAPCA2B
NCIH660
PC3
VCAP

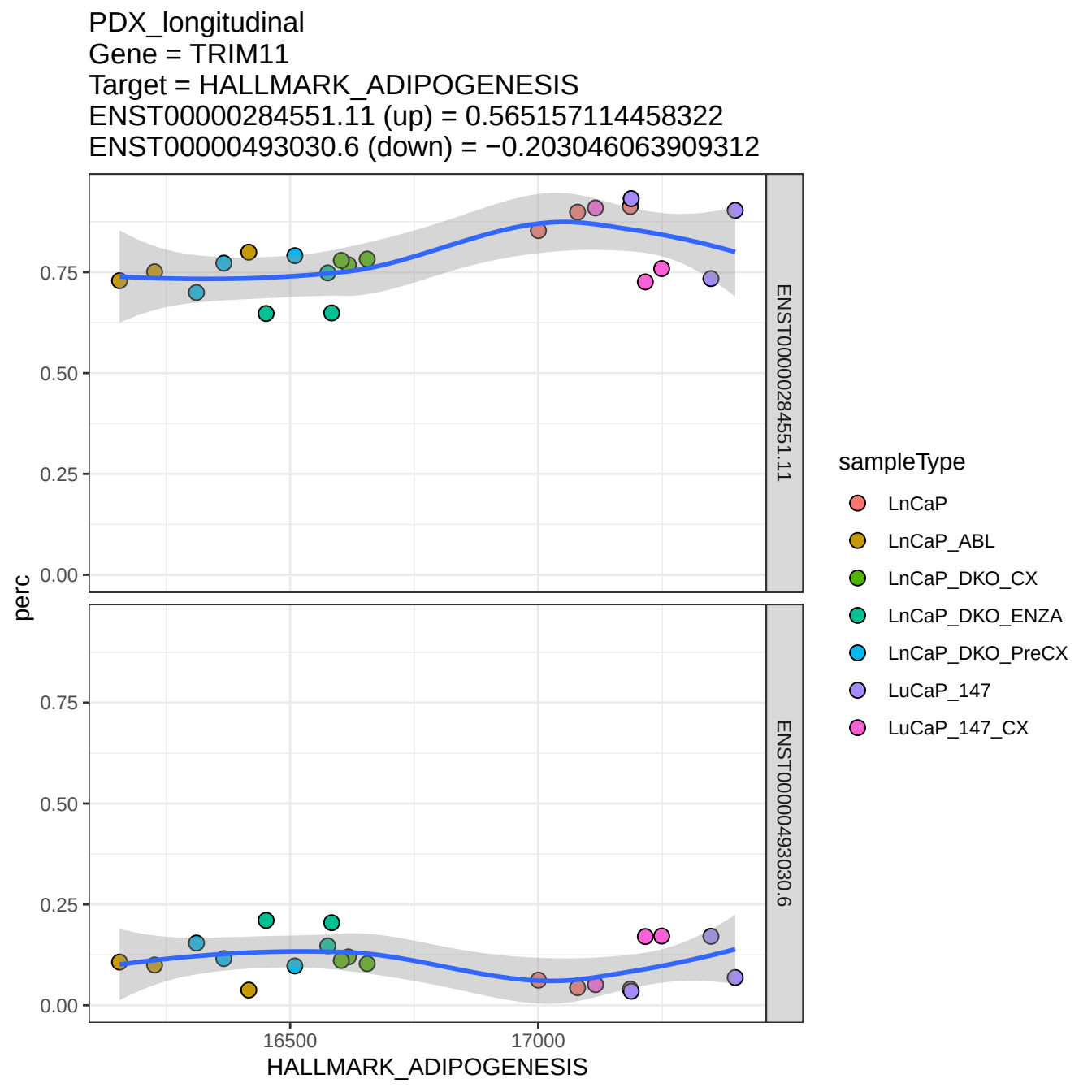

LTL_longitudinal
Gene = TRIM11
Target = HALLMARK_PEROXISOME
ENST00000284551.11 (up) = 0.341445556671331
ENST00000493030.6 (down) = 0.0828898104222817
perc
0.8
0.6
0.4
0.2
ENST00000284551.11
ENST00000493030.6
14500
15000
15500
16000
HALLMARK_PEROXISOME
sampleType
LTL331_Cx_12w
LTL331_Cx_8w
LTL331_Cx_R
LTL331_PreCx

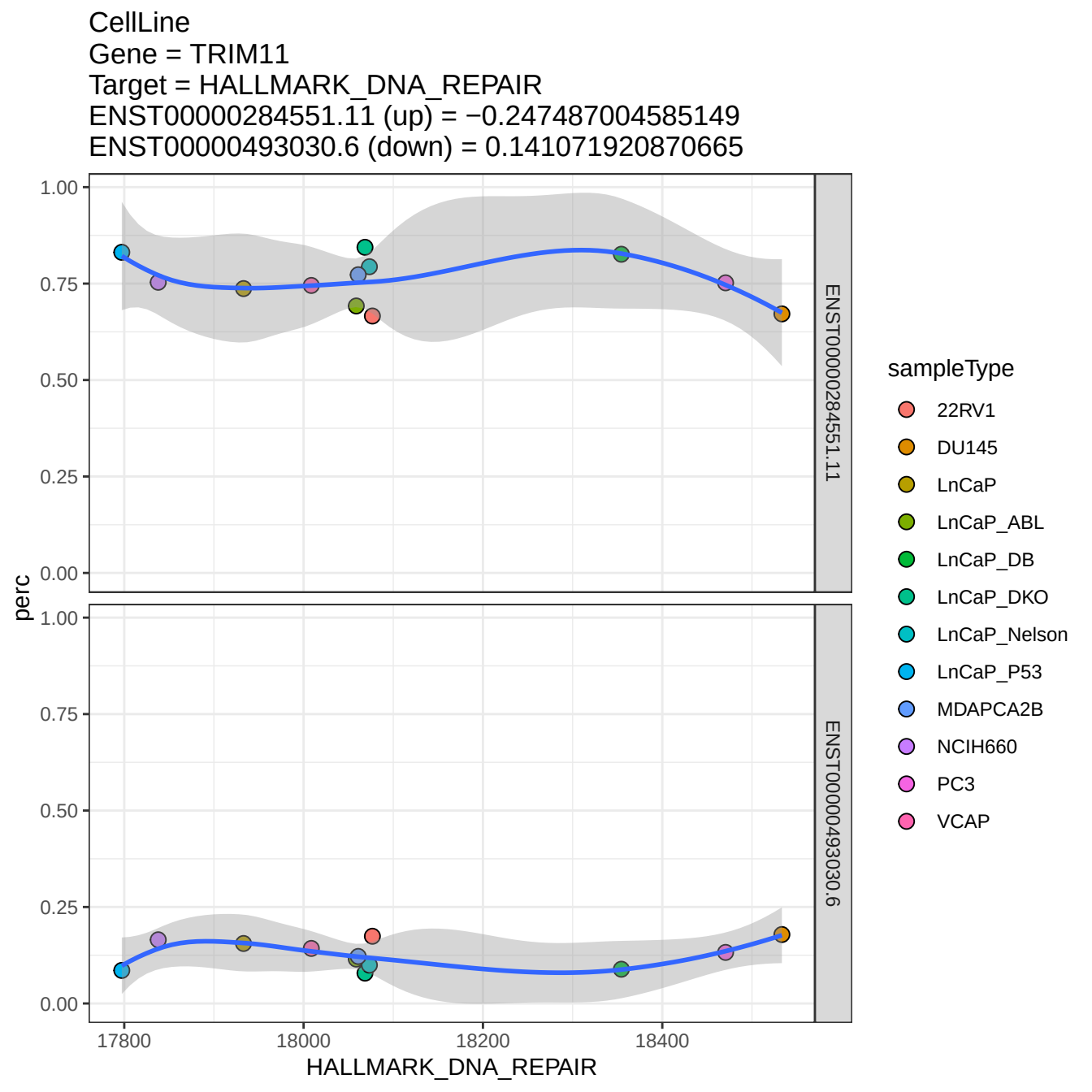
PDX_longitudinal
Gene = TRIM11
Target = HALLMARK_PEROXISOME
ENST00000284551.11 (up) = −0.222075205261521
ENST00000493030.6 (down) = 0.131712880648737

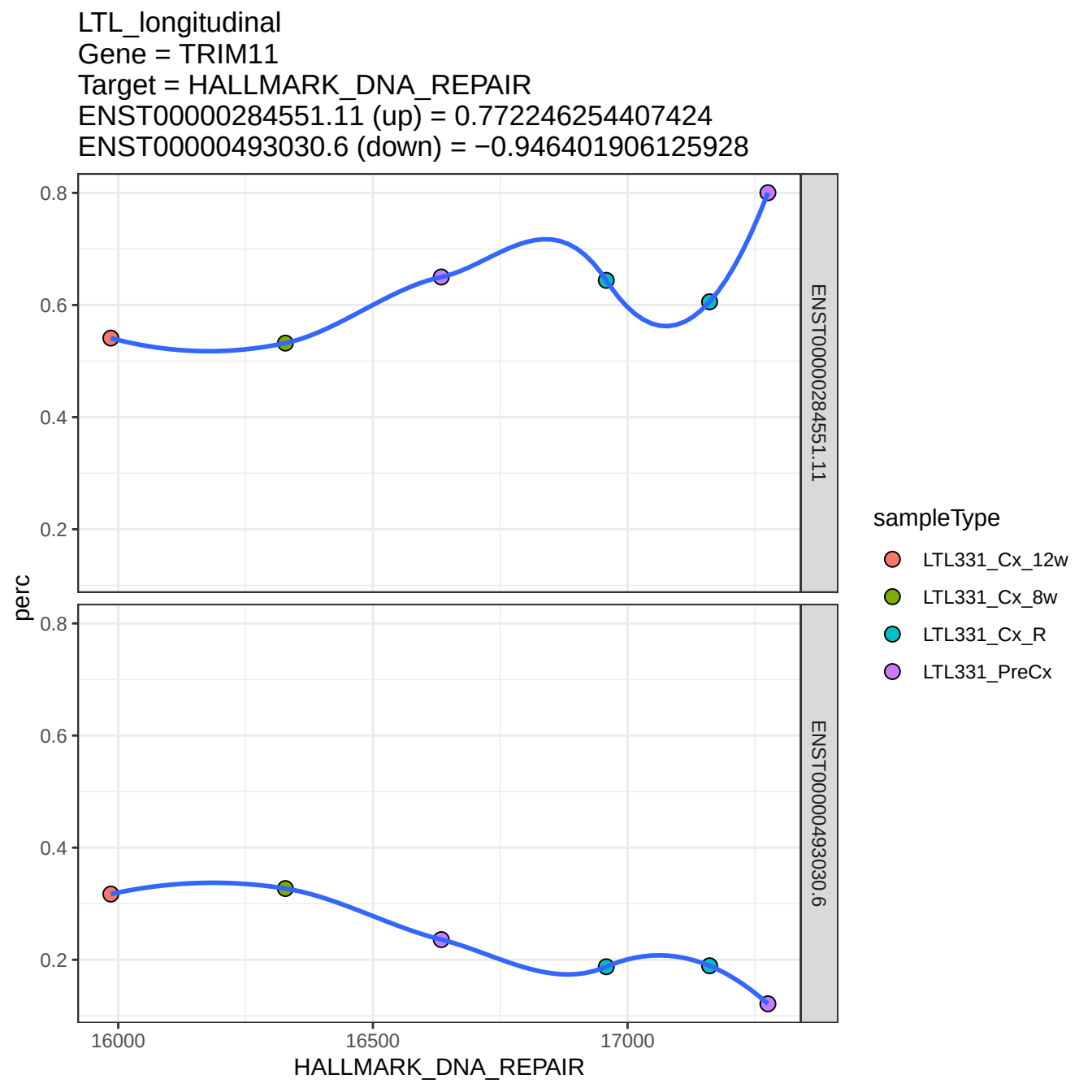


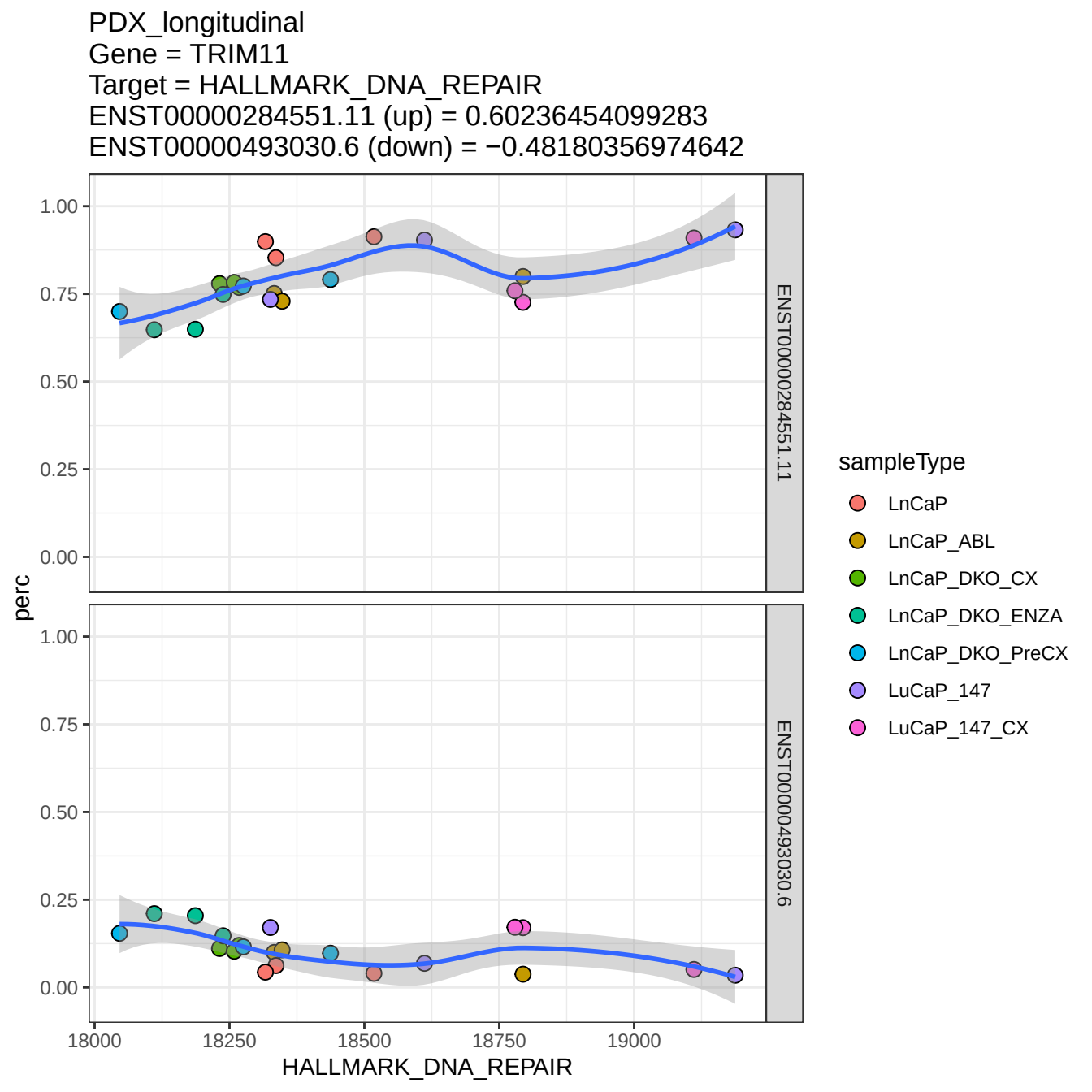

CellLine
Gene = TRIM11
Target = HALLMARK_ADIPOGENESIS
ENST00000284551.11 (up) = −0.149961371415788
ENST00000493030.6 (down) = 0.00908855287239848
perc
0.00
0.25
0.50
0.75
ENST00000284551.11
ENST00000493030.6
16000
16400
16800
HALLMARK_ADIPOGENESIS
sampleType
22RV1
DU145
LnCaP
LnCaP_ABL
LnCaP_DB
LnCaP_DKO
LnCaP_Nelson
LnCaP_P53
MDAPCA2B
NCIH660
PC3
VCAP

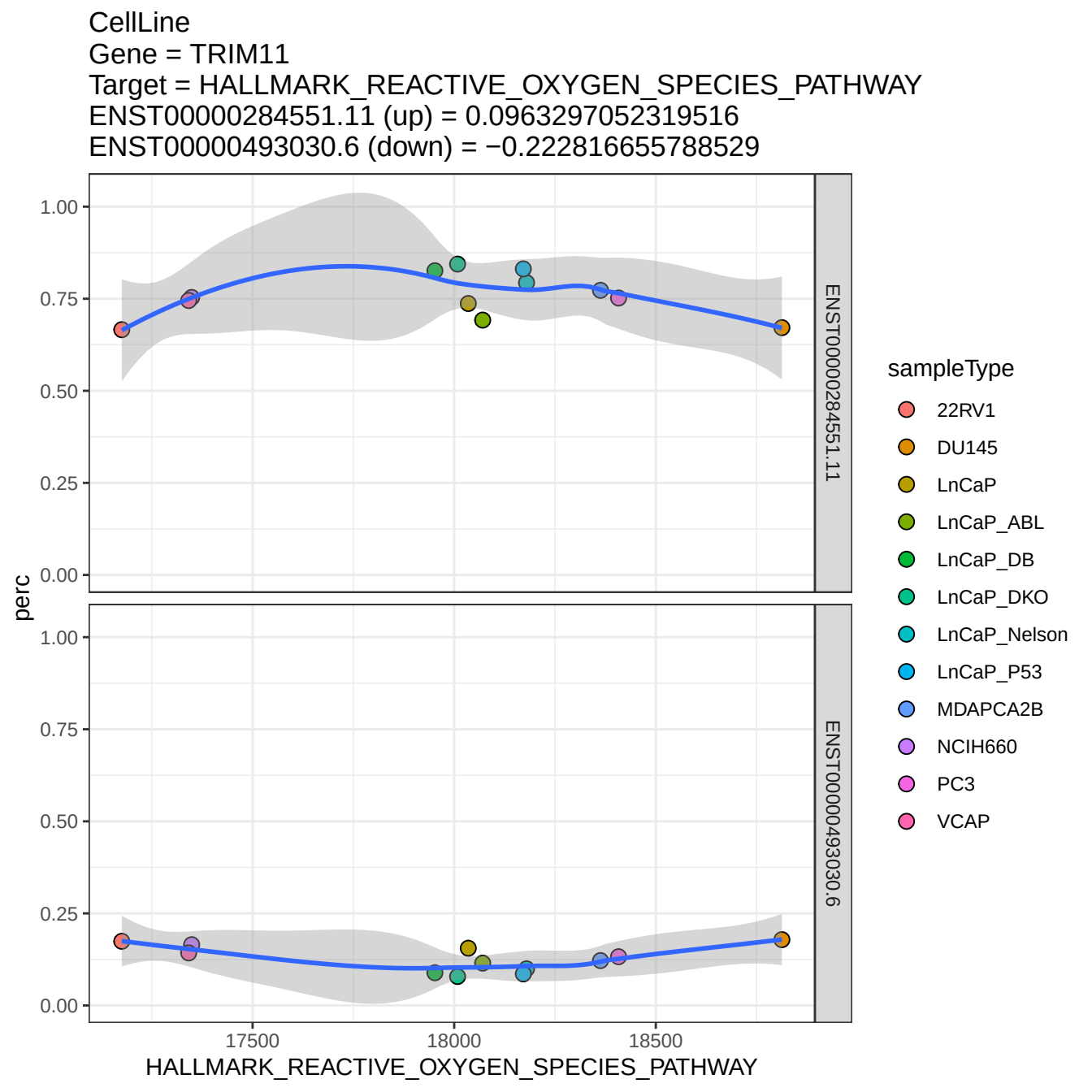

LTL_longitudinal
Gene = TRIM11
Target = HALLMARK_ADIPOGENESIS
ENST00000284551.11 (up) = 0.482498090670381
ENST00000493030.6 (down) = −0.0723356302894939
perc
0.8
0.6
0.4
0.2
ENST00000284551.11
ENST00000493030.6
14500
15000
15500
16000
HALLMARK_ADIPOGENESIS
sampleType
LTL331_Cx_12w
LTL331_Cx_8w
LTL331_Cx_R
LTL331_PreCx

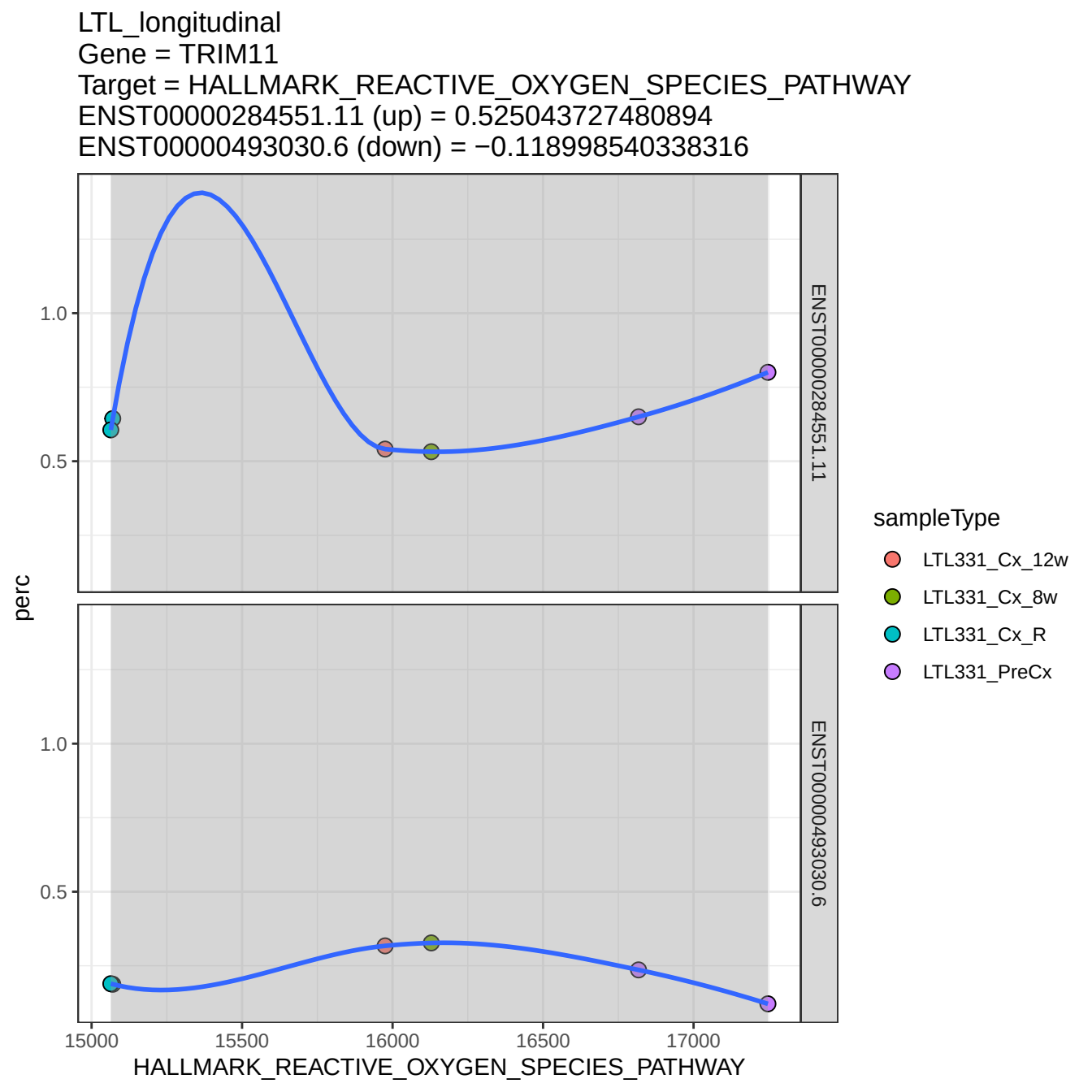

PDX_longitudinal
Gene = TRIM11
Target = HALLMARK_ADIPOGENESIS
ENST00000284551.11 (up) = 0.565157114458322
ENST00000493030.6 (down) = −0.203046063909312
perc
0.00
0.25
0.50
0.75
ENST00000284551.11
ENST00000493030.6
16500
17000
HALLMARK_ADIPOGENESIS
sampleType
LnCaP
LnCaP_ABL
LnCaP_DKO_CX
LnCaP_DKO_ENZA
LnCaP_DKO_PreCX
LuCaP_147
LuCaP_147_CX


CellLine
Gene = TRIM11
Target = HALLMARK_DNA_REPAIR
ENST00000284551.11 (up) = −0.247487004585149
ENST00000493030.6 (down) = 0.141071920870665
perc
1.00
0.75
0.50
0.25
0.00
ENST00000284551.11
ENST00000493030.6
17800
18000
18200
18400
HALLMARK_DNA_REPAIR
sampleType
22RV1
DU145
LnCaP
LnCaP_ABL
LnCaP_DB
LnCaP_DKO
LnCaP_Nelson
LnCaP_P53
MDAPCA2B
NCIH660
PC3
VCAP

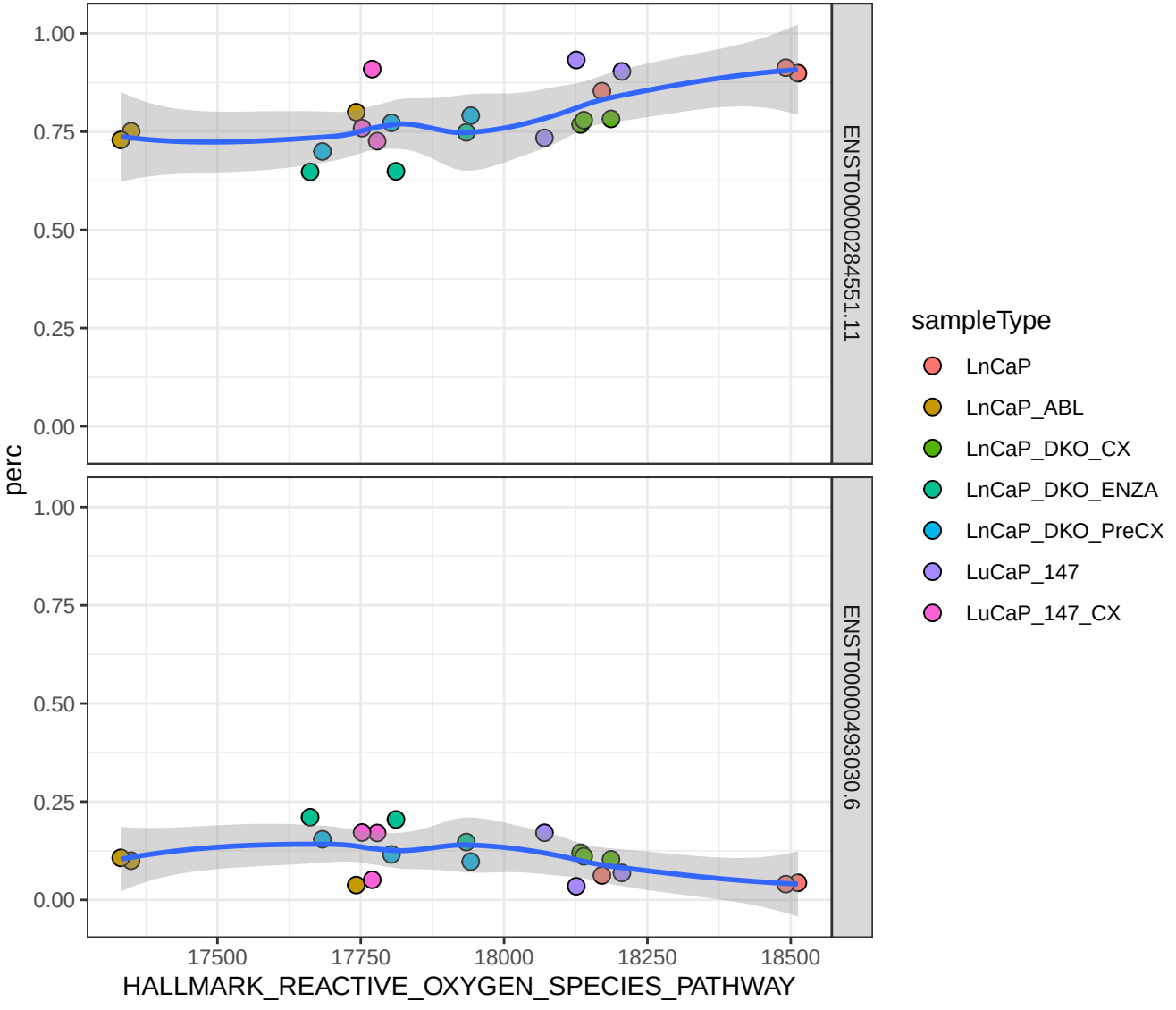

LTL_longitudinal
Gene = TRIM11
Target = HALLMARK_DNA_REPAIR
ENST00000284551.11 (up) = 0.772246254407424
ENST00000493030.6 (down) = −0.946401906125928
perc
0.8
0.6
0.4
0.2
ENST00000284551.11
ENST00000493030.6
16000
16500
17000
HALLMARK_DNA_REPAIR
sampleType
LTL331_Cx_12w
LTL331_Cx_8w
LTL331_Cx_R
LTL331_PreCx

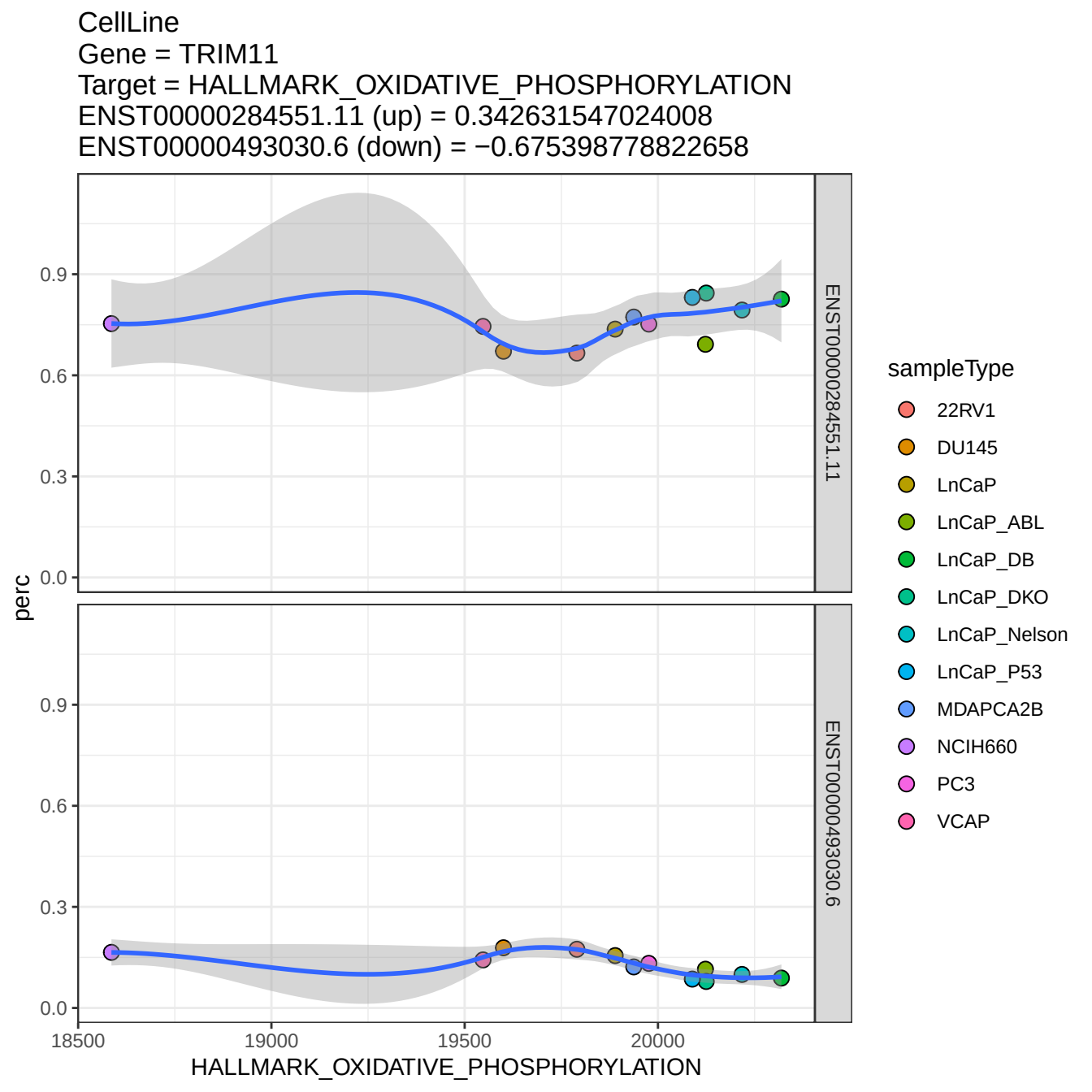

PDX_longitudinal
Gene = TRIM11
Target = HALLMARK_DNA_REPAIR
ENST00000284551.11 (up) = 0.60236454099283
ENST00000493030.6 (down) = −0.48180356974642
perc
1.00
0.75
0.50
0.25
0.00
ENST00000284551.11
ENST00000493030.6
18000
18250
18500
18750
19000
HALLMARK_DNA_REPAIR
sampleType
LnCaP
LnCaP_ABL
LnCaP_DKO_CX
LnCaP_DKO_ENZA
LnCaP_DKO_PreCX
LuCaP_147
LuCaP_147_CX


CellLine
Gene = TRIM11
Target = HALLMARK_REACTIVE_OXYGEN_SPECIES_PATHWAY
ENST00000284551.11 (up) = 0.0963297052319516
ENST00000493030.6 (down) = −0.222816655788529
perc
1.00
0.75
0.50
0.25
0.00
ENST00000284551.11
ENST00000493030.6
17500
18000
18500
HALLMARK_REACTIVE_OXYGEN_SPECIES_PATHWAY
sampleType
22RV1
DU145
LnCaP
LnCaP_ABL
LnCaP_DB
LnCaP_DKO
LnCaP_Nelson
LnCaP_P53
MDAPCA2B
NCIH660
PC3
VCAP

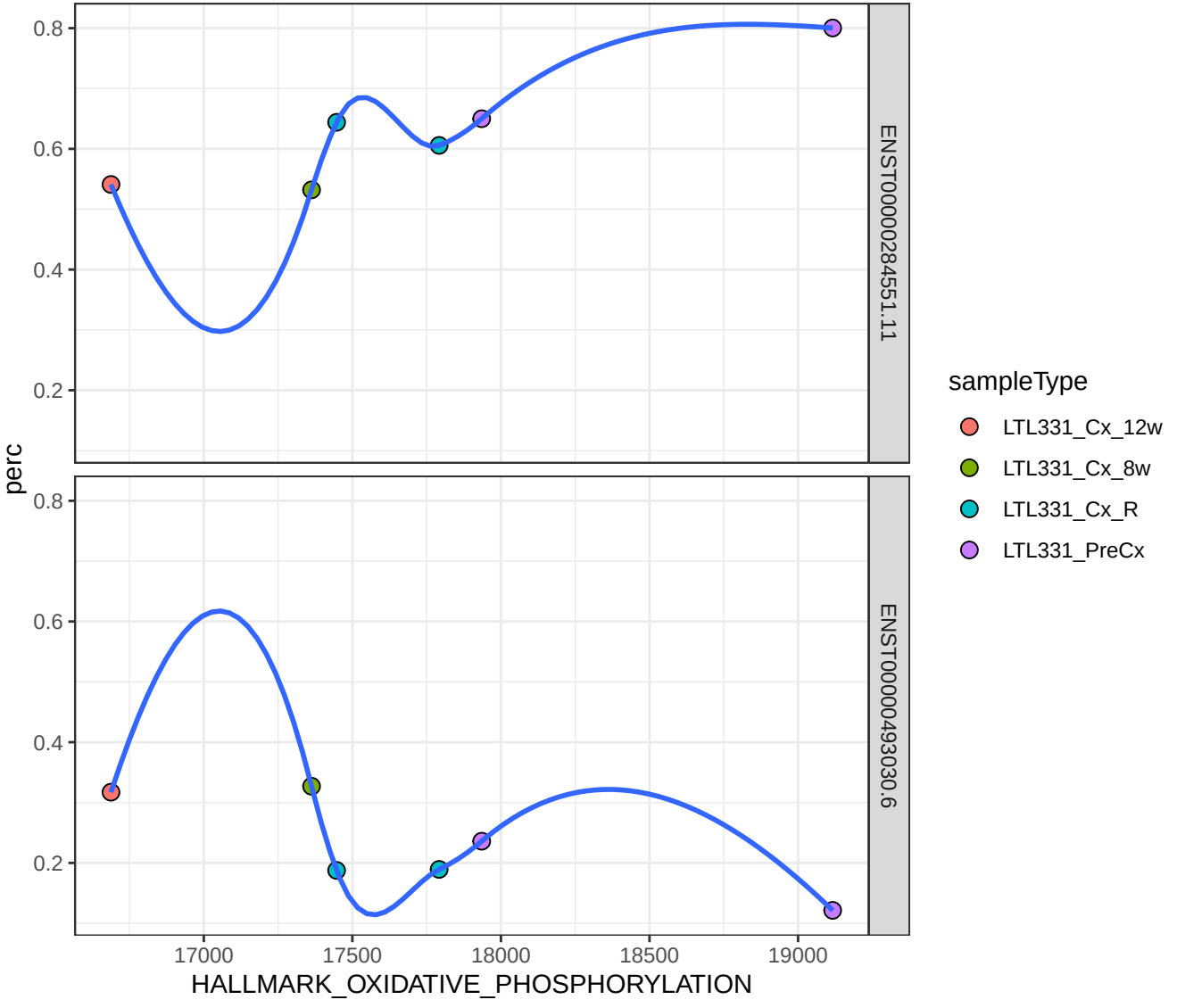

LTL_longitudinal
Gene = TRIM11
Target = HALLMARK_REACTIVE_OXYGEN_SPECIES_PATHWAY
ENST00000284551.11 (up) = 0.525043727480894
ENST00000493030.6 (down) = −0.118998540338316
ENST00000284551.11
ENST00000493030.6
perc
1.0
0.5
15000
15500
16000
16500
17000
HALLMARK_REACTIVE_OXYGEN_SPECIES_PATHWAY
sampleType
LTL331_Cx_12w
LTL331_Cx_8w
LTL331_Cx_R
LTL331_PreCx

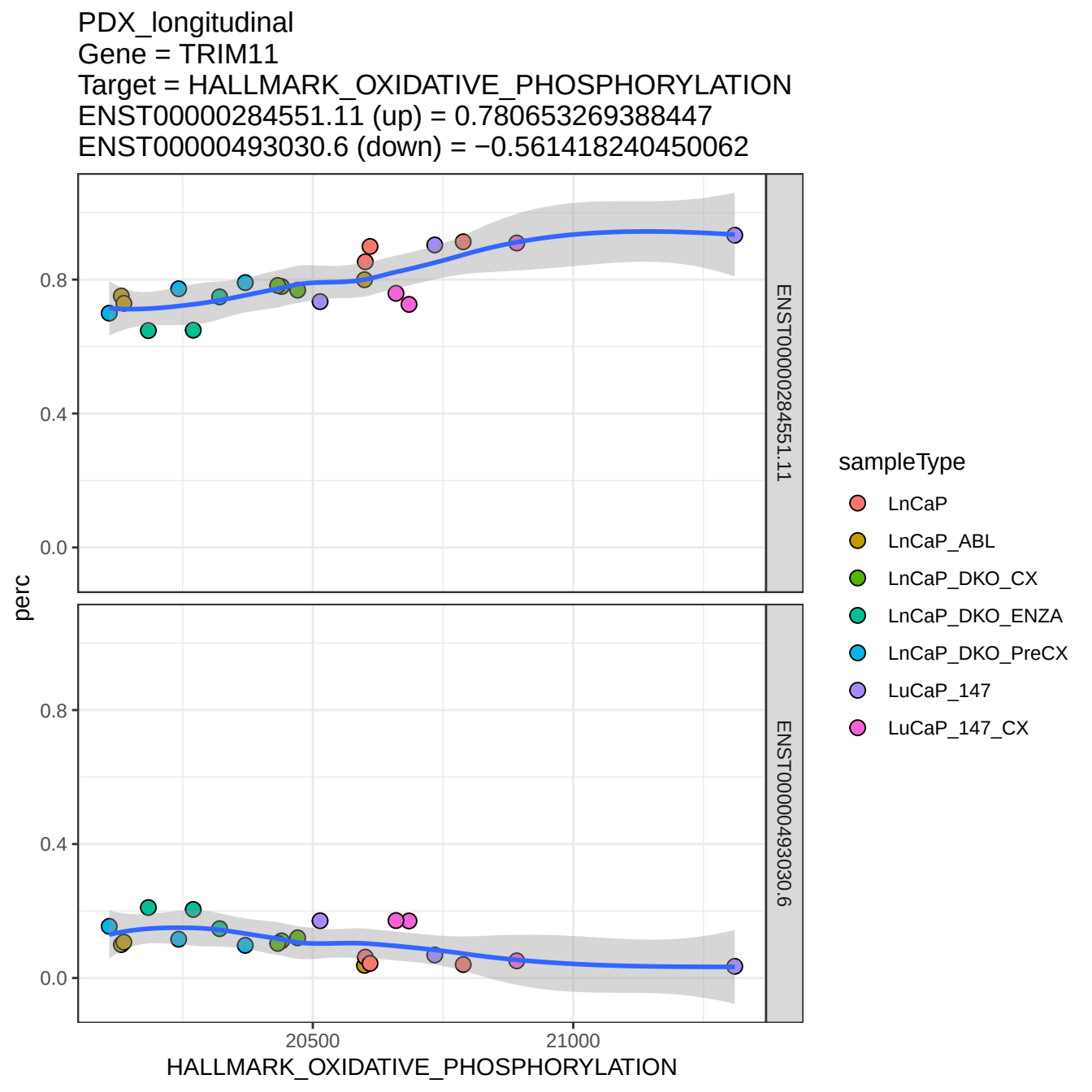
PDX_longitudinal
Gene = TRIM11
Target = HALLMARK_REACTIVE_OXYGEN_SPECIES_PATHWAY
ENST00000284551.11 (up) = 0.609499926198968
ENST00000493030.6 (down) = −0.418212017455003

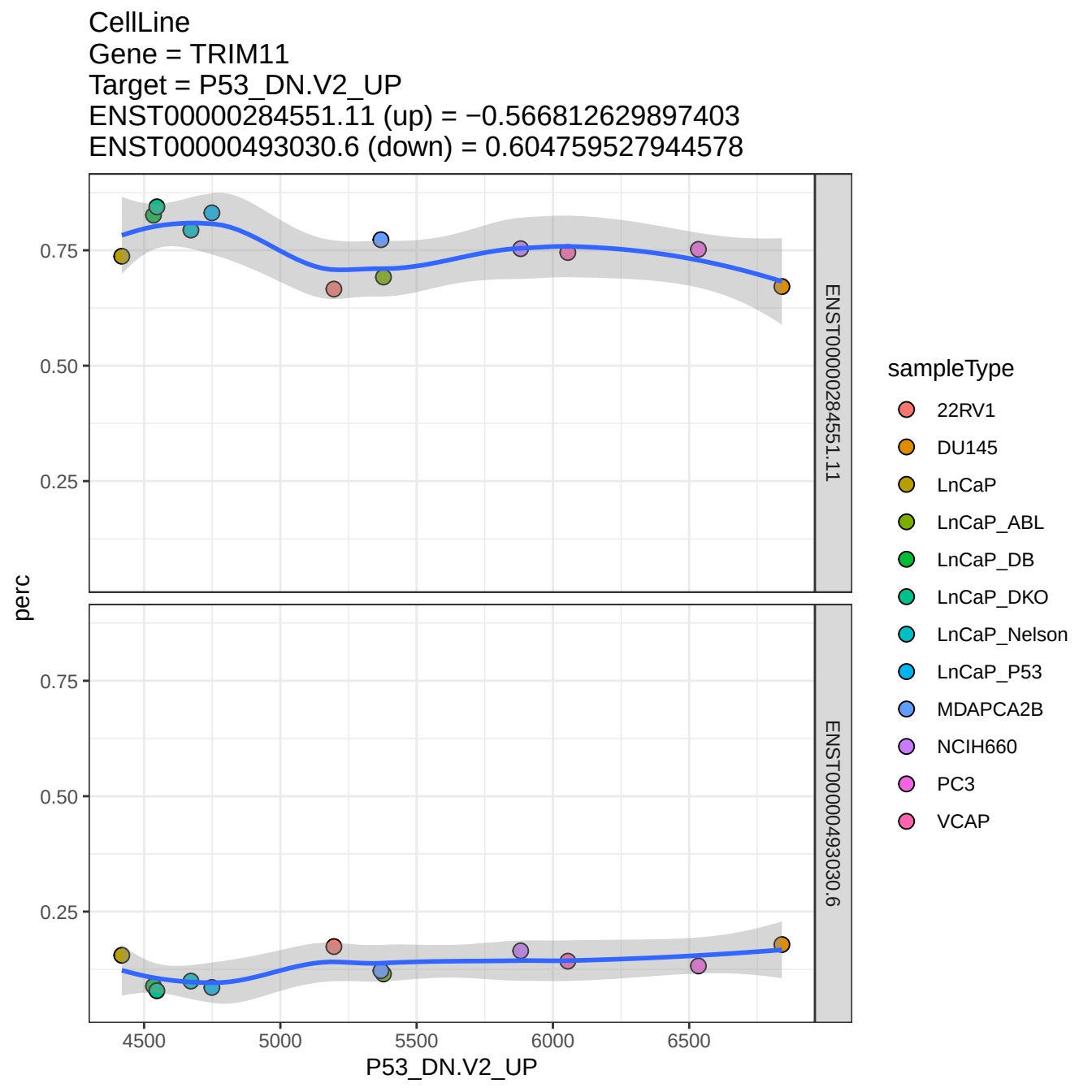


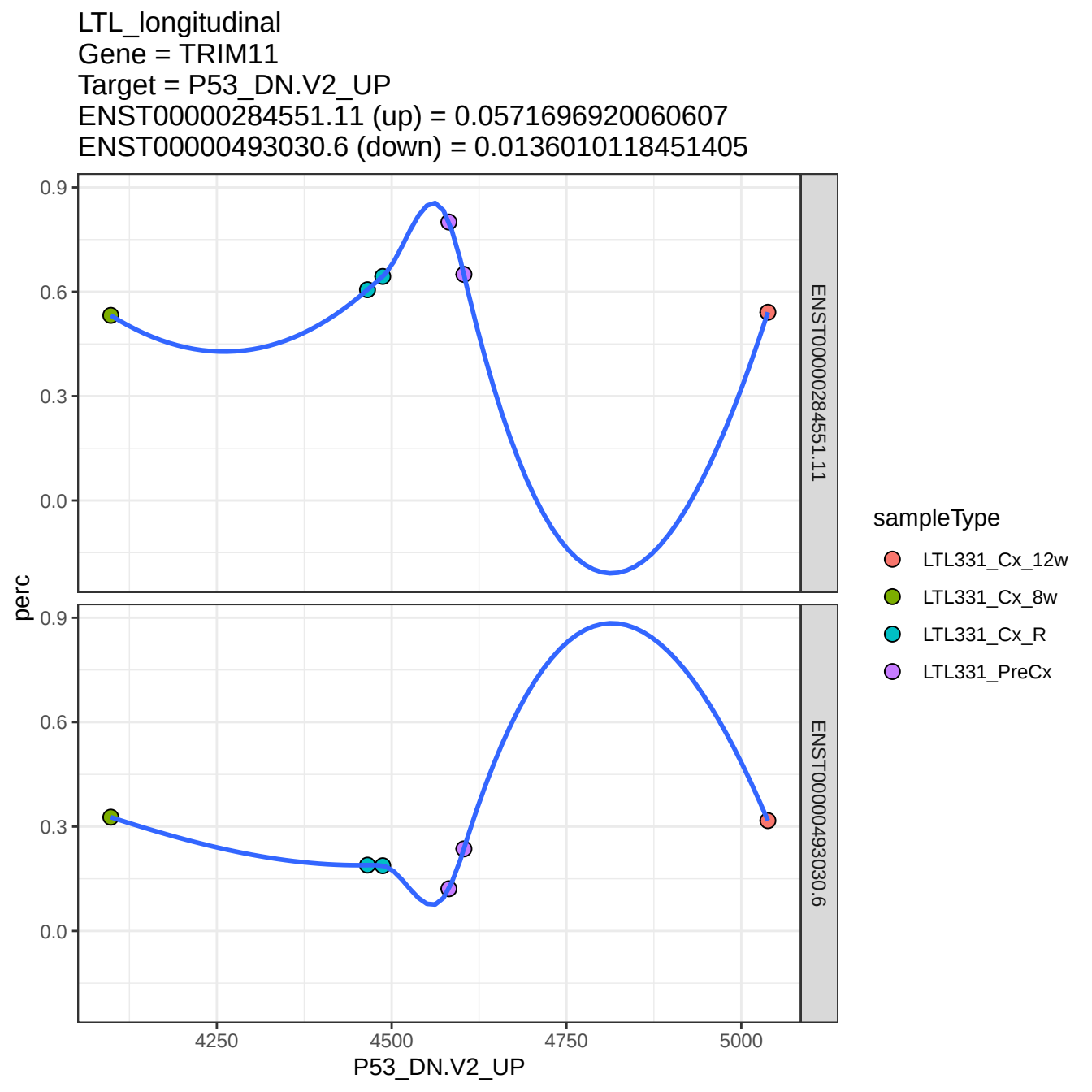

CellLine
Gene = TRIM11
Target = HALLMARK_OXIDATIVE_PHOSPHORYLATION
ENST00000284551.11 (up) = 0.342631547024008
ENST00000493030.6 (down) = −0.675398778822658
perc
0.0
0.3
0.6
0.9
ENST00000284551.11
ENST00000493030.6
18500
19000
19500
20000
HALLMARK_OXIDATIVE_PHOSPHORYLATION
sampleType
22RV1
DU145
LnCaP
LnCaP_ABL
LnCaP_DB
LnCaP_DKO
LnCaP_Nelson
LnCaP_P53
MDAPCA2B
NCIH660
PC3
VCAP

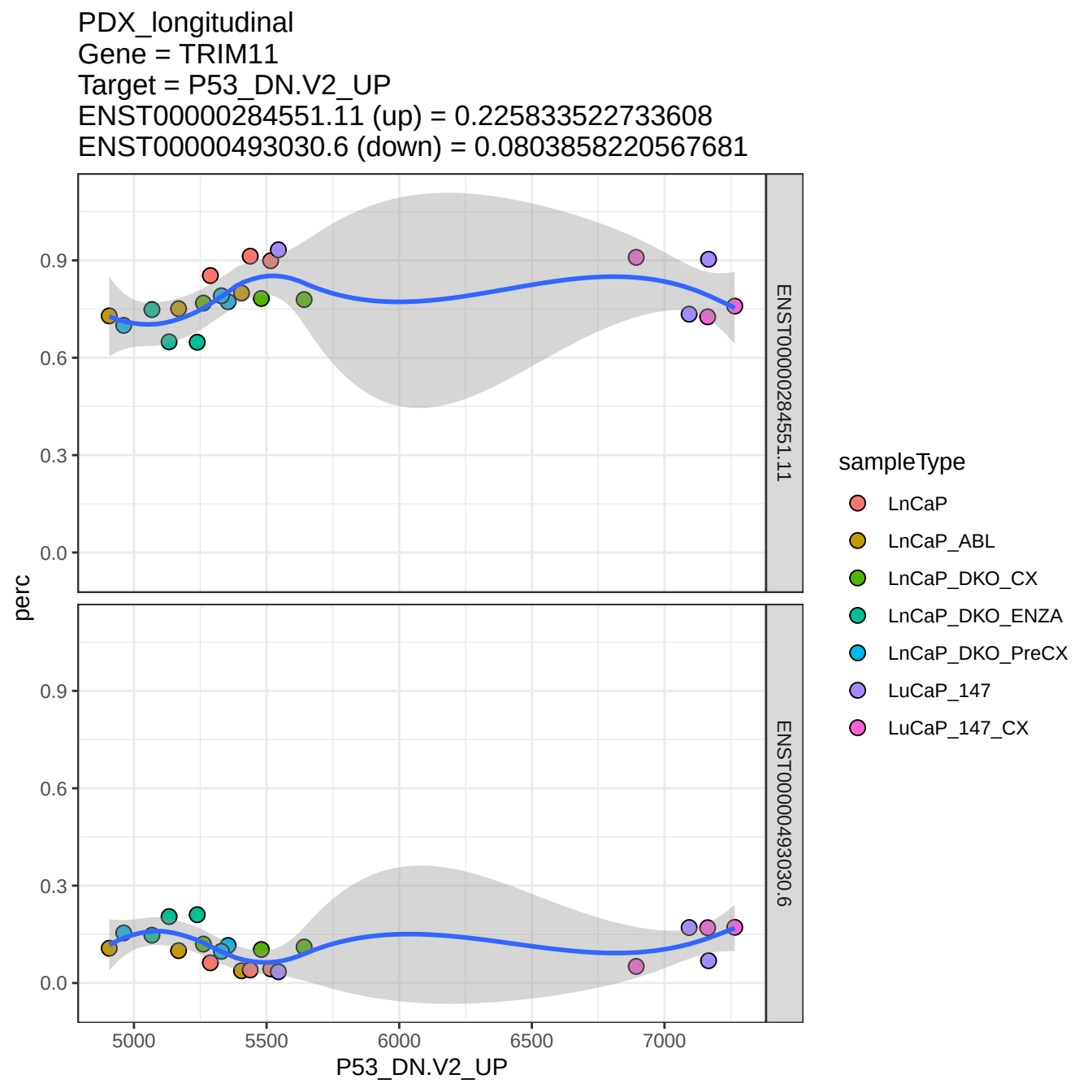
LTL_longitudinal
Gene = TRIM11
Target = HALLMARK_OXIDATIVE_PHOSPHORYLATION
ENST00000284551.11 (up) = 0.920406451537823
ENST00000493030.6 (down) = −0.819642661565318

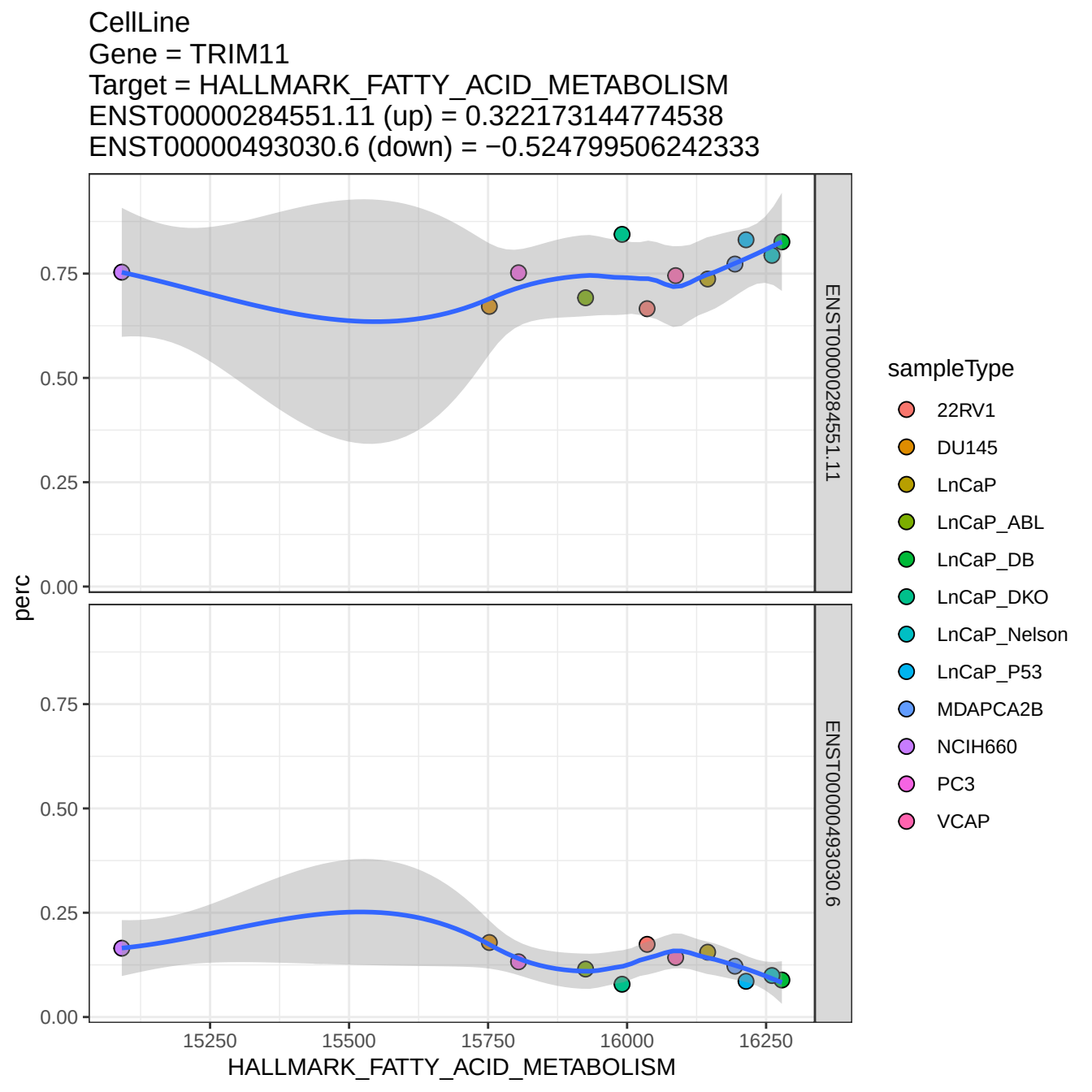


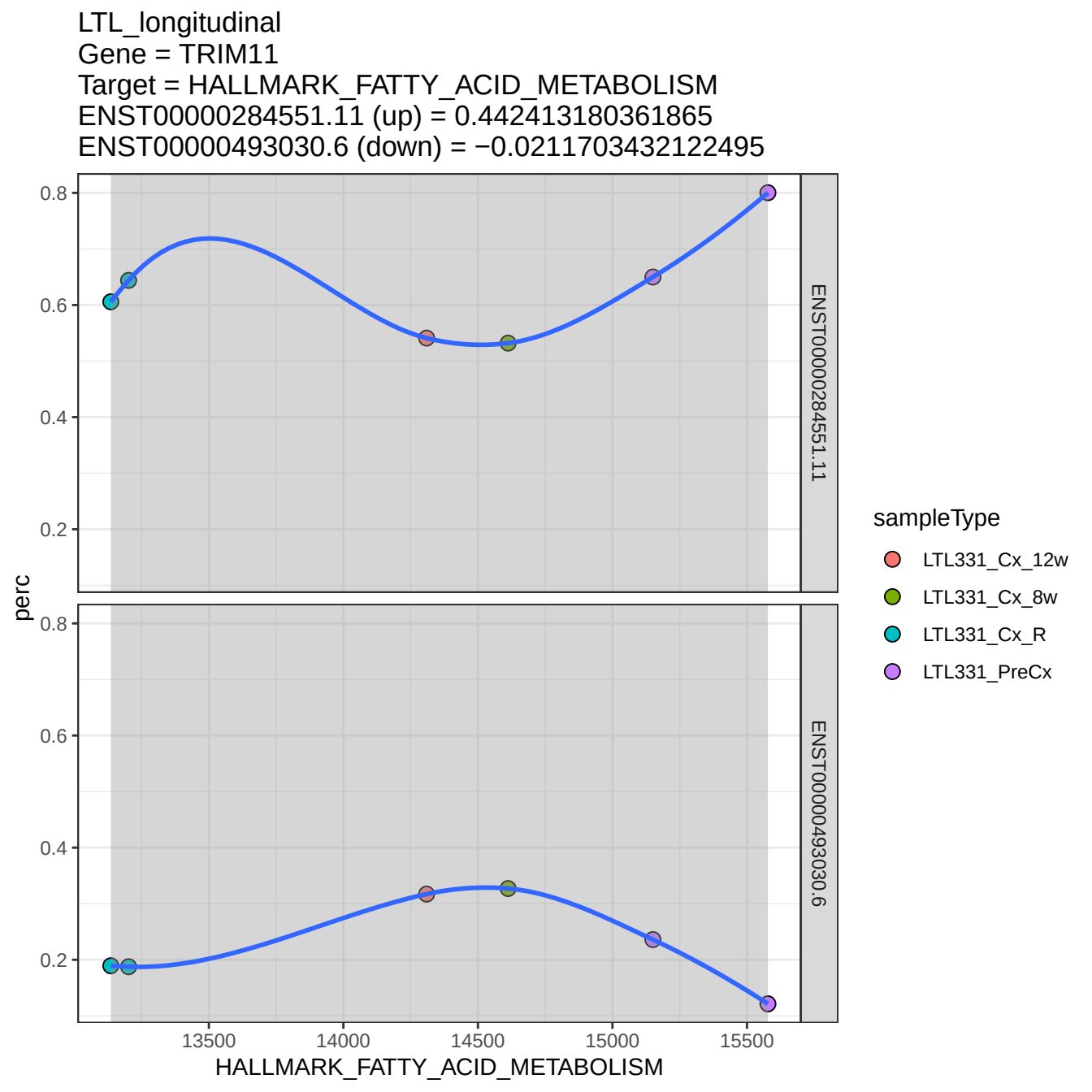

PDX_longitudinal
Gene = TRIM11
Target = HALLMARK_OXIDATIVE_PHOSPHORYLATION
ENST00000284551.11 (up) = 0.780653269388447
ENST00000493030.6 (down) = −0.561418240450062
perc
0.0
0.4
0.8
ENST00000284551.11
ENST00000493030.6
20500
21000
HALLMARK_OXIDATIVE_PHOSPHORYLATION
sampleType
LnCaP
LnCaP_ABL
LnCaP_DKO_CX
LnCaP_DKO_ENZA
LnCaP_DKO_PreCX
LuCaP_147
LuCaP_147_CX

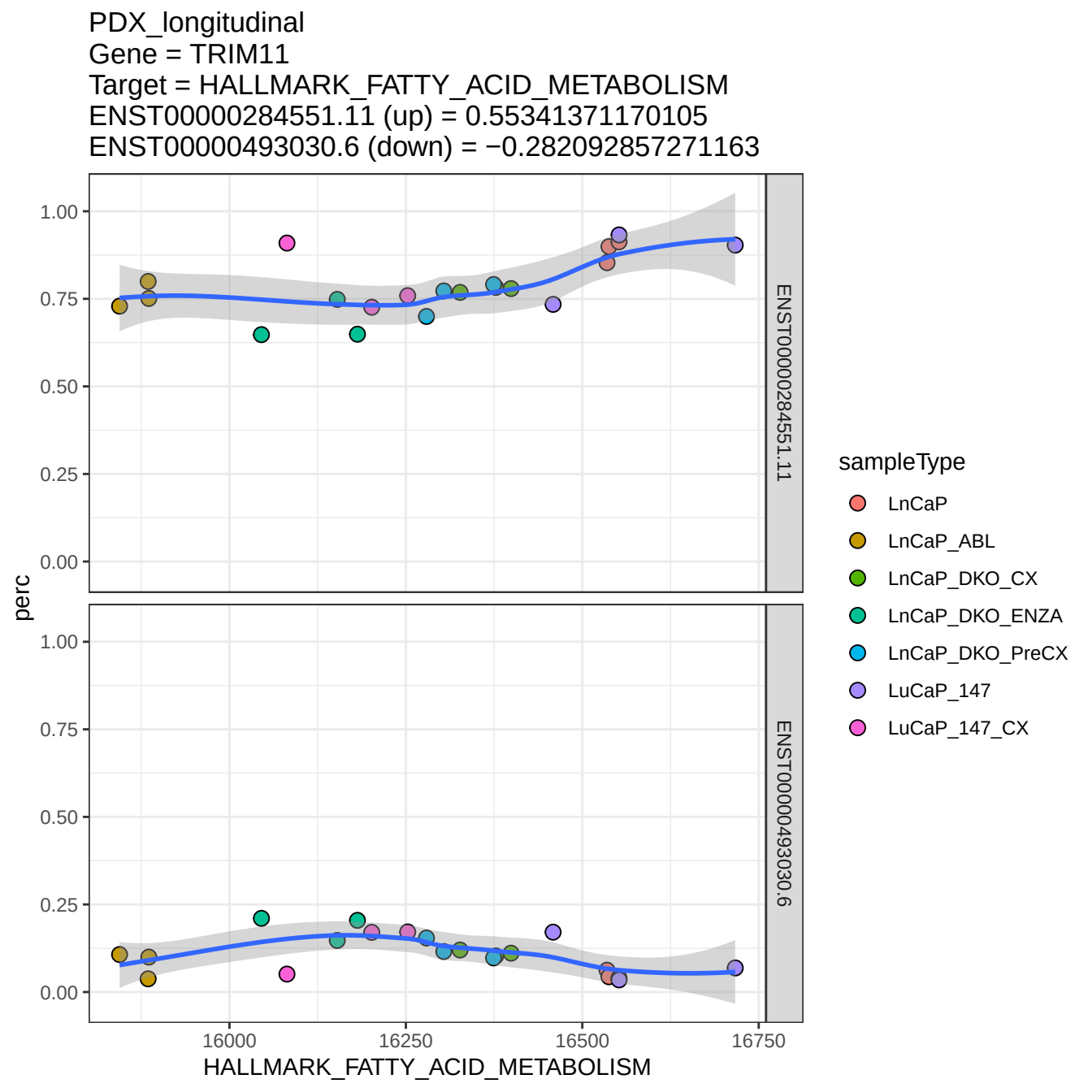

CellLine
Gene = TRIM11
Target = P53_DN.V2_UP
ENST00000284551.11 (up) = −0.566812629897403
ENST00000493030.6 (down) = 0.604759527944578
perc
0.75
0.50
0.25
ENST00000284551.11
ENST00000493030.6
4500
5000
5500
6000
6500
P53_DN.V2_UP
sampleType
22RV1
DU145
LnCaP
LnCaP_ABL
LnCaP_DB
LnCaP_DKO
LnCaP_Nelson
LnCaP_P53
MDAPCA2B
NCIH660
PC3
VCAP


LTL_longitudinal
Gene = TRIM11
Target = P53_DN.V2_UP
ENST00000284551.11 (up) = 0.0571696920060607
ENST00000493030.6 (down) = 0.0136010118451405
perc
0.9
0.6
0.3
0.0
ENST00000284551.11
ENST00000493030.6
4250
4500
4750
5000
P53_DN.V2_UP
sampleType
LTL331_Cx_12w
LTL331_Cx_8w
LTL331_Cx_R
LTL331_PreCx


PDX_longitudinal
Gene = TRIM11
Target = P53_DN.V2_UP
ENST00000284551.11 (up) = 0.225833522733608
ENST00000493030.6 (down) = 0.0803858220567681
perc
0.0
0.3
0.6
0.9
ENST00000284551.11
ENST00000493030.6
5000
5500
6000
6500
7000
P53_DN.V2_UP
sampleType
LnCaP
LnCaP_ABL
LnCaP_DKO_CX
LnCaP_DKO_ENZA
LnCaP_DKO_PreCX
LuCaP_147
LuCaP_147_CX


CellLine
Gene = TRIM11
Target = HALLMARK_FATTY_ACID_METABOLISM
ENST00000284551.11 (up) = 0.322173144774538
ENST00000493030.6 (down) = −0.524799506242333
perc
0.00
0.25
0.50
0.75
ENST00000284551.11
ENST00000493030.6
15250
15500
15750
16000
16250
HALLMARK_FATTY_ACID_METABOLISM
sampleType
22RV1
DU145
LnCaP
LnCaP_ABL
LnCaP_DB
LnCaP_DKO
LnCaP_Nelson
LnCaP_P53
MDAPCA2B
NCIH660
PC3
VCAP


LTL_longitudinal
Gene = TRIM11
Target = HALLMARK_FATTY_ACID_METABOLISM
ENST00000284551.11 (up) = 0.442413180361865
ENST00000493030.6 (down) = −0.0211703432122495
perc
0.8
0.6
0.4
0.2
ENST00000284551.11
ENST00000493030.6
13500
14000
14500
15000
15500
HALLMARK_FATTY_ACID_METABOLISM
sampleType
LTL331_Cx_12w
LTL331_Cx_8w
LTL331_Cx_R
LTL331_PreCx


PDX_longitudinal
Gene = TRIM11
Target = HALLMARK_FATTY_ACID_METABOLISM
ENST00000284551.11 (up) = 0.55341371170105
ENST00000493030.6 (down) = −0.282092857271163
perc
1.00
0.75
0.50
0.25
0.00
ENST00000284551.11
ENST00000493030.6
16000
16250
16500
16750
HALLMARK_FATTY_ACID_METABOLISM
sampleType
LnCaP
LnCaP_ABL
LnCaP_DKO_CX
LnCaP_DKO_ENZA
LnCaP_DKO_PreCX
LuCaP_147
LuCaP_147_CX