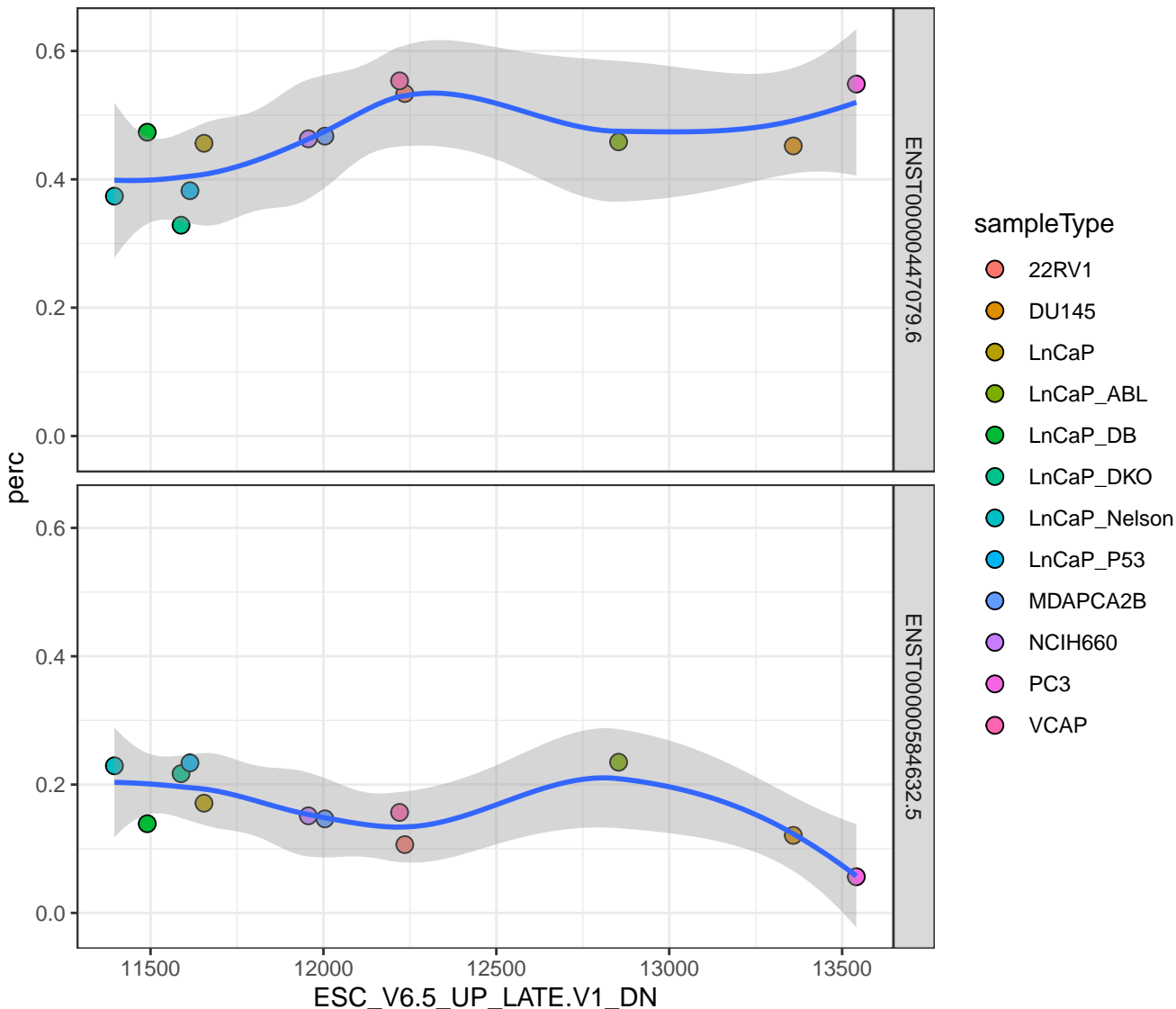


CellLine
 Gene = CDK12
 Target = ESC_V6.5_UP_LATE.V1_DN
 ENST00000447079.6 (up) = 0.535714148966806
 ENST00000584632.5 (down) = -0.567841507861841



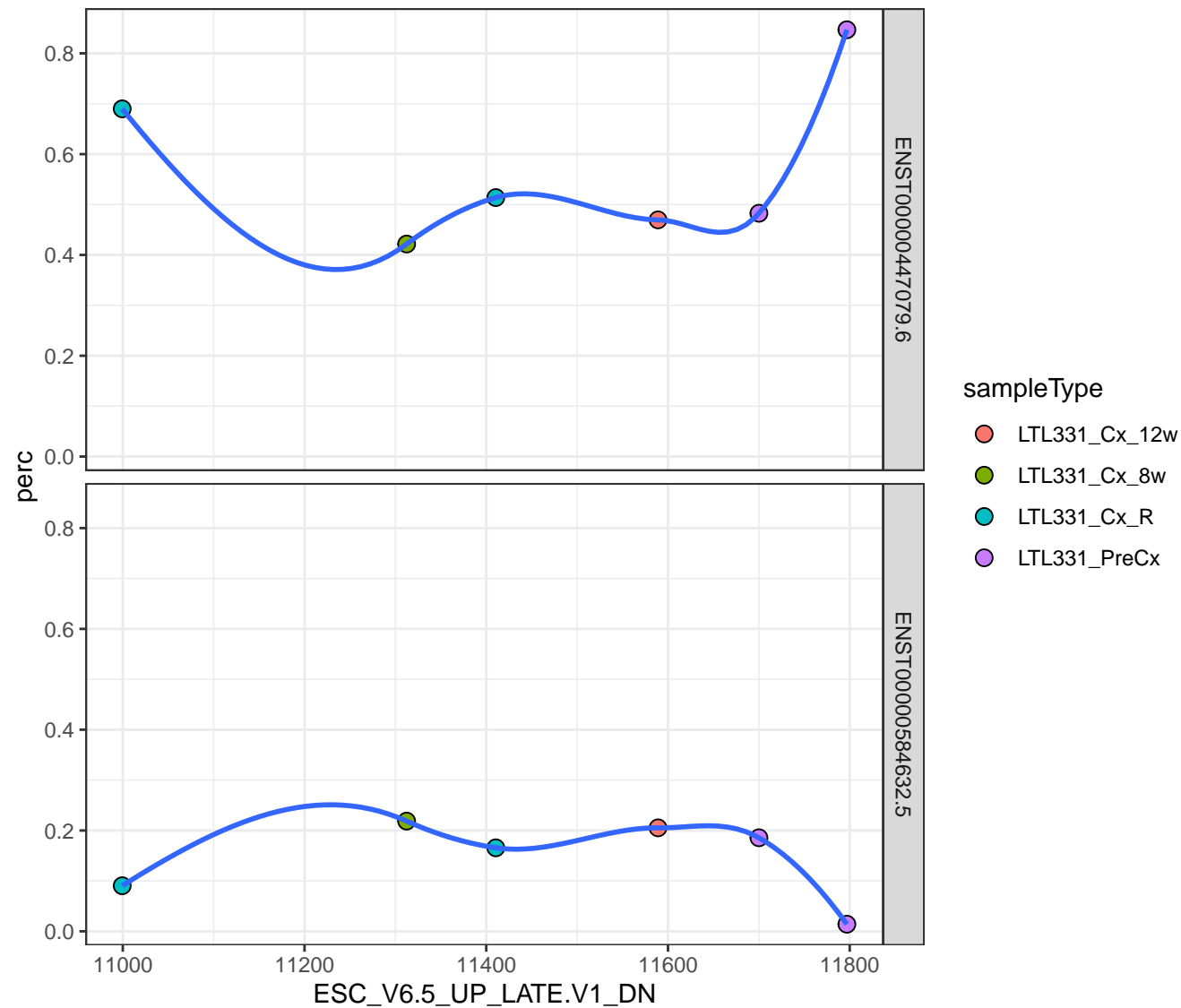
LTL_longitudinal

Gene = CDK12

Target = ESC_V6.5_UP_LATE.V1_DN

ENST00000447079.6 (up) = 0.120697366503852

ENST00000584632.5 (down) = -0.116528012129962



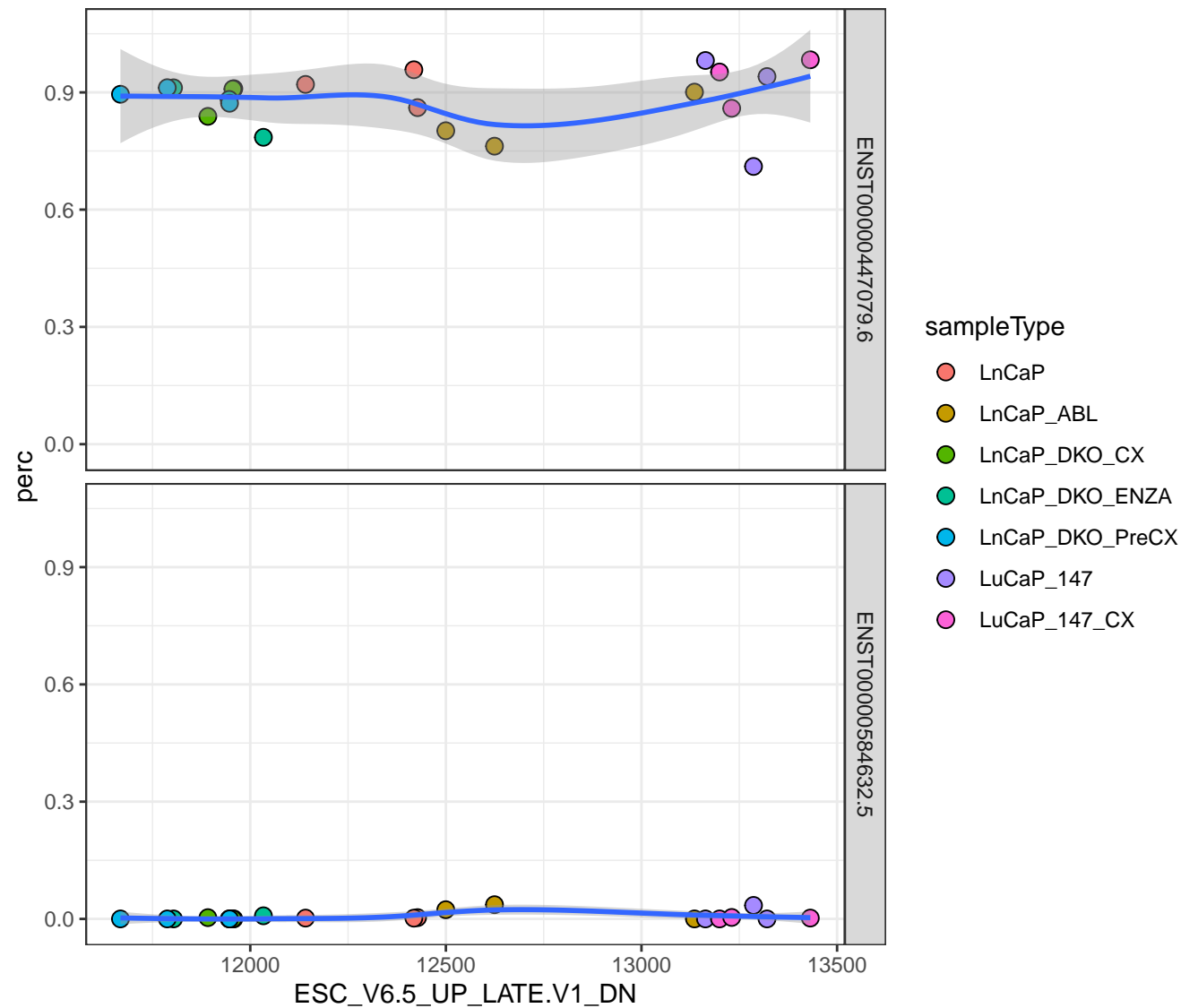
PDX_longitudinal

Gene = CDK12

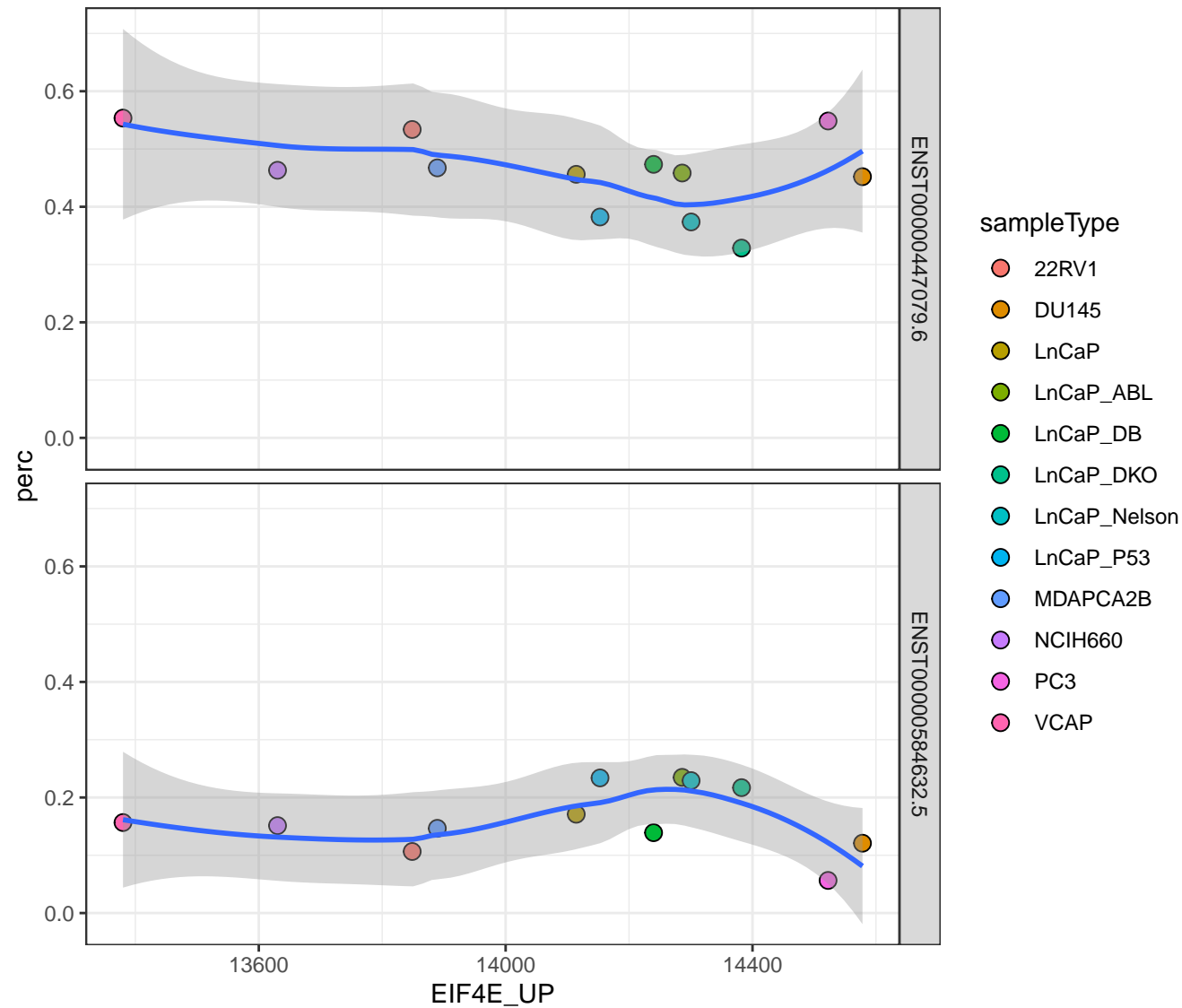
Target = ESC_V6.5_UP_LATE.V1_DN

ENST00000447079.6 (up) = 0.0909085400440463

ENST00000584632.5 (down) = 0.239700179469419



ENST00000584632.5 (down) = 0.0219531615748838



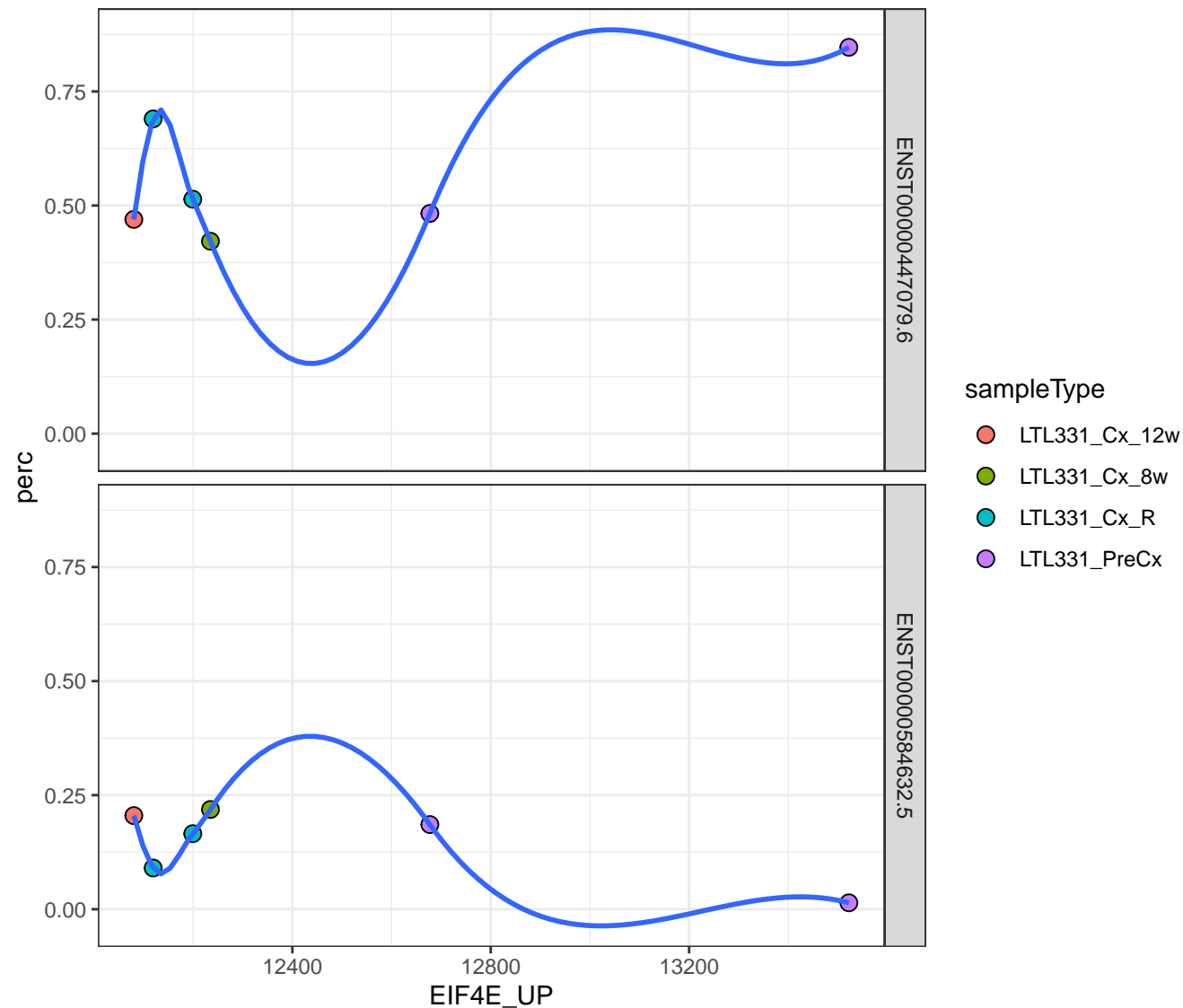
LTL_longitudinal

Gene = CDK12

Target = EIF4E_UP

ENST00000447079.6 (up) = 0.702408403681342

ENST00000584632.5 (down) = -0.710915662163961



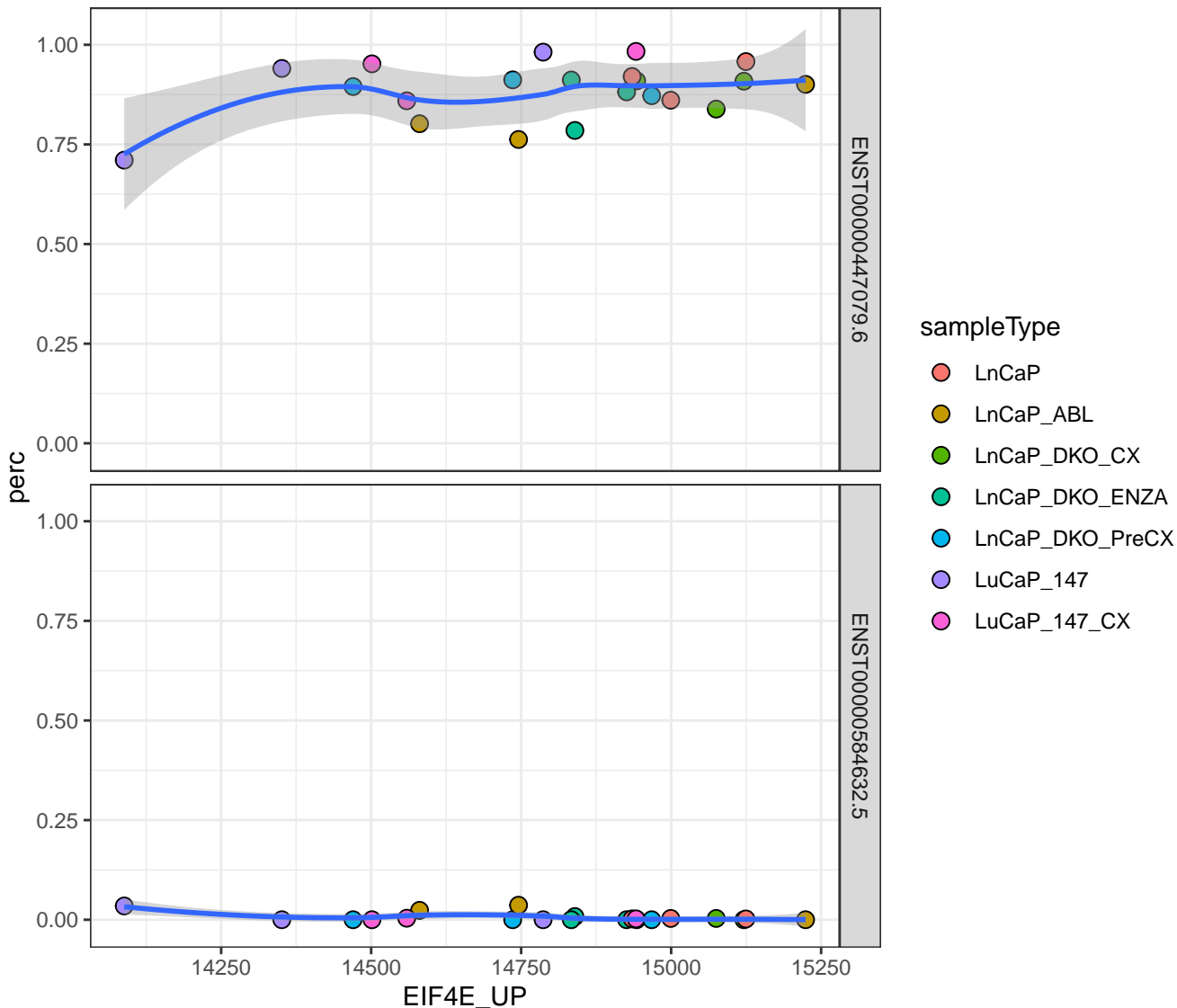
PDX_longitudinal

Gene = CDK12

Target = EIF4E_UP

ENST00000447079.6 (up) = 0.345188461529768

ENST00000584632.5 (down) = -0.457416743590816



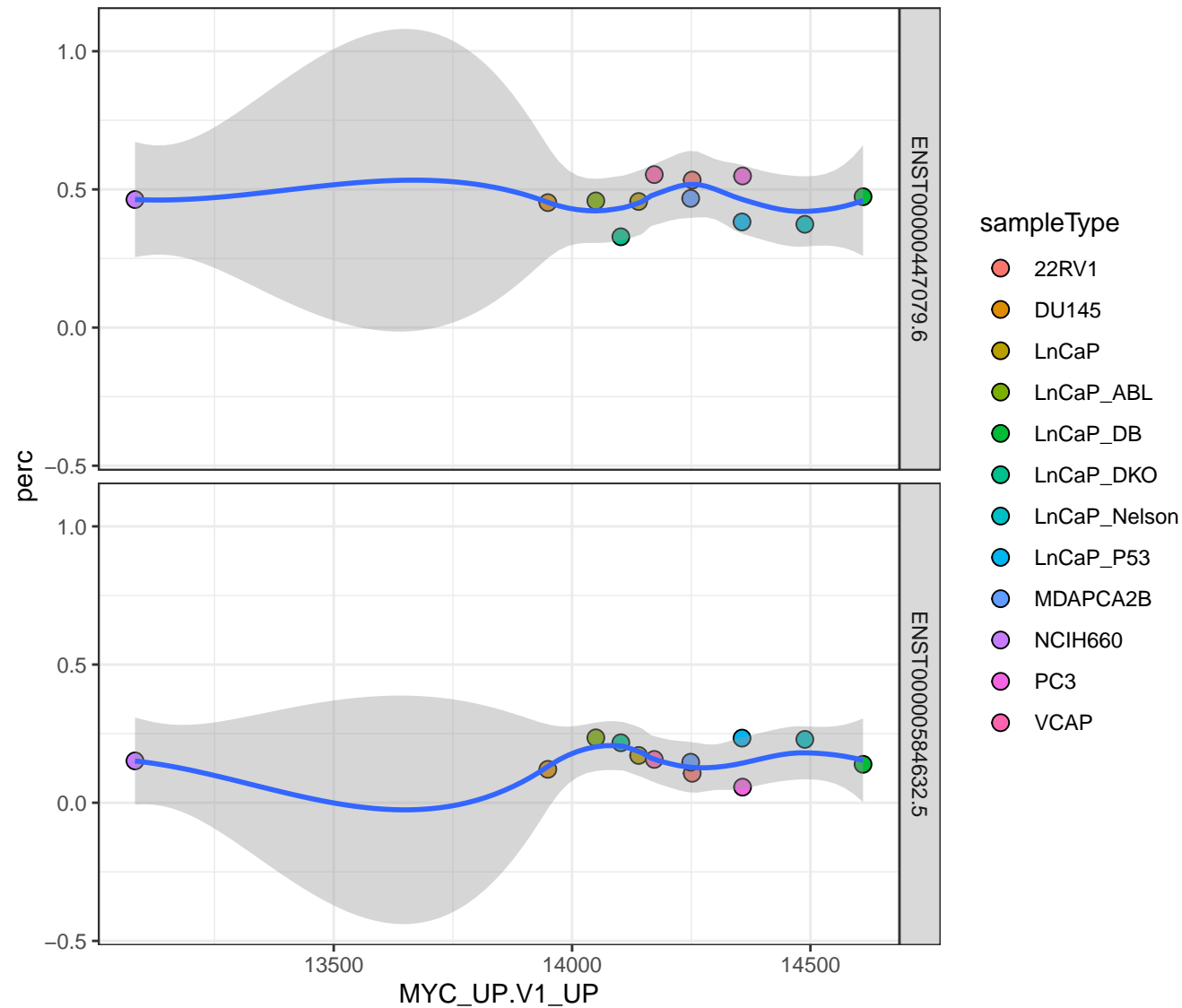
CellLine

Gene = CDK12

Target = MYC_UP.V1_UP

ENST00000447079.6 (up) = -0.019221166778354

ENST00000584632.5 (down) = 0.0311874836762465



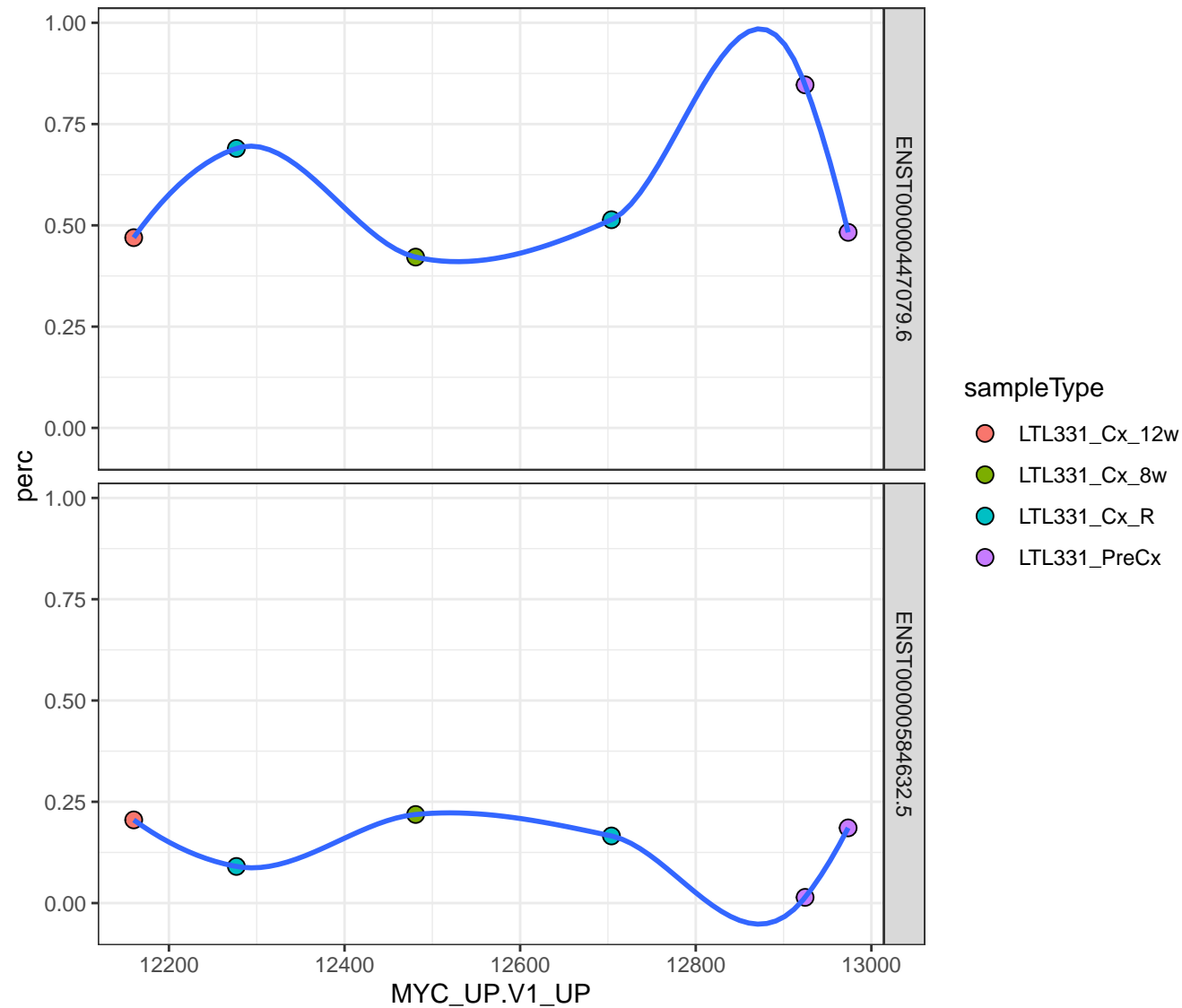
LTL_longitudinal

Gene = CDK12

Target = MYC_UP.V1_UP

ENST00000447079.6 (up) = 0.270613124464655

ENST00000584632.5 (down) = -0.320809129900247



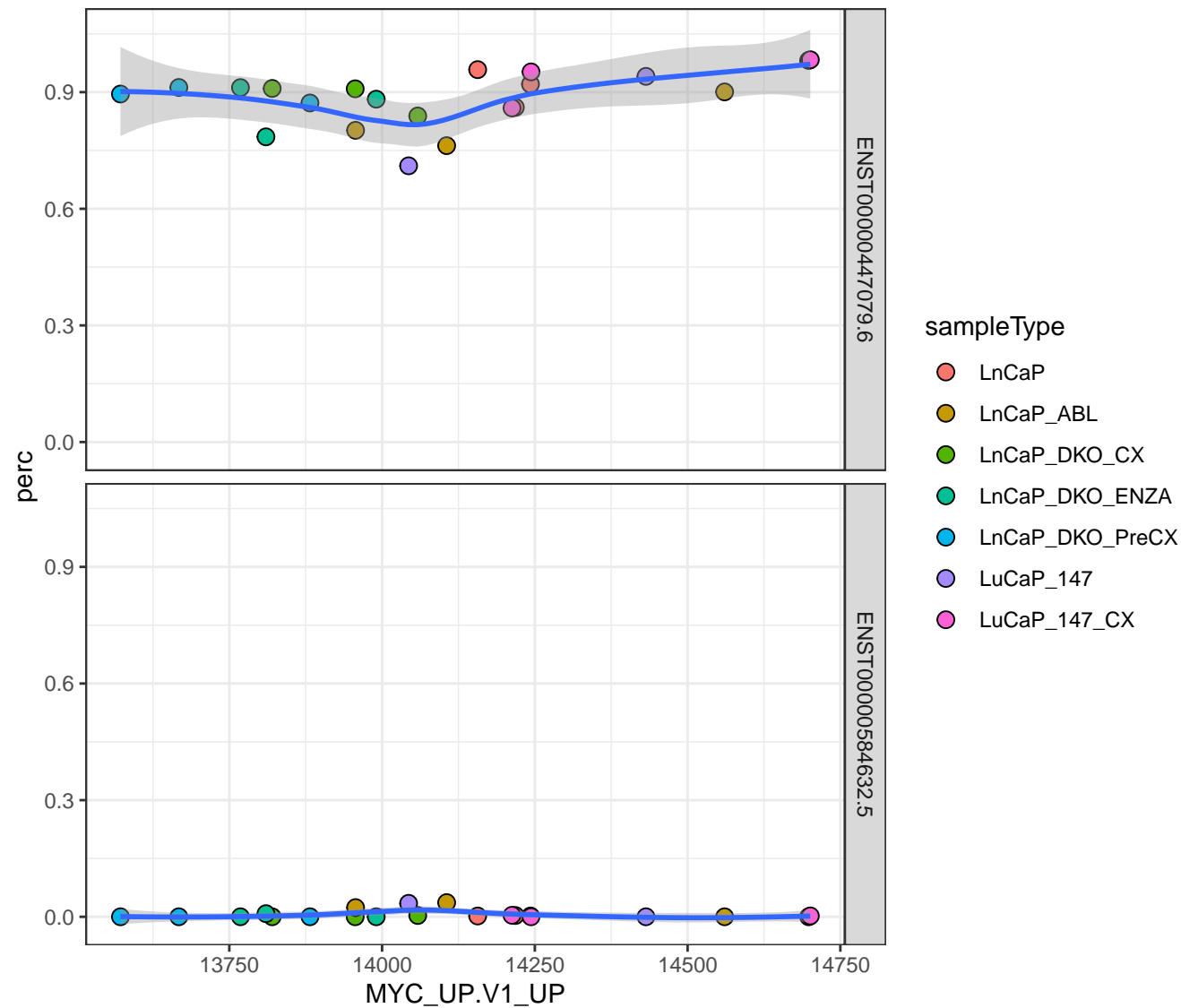
PDX_longitudinal

Gene = CDK12

Target = MYC_UP.V1_UP

ENST00000447079.6 (up) = 0.390927447186091

ENST00000584632.5 (down) = -0.0715085317777073



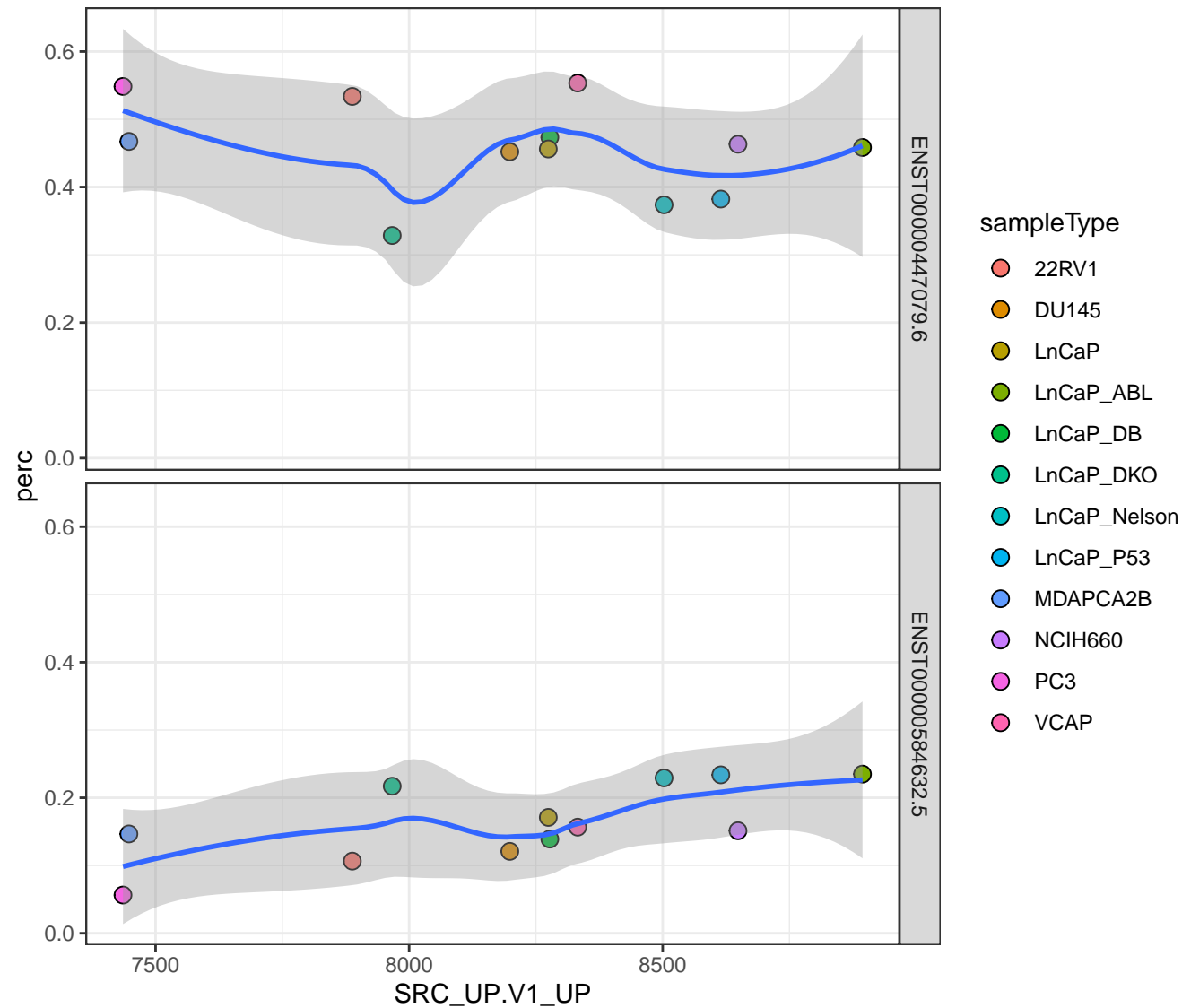
CellLine

Gene = CDK12

Target = SRC_UP.V1_UP

ENST00000447079.6 (up) = -0.316937780806682

ENST00000584632.5 (down) = 0.676807766302426



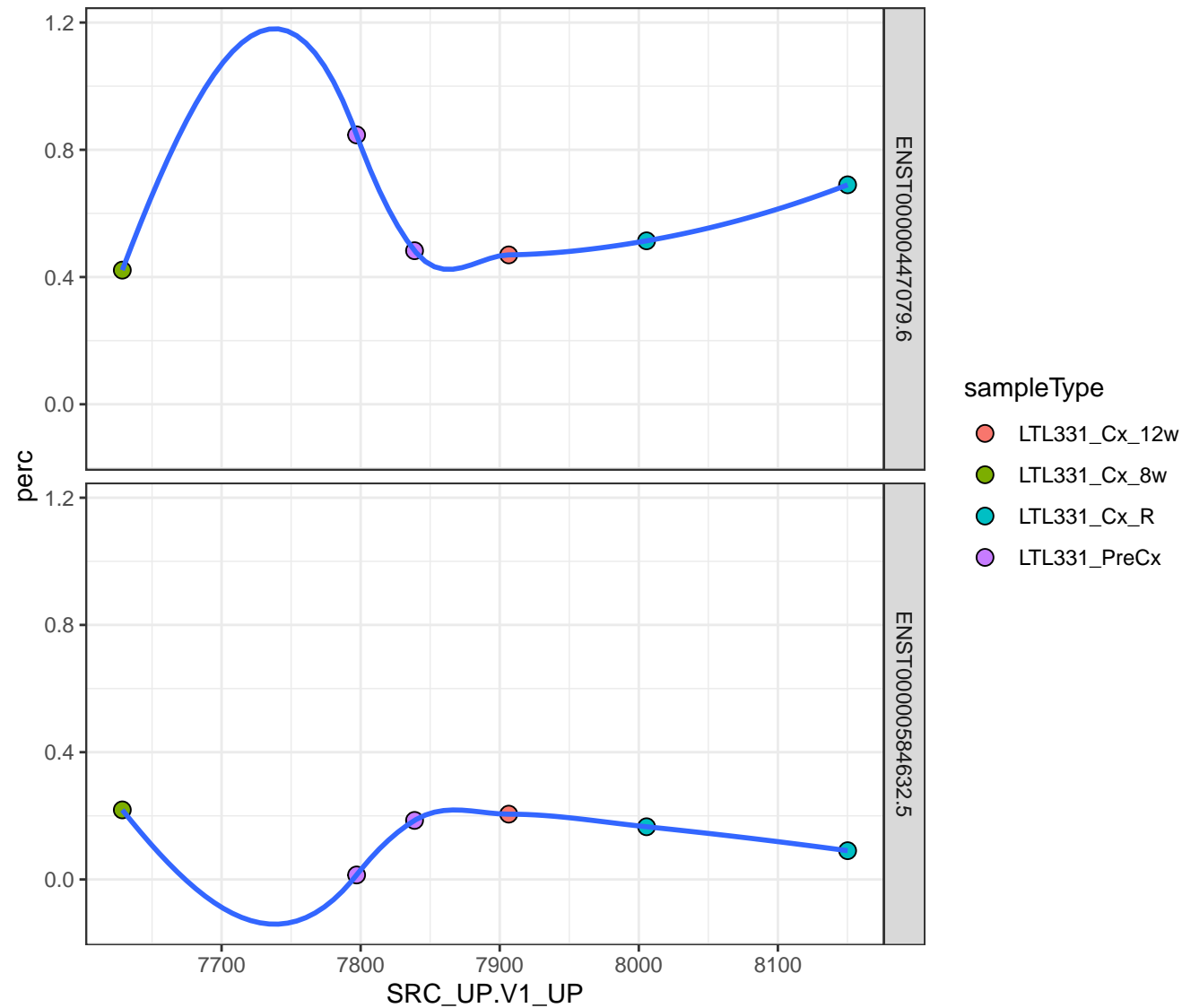
LTL_longitudinal

Gene = CDK12

Target = SRC_UP.V1_UP

ENST00000447079.6 (up) = 0.276614234946733

ENST00000584632.5 (down) = -0.28172655261088



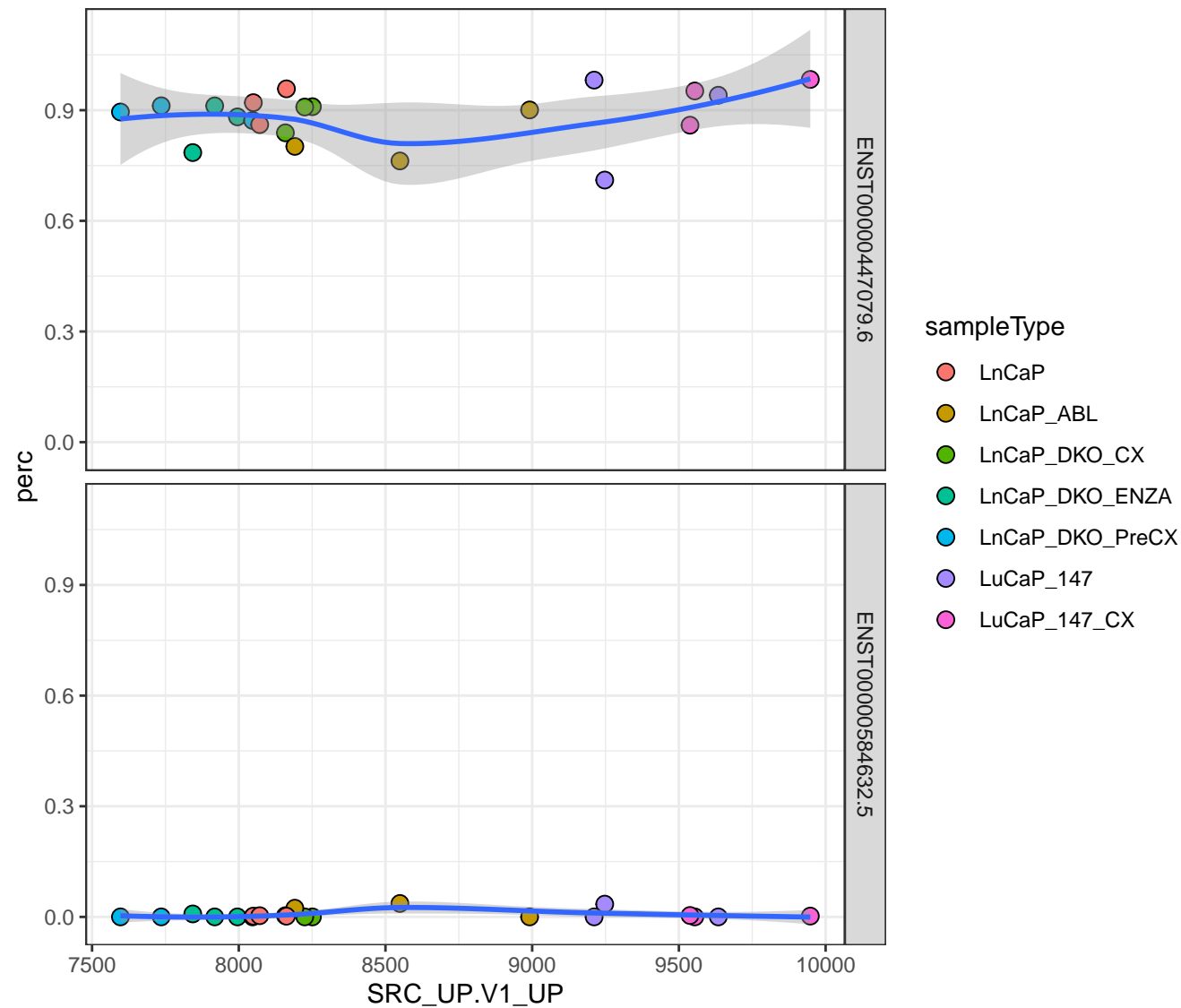
PDX_longitudinal

Gene = CDK12

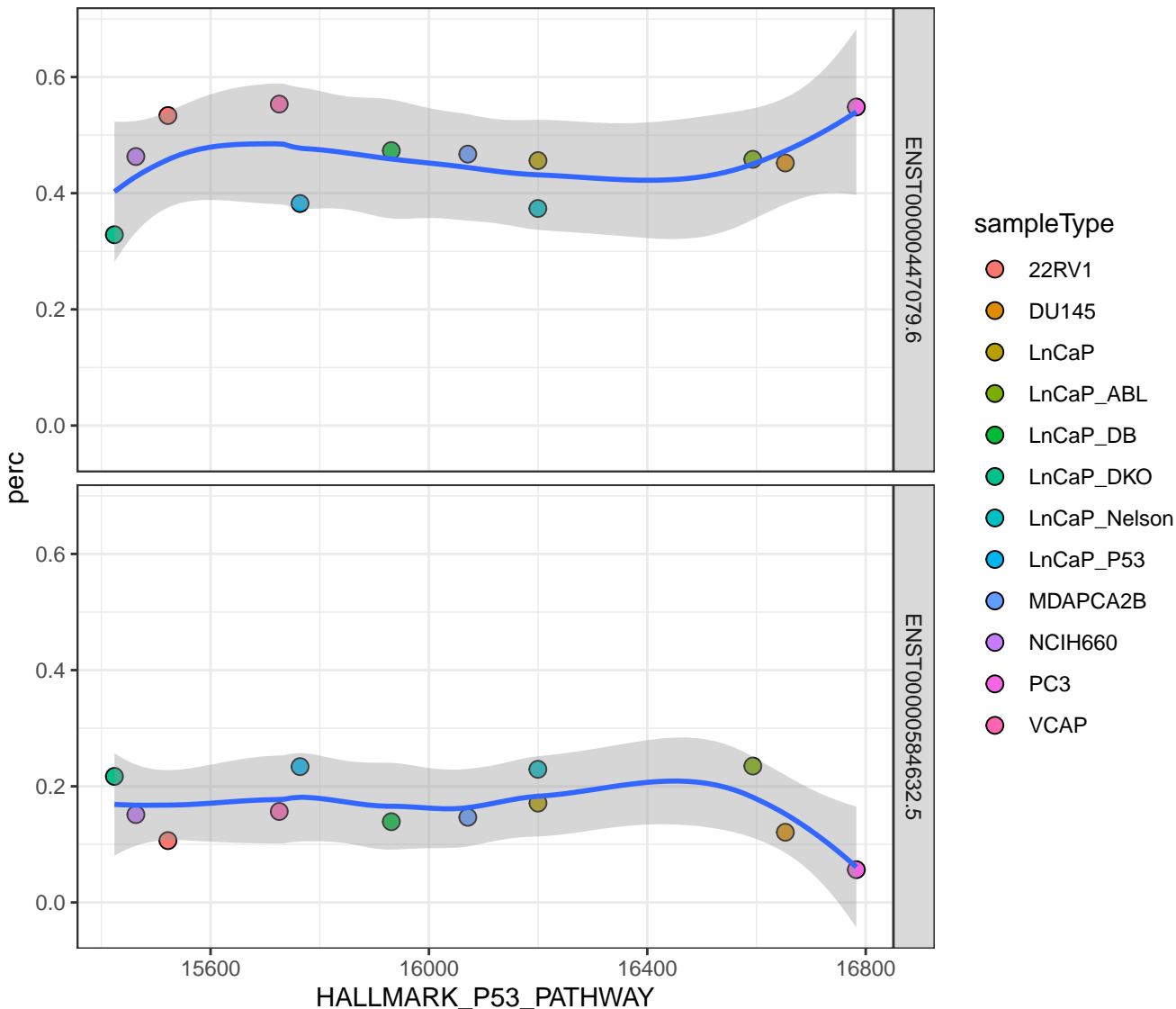
Target = SRC_UP.V1_UP

ENST00000447079.6 (up) = 0.210872844990586

ENST00000584632.5 (down) = 0.097241937868499



CellLine
 Gene = CDK12
 Target = HALLMARK_P53_PATHWAY
 ENST00000447079.6 (up) = 0.21372074292562
 ENST00000584632.5 (down) = -0.225713140575635



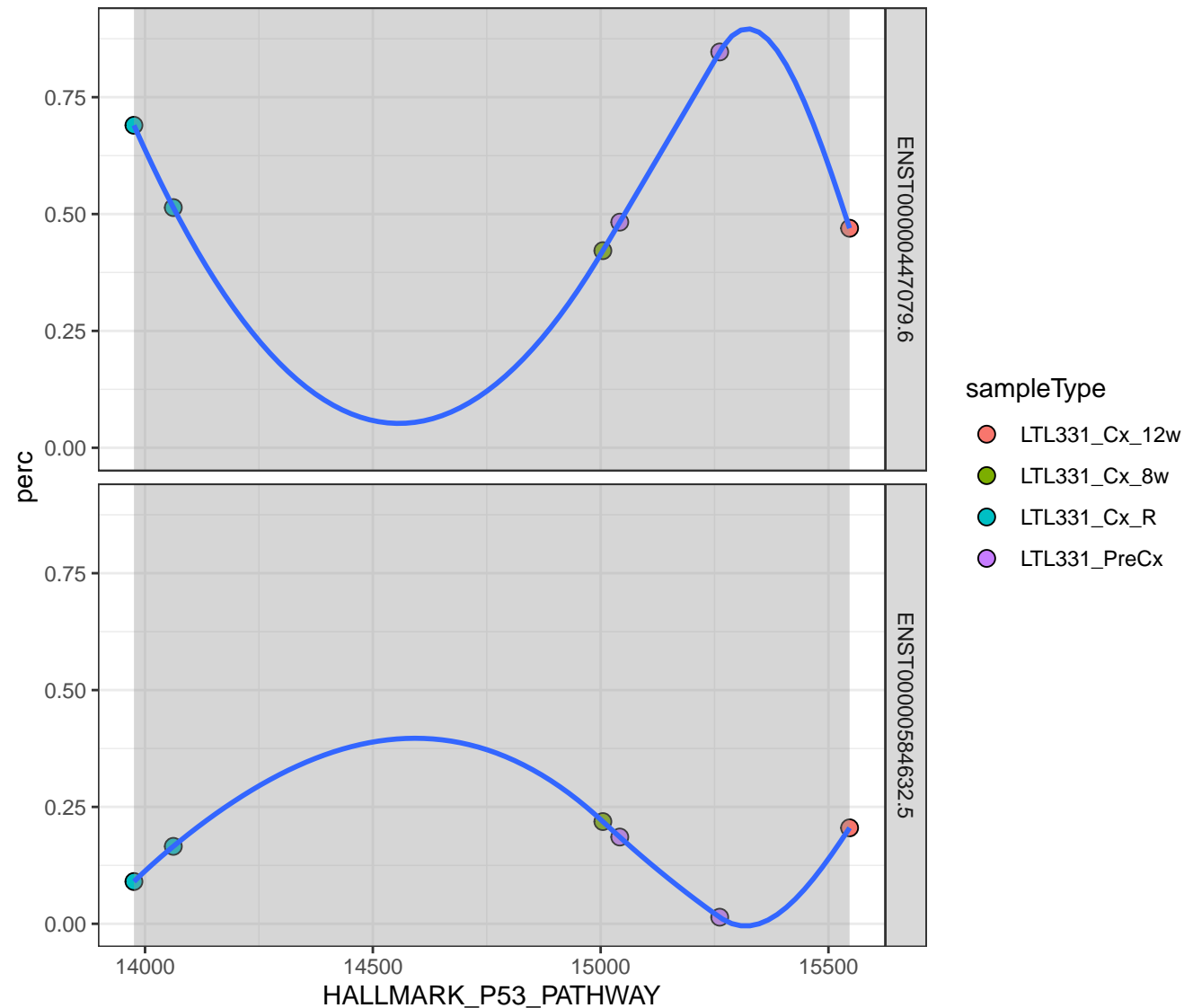
LTL_longitudinal

Gene = CDK12

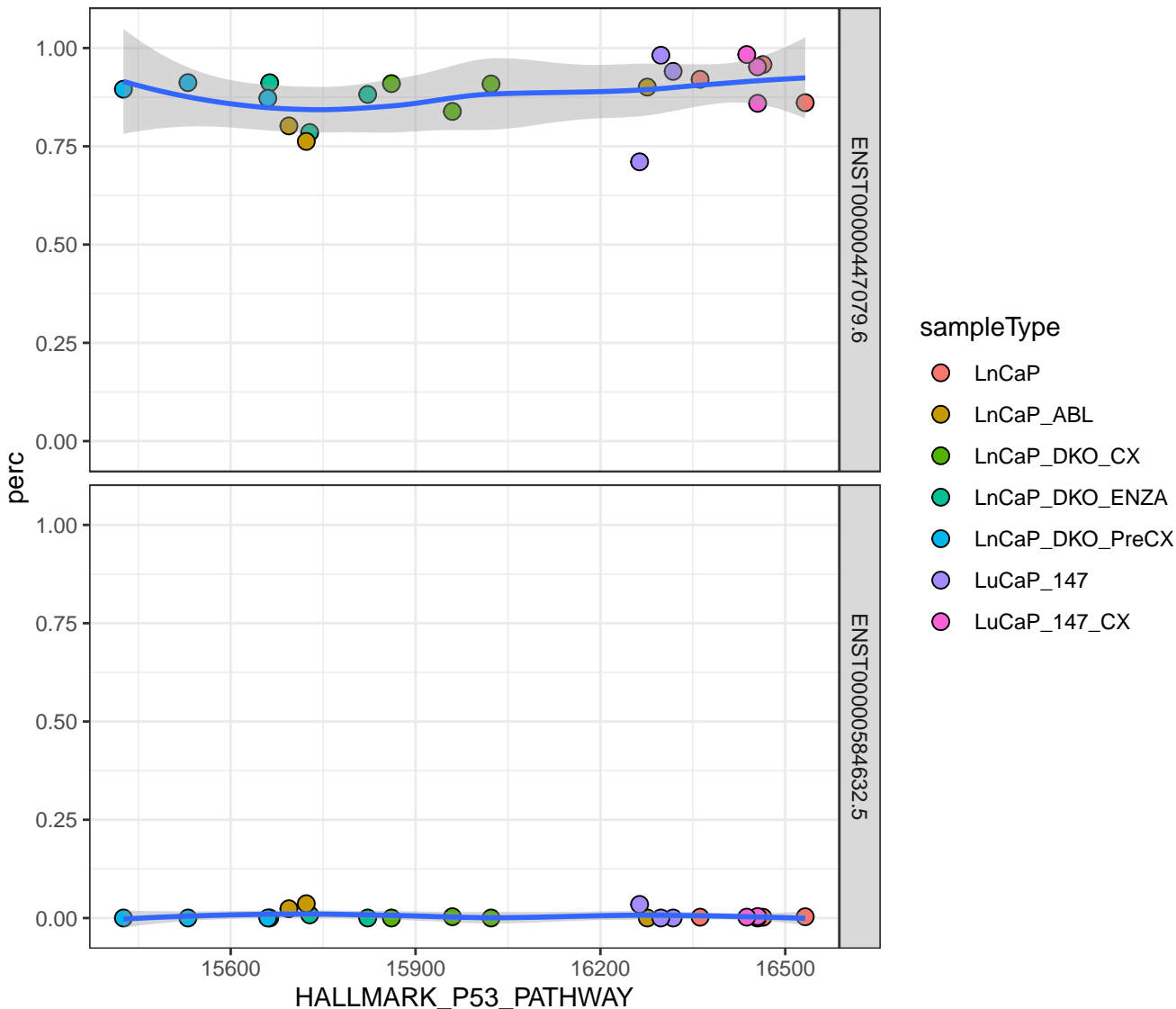
Target = HALLMARK_P53_PATHWAY

ENST00000447079.6 (up) = -0.106271765526606

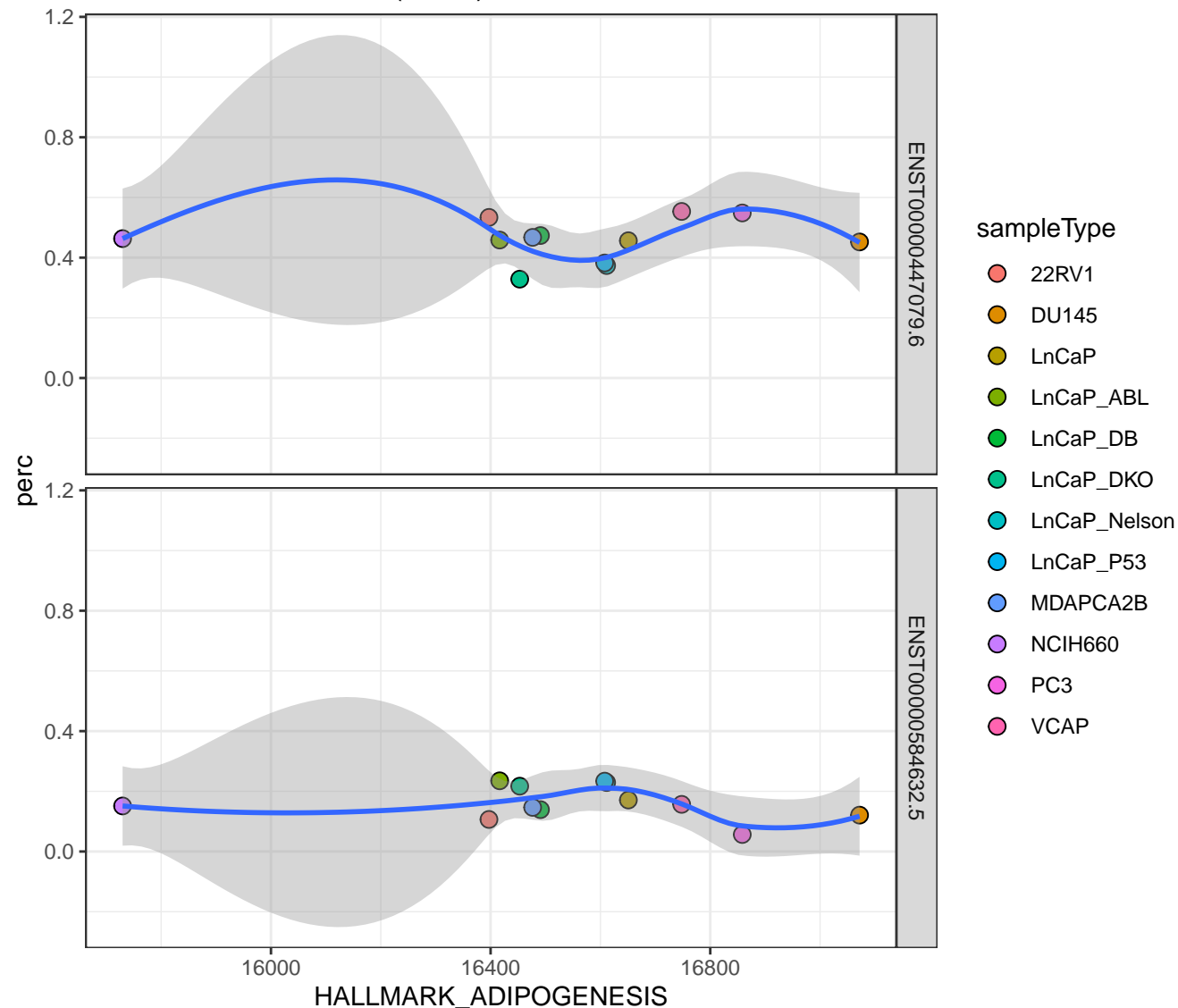
ENST00000584632.5 (down) = 0.152968646943937



PDX_longitudinal
Gene = CDK12
Target = HALLMARK_P53_PATHWAY
ENST00000447079.6 (up) = 0.320702550875552
ENST00000584632.5 (down) = -0.12112137859022



CellLine
 Gene = CDK12
 Target = HALLMARK_ADIPOGENESIS
 ENST00000447079.6 (up) = 0.117238683024871
 ENST00000584632.5 (down) = -0.204567047908495



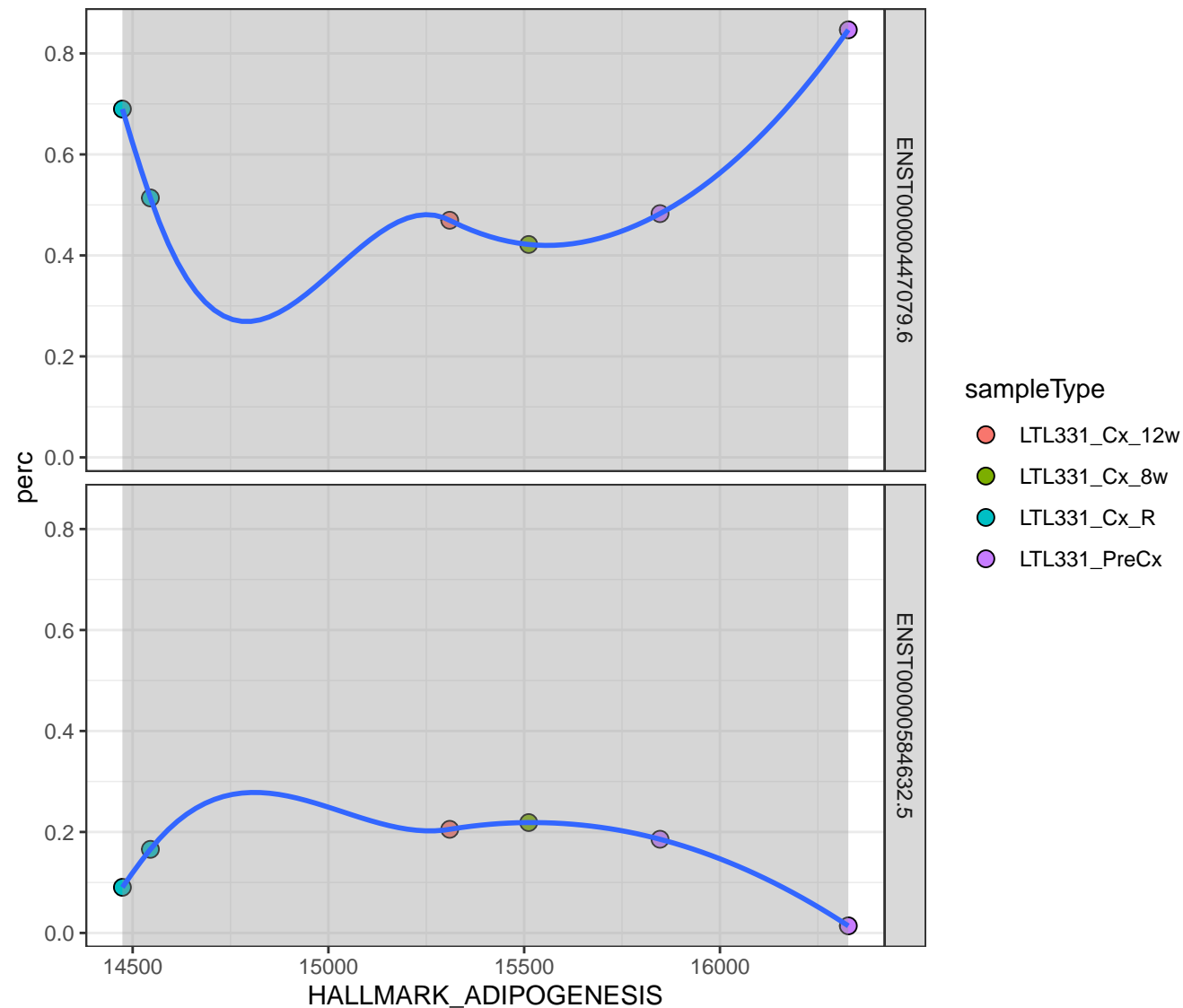
LTL_longitudinal

Gene = CDK12

Target = HALLMARK_ADIPOGENESIS

ENST00000447079.6 (up) = 0.248027560133093

ENST00000584632.5 (down) = -0.233209724112101



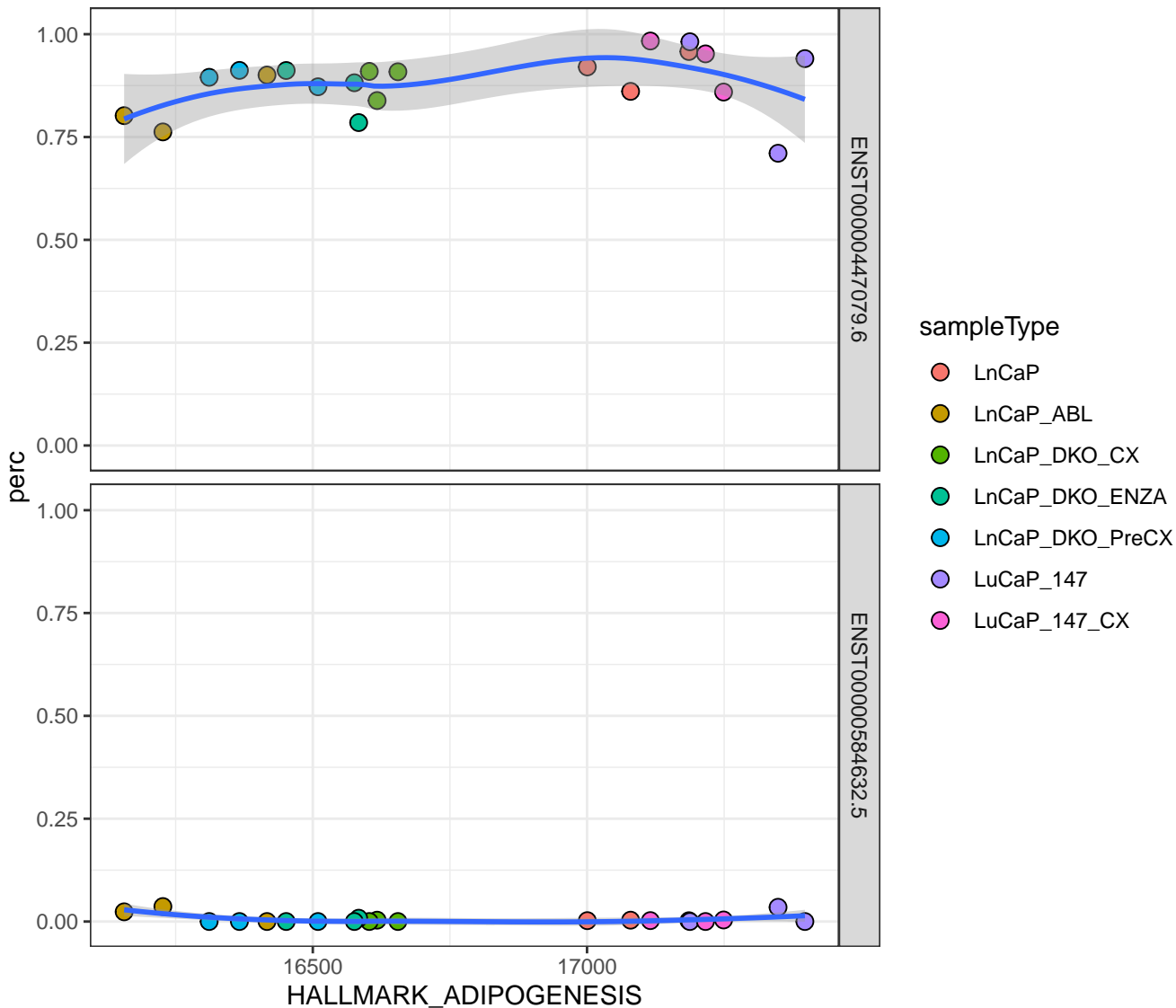
PDX_longitudinal

Gene = CDK12

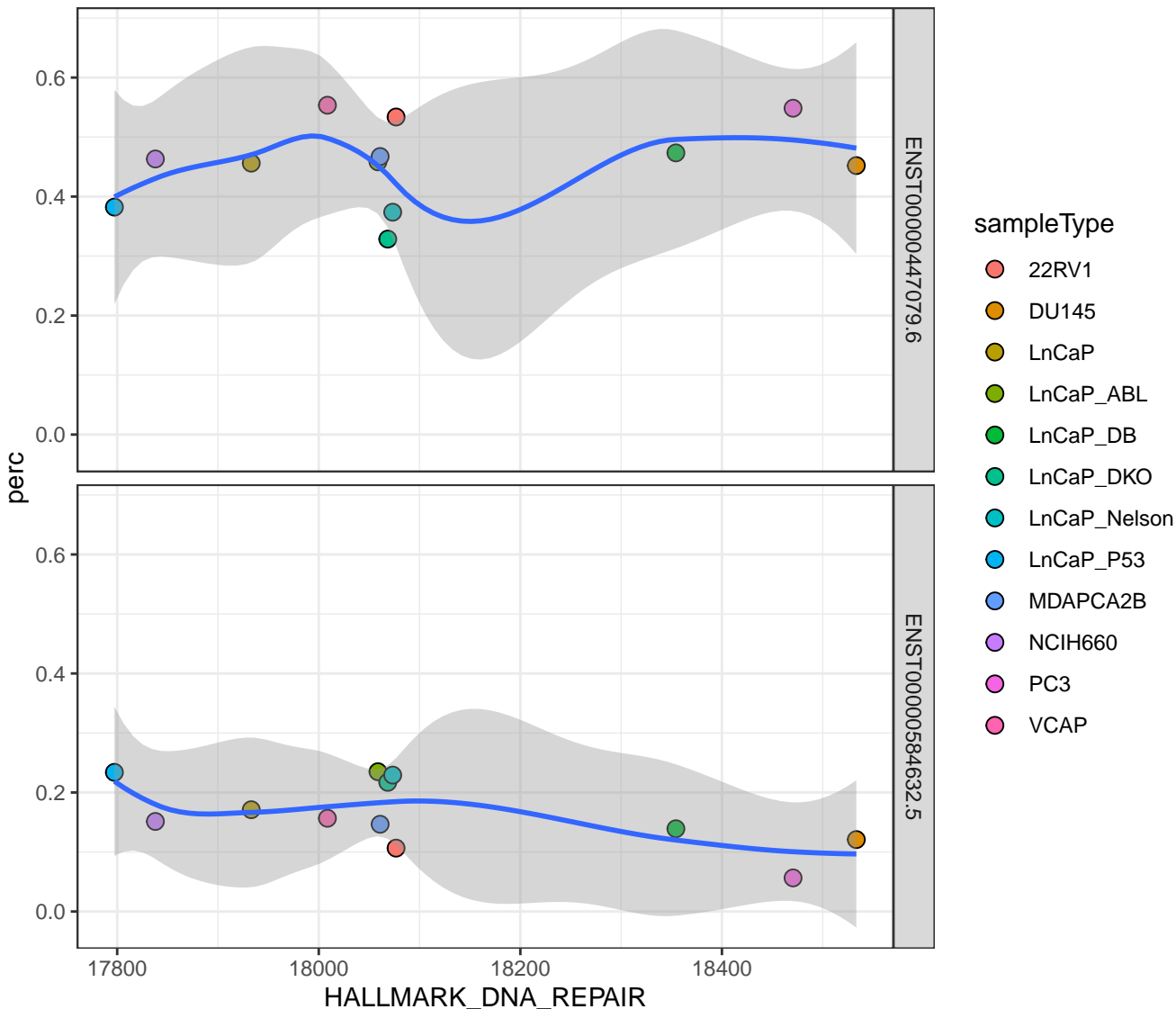
Target = HALLMARK_ADIPOGENESIS

ENST00000447079.6 (up) = 0.29191807857943

ENST00000584632.5 (down) = -0.131010077315246



CellLine
Gene = CDK12
Target = HALLMARK_DNA_REPAIR
ENST00000447079.6 (up) = 0.294953803974921
ENST00000584632.5 (down) = -0.609126453437152



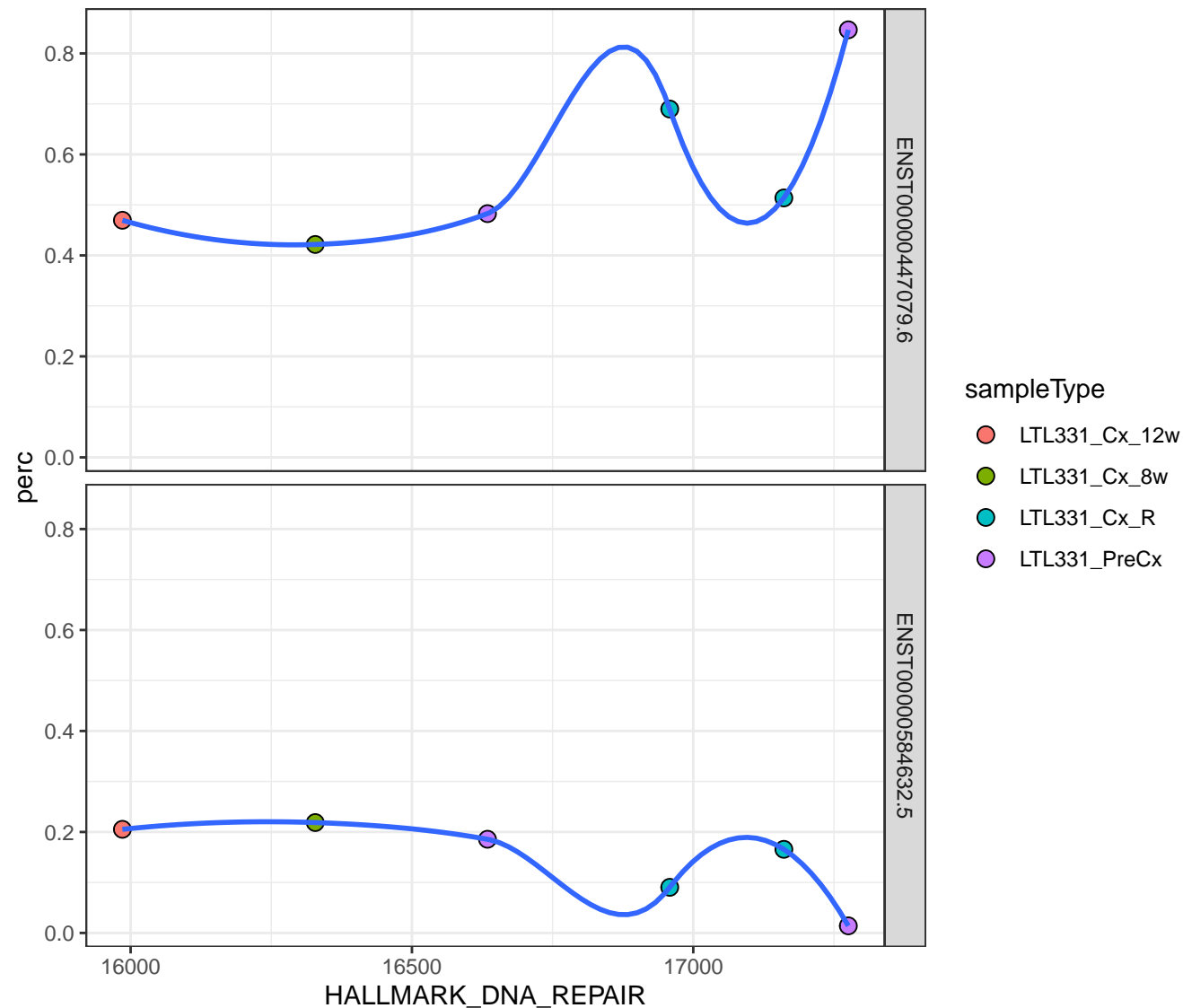
LTL_longitudinal

Gene = CDK12

Target = HALLMARK_DNA_REPAIR

ENST00000447079.6 (up) = 0.722950850813561

ENST00000584632.5 (down) = -0.772506203309161



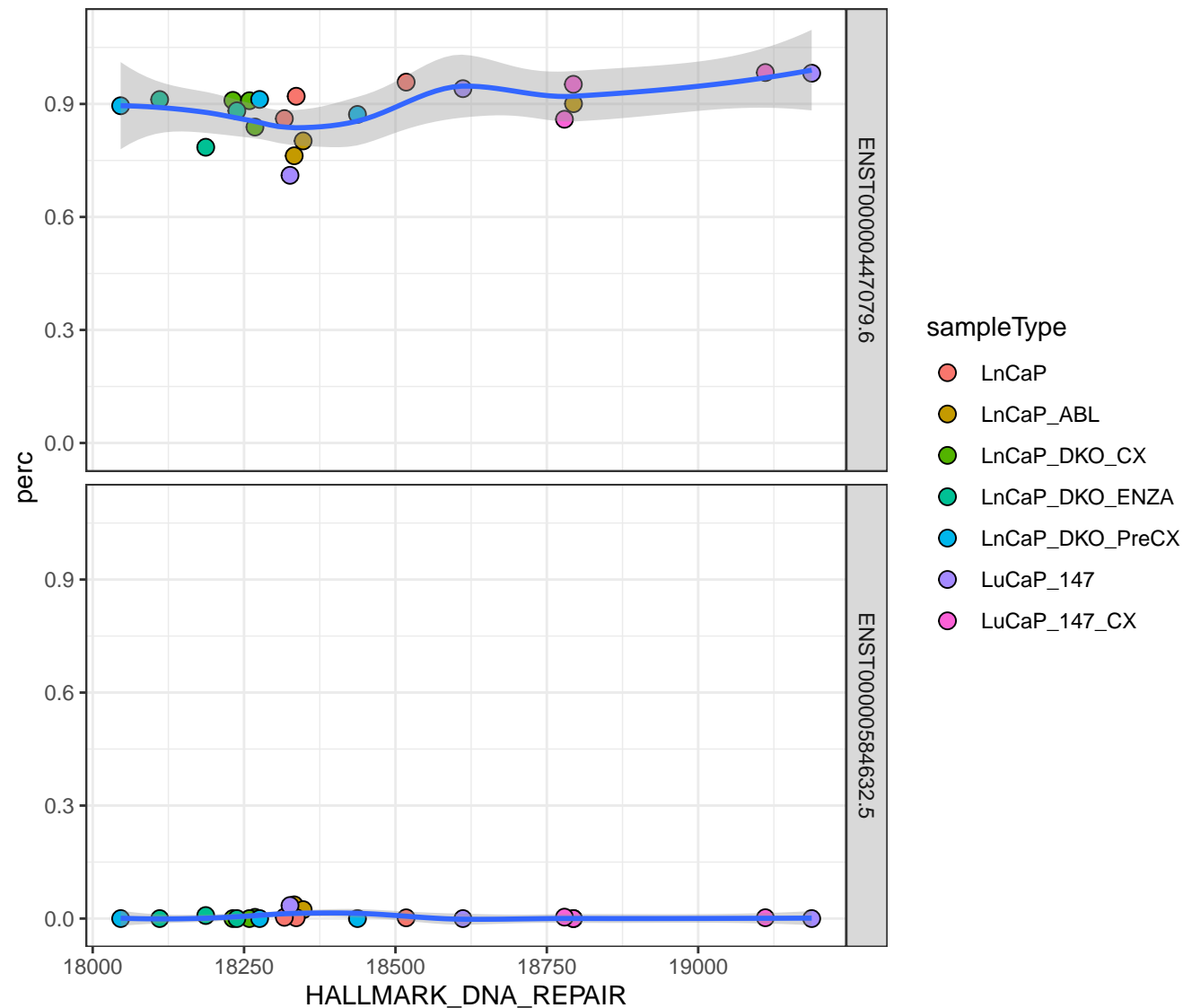
PDX_longitudinal

Gene = CDK12

Target = HALLMARK_DNA_REPAIR

ENST00000447079.6 (up) = 0.495308151822167

ENST00000584632.5 (down) = -0.168378103457534



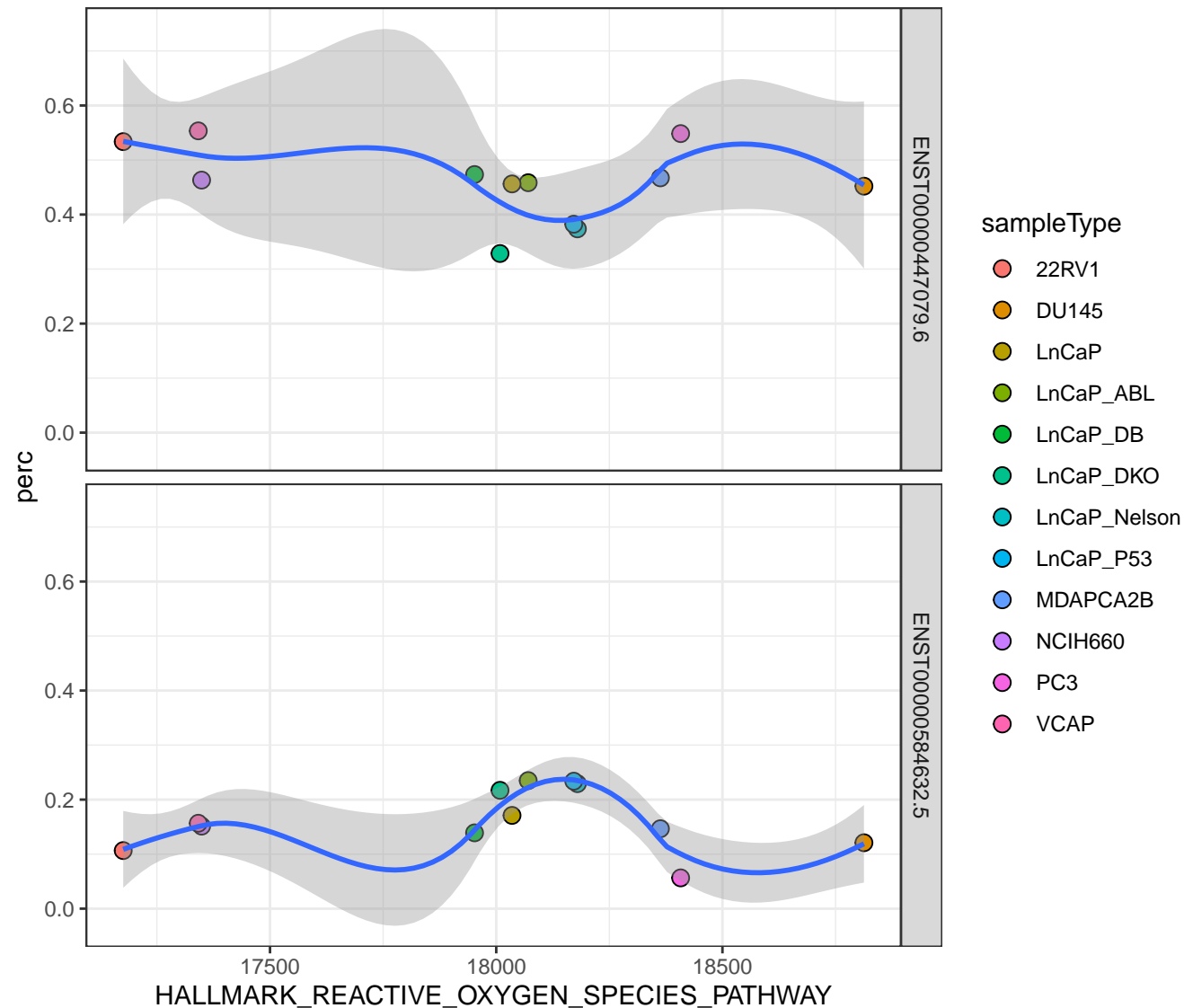
CellLine

Gene = CDK12

Target = HALLMARK_REACTIVE_OXYGEN_SPECIES_PATHWAY

ENST00000447079.6 (up) = -0.334305883428962

ENST00000584632.5 (down) = 0.0194141815893987



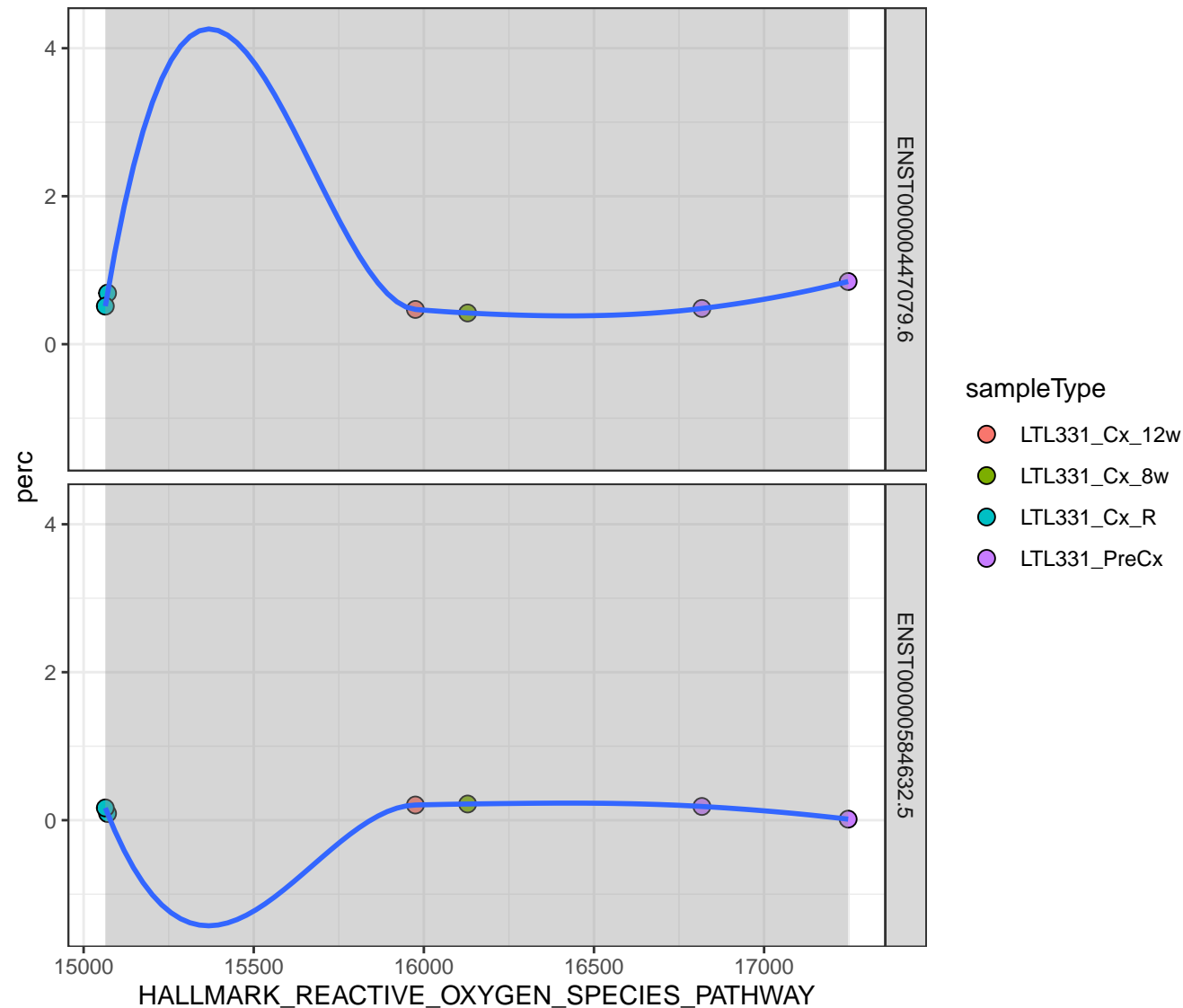
LTL_longitudinal

Gene = CDK12

Target = HALLMARK_REACTIVE_OXYGEN_SPECIES_PATHWAY

ENST00000447079.6 (up) = 0.271963109322279

ENST00000584632.5 (down) = -0.259082174699576



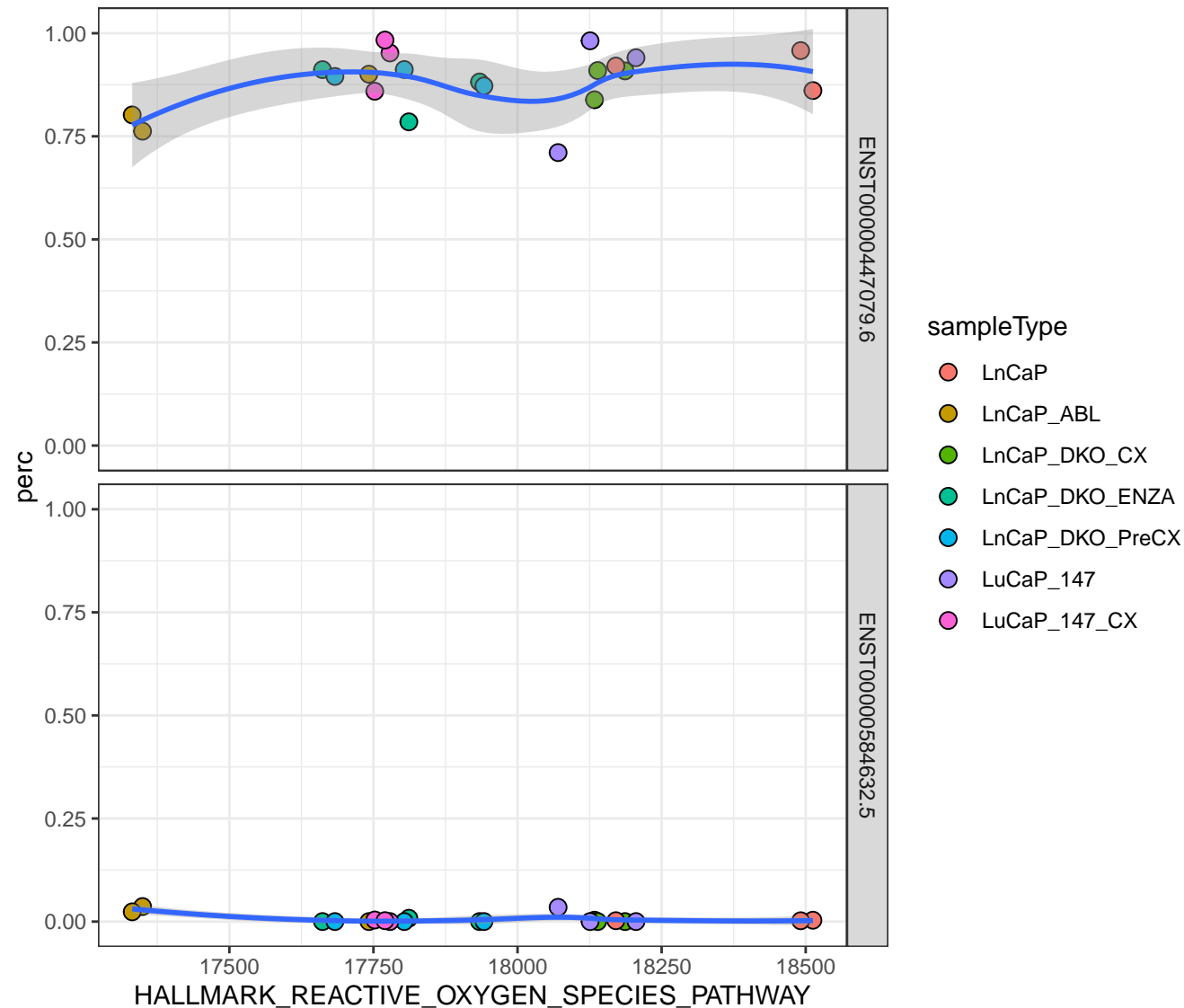
PDX_longitudinal

Gene = CDK12

Target = HALLMARK_REACTIVE_OXYGEN_SPECIES_PATHWAY

ENST00000447079.6 (up) = 0.317349540850123

ENST00000584632.5 (down) = -0.405907988081314



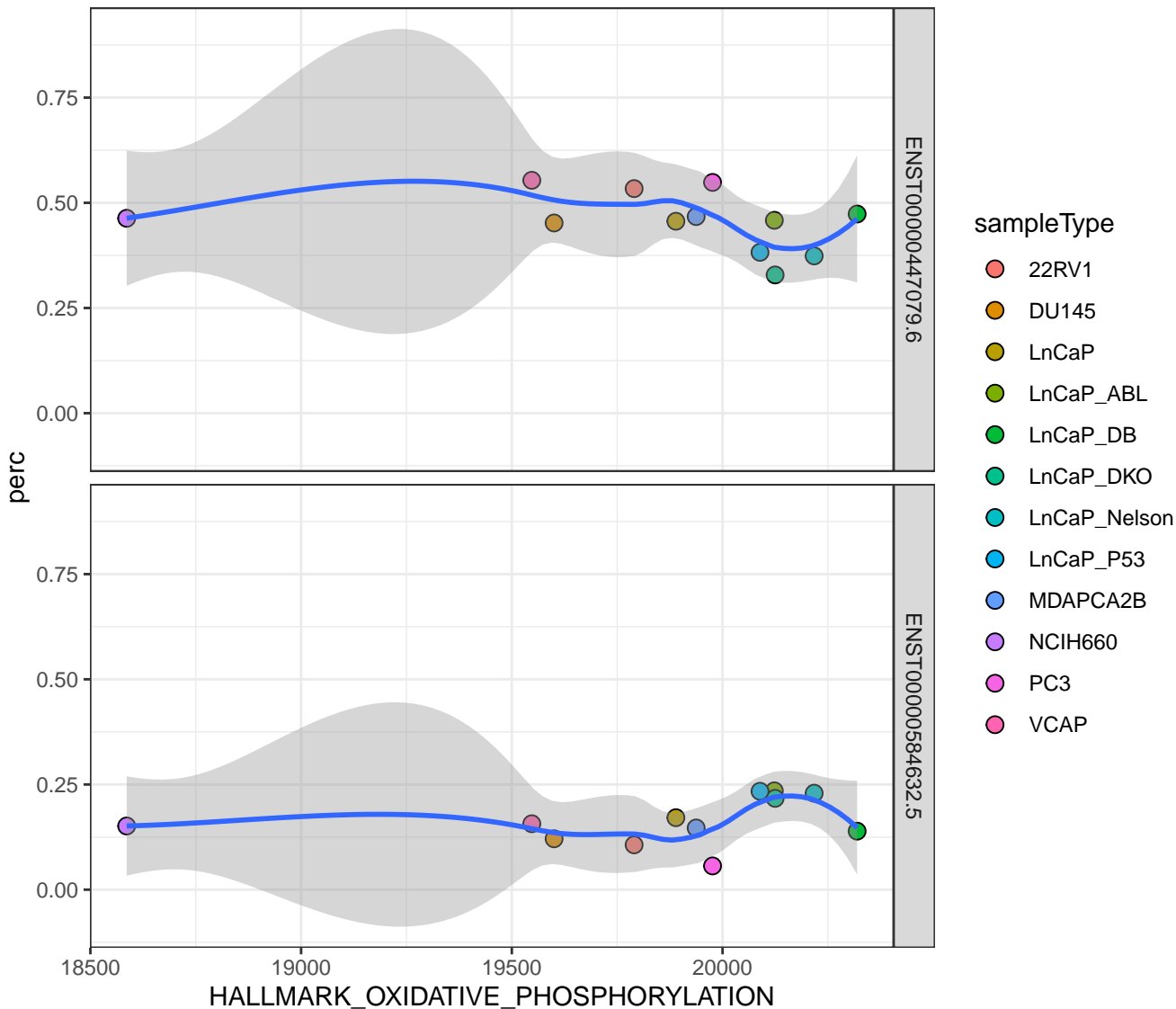
CellLine

Gene = CDK12

Target = HALLMARK_OXIDATIVE_PHOSPHORYLATION

ENST00000447079.6 (up) = -0.293939253764671

ENST00000584632.5 (down) = 0.281521493279512



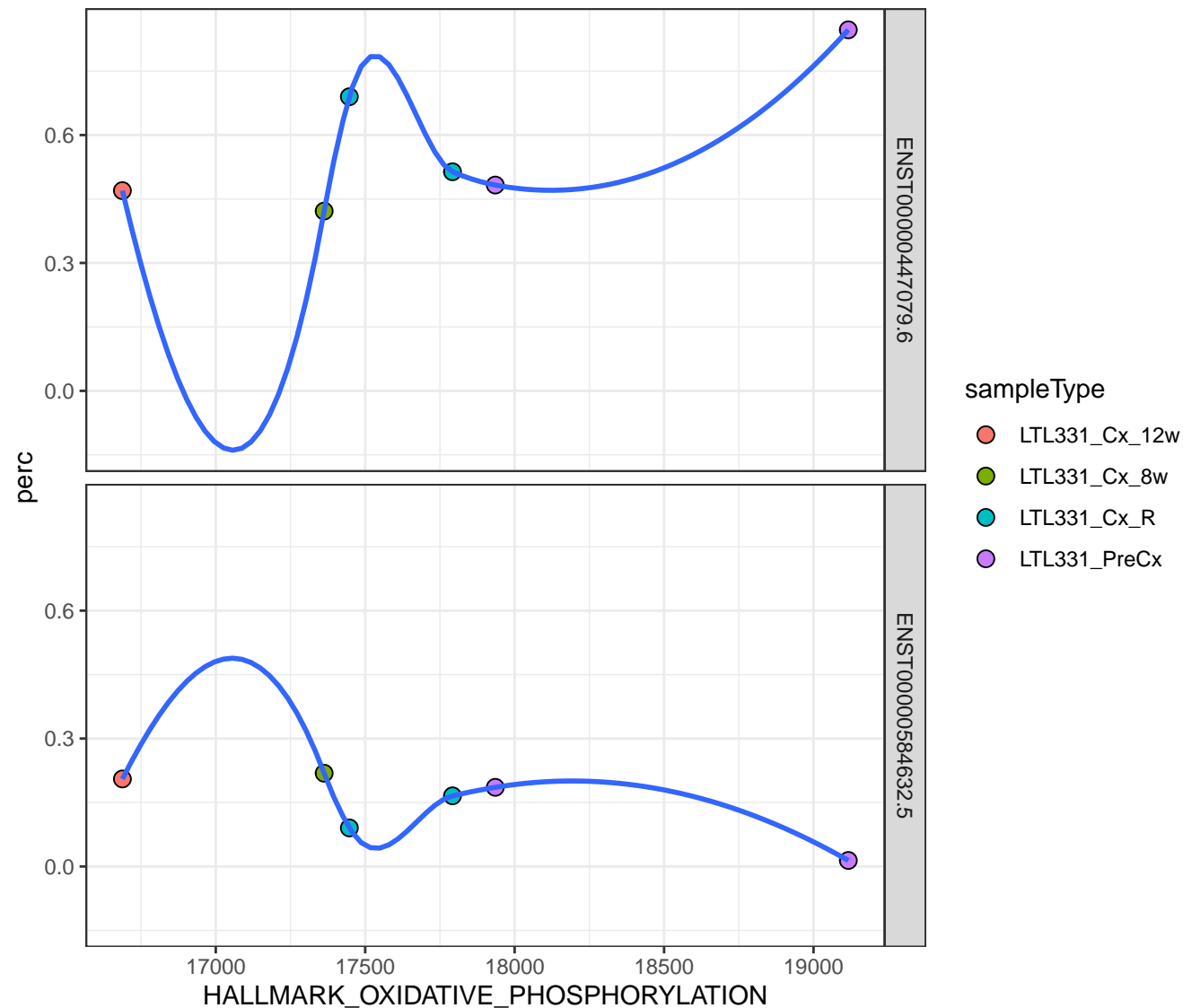
LTL_longitudinal

Gene = CDK12

Target = HALLMARK_OXIDATIVE_PHOSPHORYLATION

ENST00000447079.6 (up) = 0.737114768782182

ENST00000584632.5 (down) = -0.770586868355248



PDX_longitudinal

Gene = CDK12

Target = HALLMARK_OXIDATIVE_PHOSPHORYLATION

ENST00000447079.6 (up) = 0.566891958475669

ENST00000584632.5 (down) = -0.315042367656922

