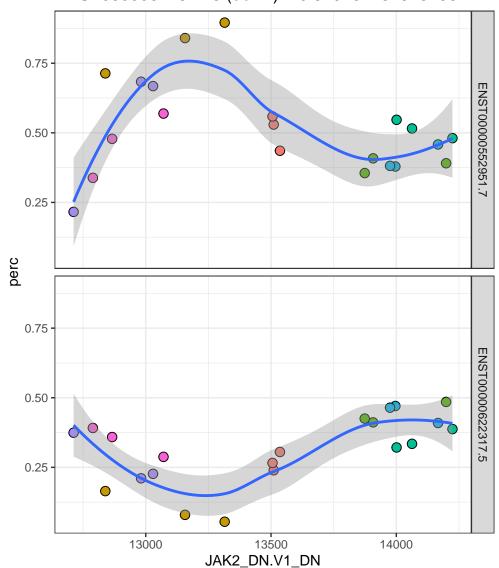


LTL_longitudinal Gene = NOPCHAP1 Target = JAK2_DN.V1_DN ENST00000552951.7 (up) = 0.85141321560867 ENST00000622317.5 (down) = -0.869928468645149 1.00 0.75 ENST00000552951.7 0.50 0.25 sampleType LTL331_Cx_12w 0.00 bec 1.00 LTL331_Cx_8w LTL331_Cx_R LTL331_PreCx 0.75 ENST00000622317.5 0.50 0.25 0.00 13750 14000 14250 14500 JAK2_DN.V1_DN

PDX_longitudinal
Gene = NOPCHAP1
Target = JAK2_DN.V1_DN
ENST00000552951.7 (up) = -0.269796012370067
ENST00000622317.5 (down) = 0.529767234579456



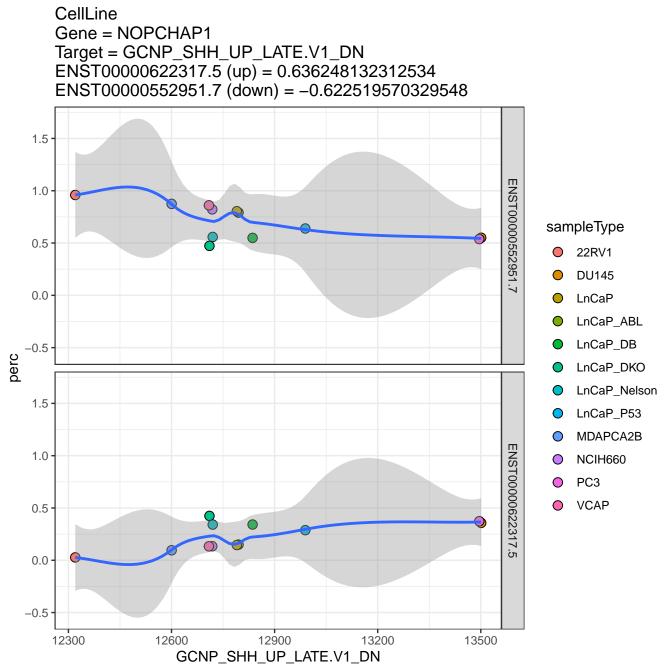
sampleType

- LnCaP
- LnCaP_ABL

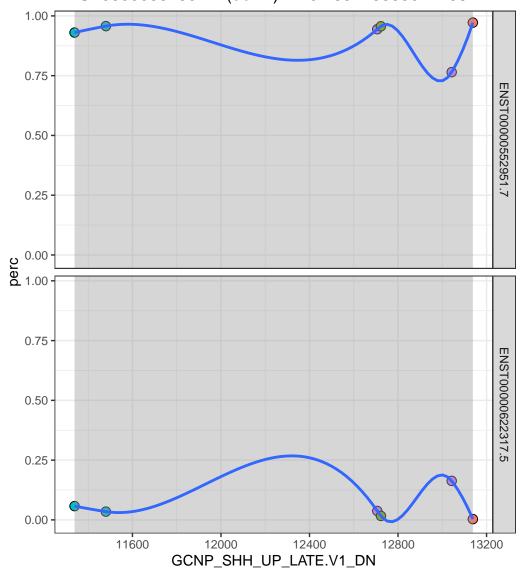
LnCaP_DKO_CX

- LnCaP_DKO_ENZA
- LnCaP_DKO_PreCX
 - LuCaP_147

LuCaP_147_CX



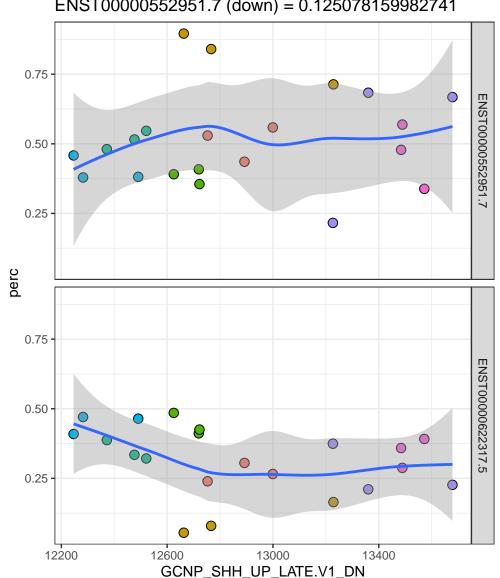
LTL_longitudinal
Gene = NOPCHAP1
Target = GCNP_SHH_UP_LATE.V1_DN
ENST00000622317.5 (up) = 0.129353849312617
ENST00000552951.7 (down) = -0.283778309022238



sampleType

LTL331_Cx_12w

LTL331_Cx_8w LTL331_Cx_R LTL331_PreCx PDX_longitudinal Gene = NOPCHAP1 Target = GCNP_SHH_UP_LATE.V1_DN ENST00000622317.5 (up) = -0.30995322239491 ENST00000552951.7 (down) = 0.125078159982741



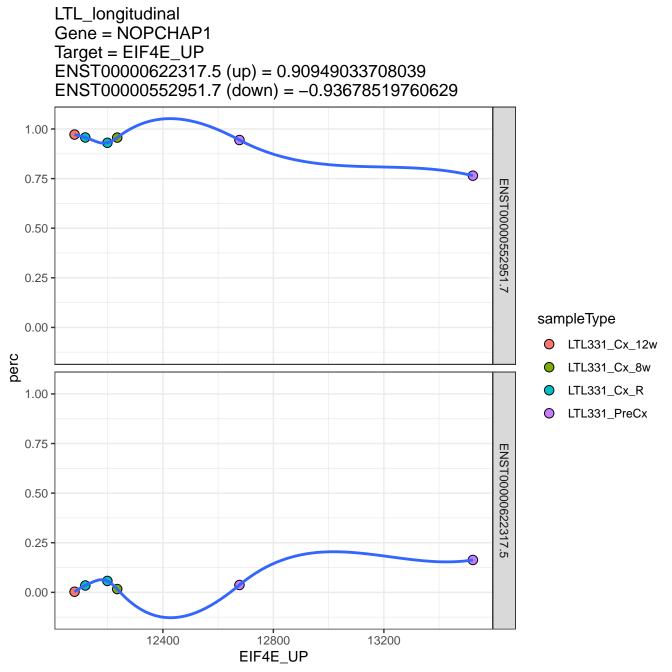
sampleType

- LnCaP
- LnCaP_ABL

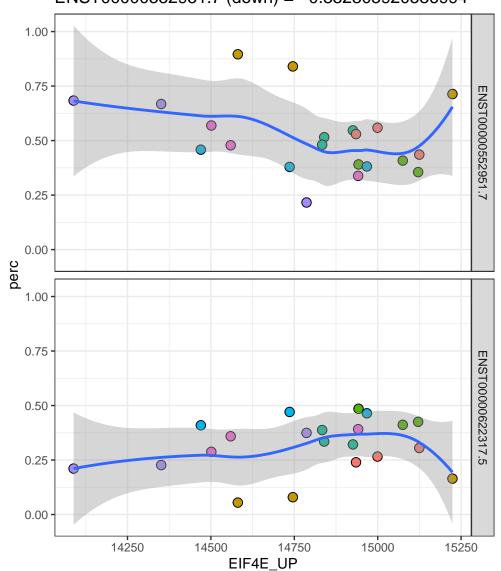
 LnCaP_DKO_CX
- LnCaP_DKO_ENZALnCaP_DKO_PreCX
- LuCaP_147
- LuCaP_147_CX

Gene = NOPCHAP1 Target = EIF4E_UP ENST00000622317.5 (up) = 0.74924226910527ENST00000552951.7 (down) = -0.7783776218633990 0.8 ENST00000552951.7 sampleType 22RV1 0.4 DU145 LnCaP LnCaP_ABL 0.0 LnCaP_DB perc LnCaP_DKO LnCaP_Nelson LnCaP_P53 MDAPCA2B 0.8 ENST00000622317.5 NCIH660 PC3 **VCAP** 0.4 0.0 13600 14000 14400 EIF4E_UP

CellLine



PDX_longitudinal Gene = NOPCHAP1 Target = EIF4E_UP ENST00000622317.5 (up) = 0.257470011869658 ENST00000552951.7 (down) = -0.352305920536994



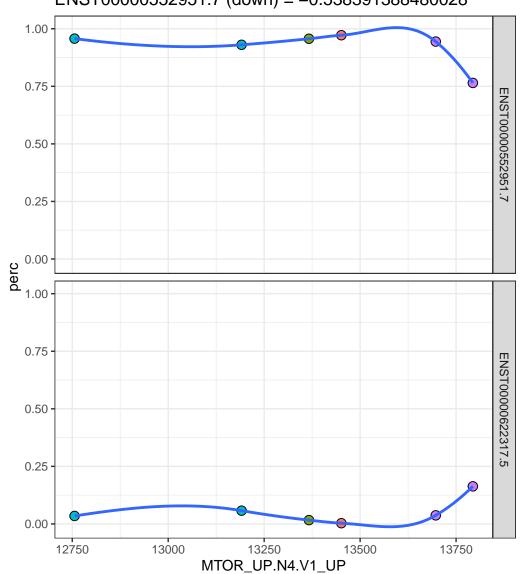
sampleType

- LnCaP
- LnCaP_ABL
- LnCaP_DKO_CXLnCaP_DKO_ENZA
- LnCaP_DKO_PreCX
- LuCaP_147
- LuCaP_147_CX

CellLine Gene = NOPCHAP1 Target = MTOR_UP.N4.V1_UP ENST00000622317.5 (up) = 0.510648730187374ENST00000552951.7 (down) = -0.514214430261314ENST00000552951.7 sampleType 22RV1 **DU145** 0 LnCaP LnCaP_ABL LnCaP_DB perc LnCaP_DKO LnCaP_Nelson LnCaP_P53 MDAPCA2B ENST00000622317.5 NCIH660 PC3 **VCAP** 0 0 14500 15000 15500

MTOR_UP.N4.V1_UP

LTL_longitudinal
Gene = NOPCHAP1
Target = MTOR_UP.N4.V1_UP
ENST00000622317.5 (up) = 0.446139898305638
ENST00000552951.7 (down) = -0.538391388480028



sampleType

LTL331_Cx_12w

LTL331_Cx_8w

LTL331_Cx_R LTL331_PreCx

PDX_longitudinal Gene = NOPCHAP1 Target = MTOR_UP.N4.V1_UP ENST00000622317.5 (up) = 0.564379481049295ENST00000552951.7 (down) = -0.5365264748143551.00 ENST00000552951.7 0.75 -0 0 0.50 -0.25 sampleType 0 LnCaP 0.00 LnCaP_ABL LnCaP_DKO_CX perc LnCaP_DKO_ENZA LnCaP_DKO_PreCX 1.00 LuCaP_147 LuCaP_147_CX ENST00000622317.5 0.75 -0.50

15600

0.25

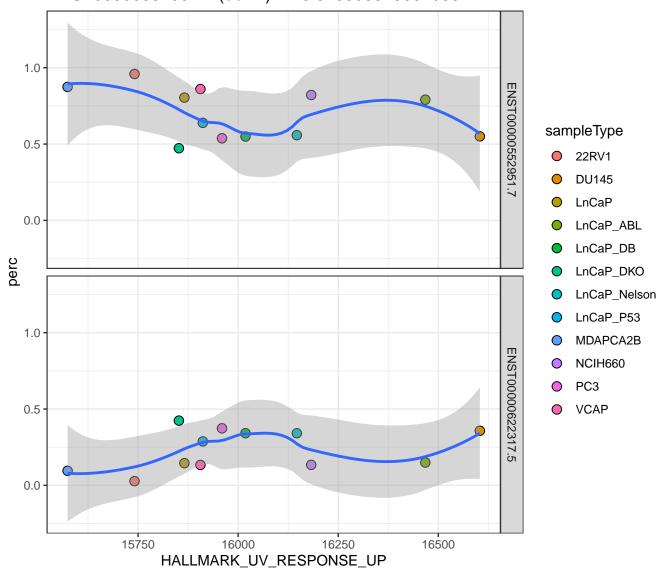
0.00

15000

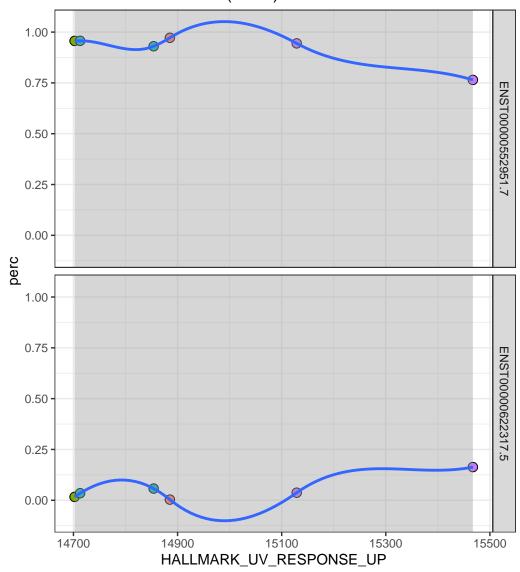
15300

MTOR_UP.N4.V1_UP

CellLine
Gene = NOPCHAP1
Target = HALLMARK_UV_RESPONSE_UP
ENST00000622317.5 (up) = 0.304119701989685
ENST00000552951.7 (down) = -0.326303940667086



LTL_longitudinal
Gene = NOPCHAP1
Target = HALLMARK_UV_RESPONSE_UP
ENST00000622317.5 (up) = 0.834076804089033
ENST00000552951.7 (down) = -0.861438914680929



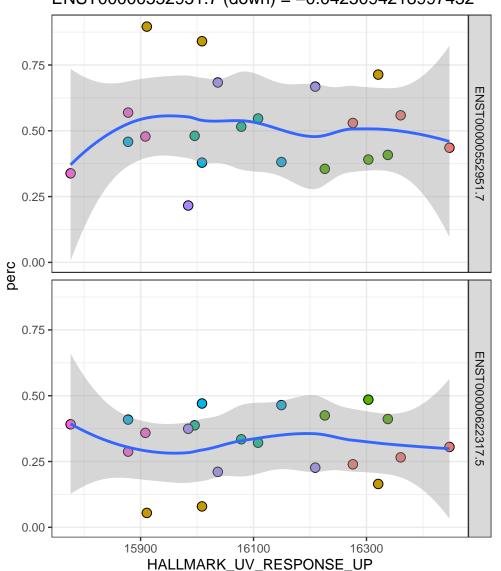
sampleType

LTL331_Cx_12w

LTL331_Cx_8w LTL331_Cx_R

LTL331_PreCx

PDX_longitudinal Gene = NOPCHAP1 Target = HALLMARK_UV_RESPONSE_UP ENST00000622317.5 (up) = 0.0141061831232131 ENST00000552951.7 (down) = -0.0425094218997432

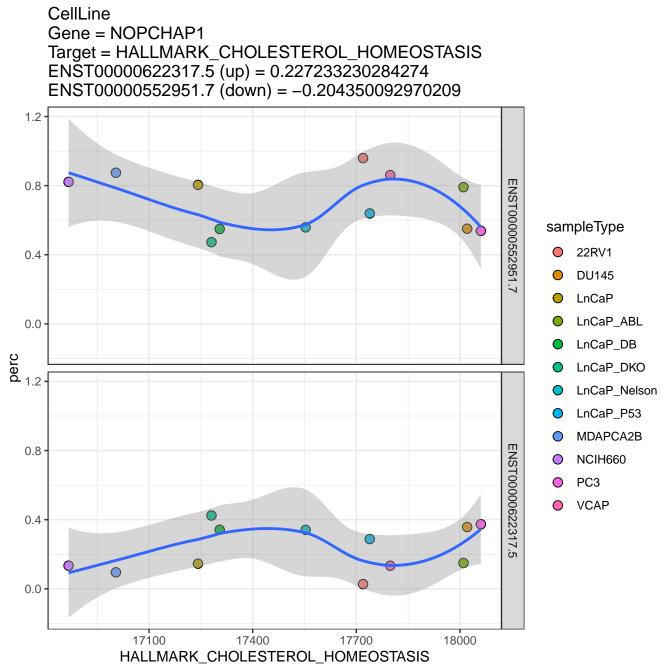


sampleType

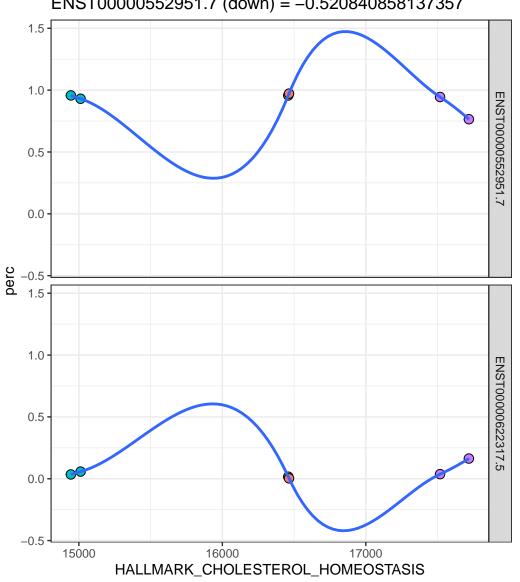
- LnCaP
- LnCaP_ABL

LnCaP_DKO_CX

- LnCaP_DKO_ENZA
- LnCaP_DKO_PreCXLuCaP_147
- LuCaP_147_CX



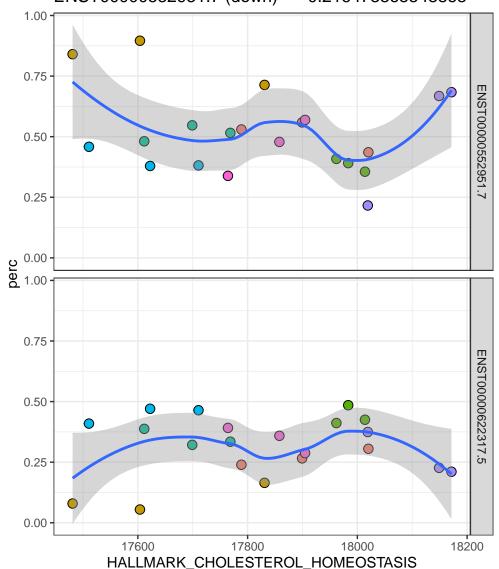
LTL_longitudinal
Gene = NOPCHAP1
Target = HALLMARK_CHOLESTEROL_HOMEOSTASIS
ENST00000622317.5 (up) = 0.417193853662553
ENST00000552951.7 (down) = -0.520840858137357



sampleType

- LTL331_Cx_12wLTL331_Cx_8w
- LTL331_Cx_R
 -) LTL331_PreCx

PDX_longitudinal Gene = NOPCHAP1 Target = HALLMARK_CHOLESTEROL_HOMEOSTASIS ENST00000622317.5 (up) = 0.0863102469983777 ENST00000552951.7 (down) = -0.216473365543868



sampleType

- LnCaP
- LnCaP_ABL

LnCaP_DKO_CX

- LnCaP_DKO_ENZA
- LnCaP_DKO_PreCX
 - LuCaP_147
- LuCaP_147_CX

CellLine Gene = NOPCHAP1 Target = SRC_UP.V1_UP ENST00000622317.5 (up) = -0.0642405728201111ENST00000552951.7 (down) = 0.03779923117505961.0 ENST00000552951.7 0 sampleType 0.5 22RV1 DU145 LnCaP 0.0 LnCaP_ABL LnCaP_DB LnCaP_DKO LnCaP_Nelson LnCaP_P53 1.0 MDAPCA2B ENST00000622317.5 NCIH660 PC3 0.5 **VCAP** 00 0 0.0

8500

perc

7500

8000

SRC_UP.V1_UP

LTL_longitudinal Gene = NOPCHAP1 Target = SRC_UP.V1_UP ENST00000622317.5 (up) = -0.0964567374131289 ENST00000552951.7 (down) = 0.2198749630383561.00 -0.75 ENST00000552951.7 0.50 -0.25 sampleType LTL331_Cx_12w 0.00 d 1.00 LTL331_Cx_8w LTL331_Cx_R LTL331_PreCx 0.75 ENST00000622317.5 0.50

0.25

0.00

7700

7800

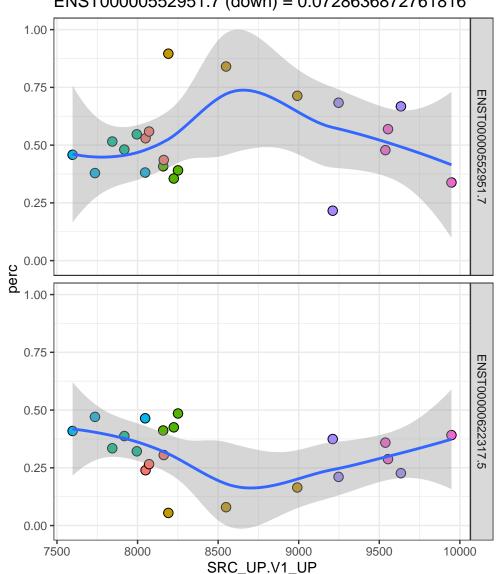
7900

SRC_UP.V1_UP

8000

8100

PDX_longitudinal Gene = NOPCHAP1 Target = SRC_UP.V1_UP ENST00000622317.5 (up) = -0.223670627034703 ENST00000552951.7 (down) = 0.0728636872761816



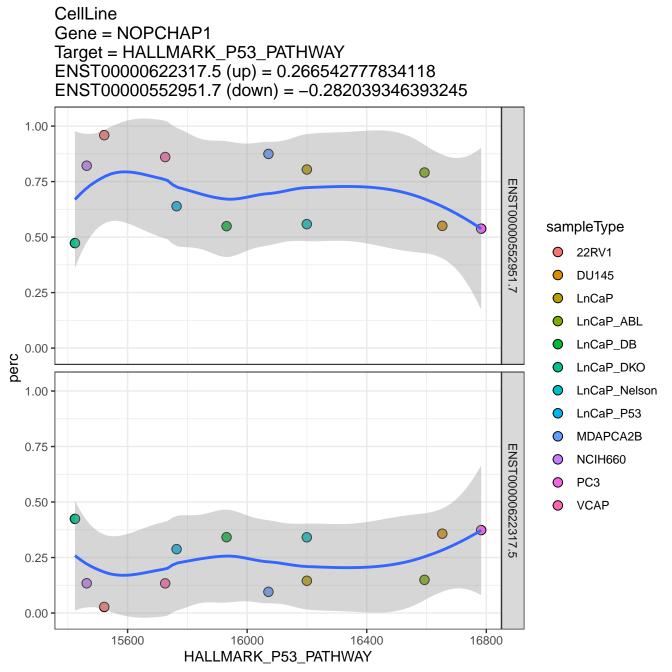
sampleType

- LnCaP
- LnCaP_ABL
- LnCaP_DKO_ENZA

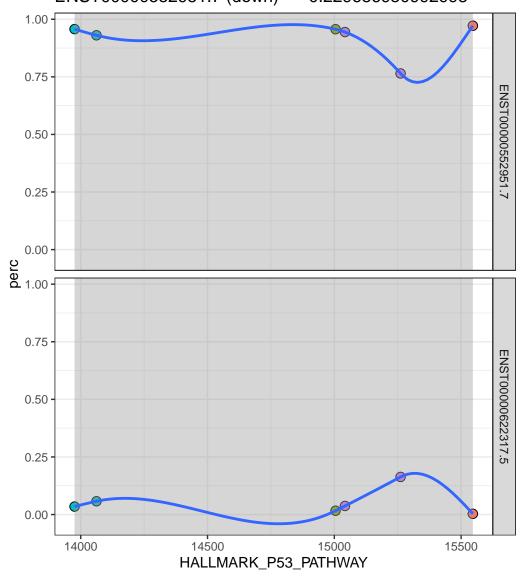
LnCaP_DKO_CX

LnCaP_DKO_PreCX

- LuCaP_147
- LuCaP_147_CX



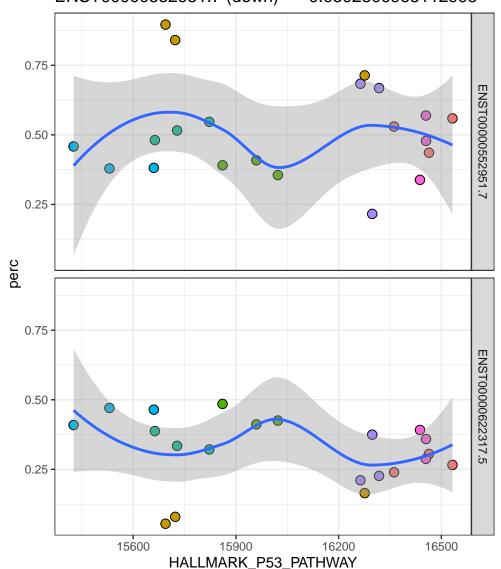
LTL_longitudinal
Gene = NOPCHAP1
Target = HALLMARK_P53_PATHWAY
ENST00000622317.5 (up) = 0.0761693277445701
ENST00000552951.7 (down) = -0.229686950992998



sampleType

LTL331_Cx_12w

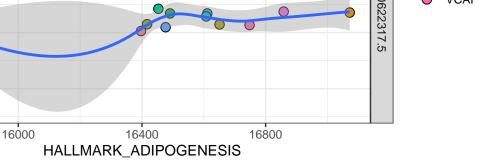
LTL331_Cx_8w LTL331_Cx_R LTL331_PreCx PDX_longitudinal Gene = NOPCHAP1 Target = HALLMARK_P53_PATHWAY ENST00000622317.5 (up) = -0.213770725557526 ENST00000552951.7 (down) = -0.0502800968112995



sampleType

- LnCaP
- LnCaP_ABL
- LnCaP_DKO_CXLnCaP_DKO_ENZA
- LnCaP_DKO_PreCX
- LuCaP_147
- LuCaP_147_CX

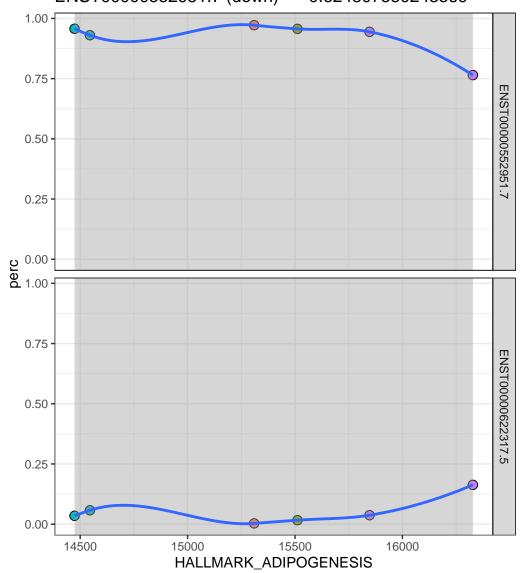
CellLine Gene = NOPCHAP1 Target = HALLMARK_ADIPOGENESIS ENST00000622317.5 (up) = 0.419573147164392 ENST00000552951.7 (down) = -0.389684549294756 2 . ENST00000552951.7 sampleType 0 22RV1 0 **DU145** LnCaP LnCaP_ABL -1 LnCaP_DB perc LnCaP_DKO LnCaP_Nelson LnCaP_P53 2. MDAPCA2B ENST00000622317.5 NCIH660 PC3 1 **VCAP**



0

-1

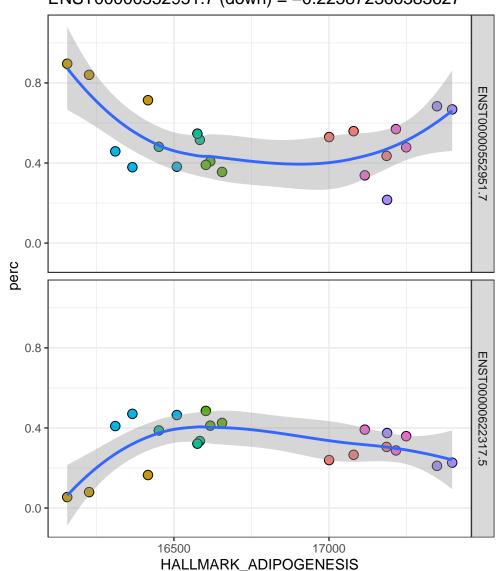
LTL_longitudinal
Gene = NOPCHAP1
Target = HALLMARK_ADIPOGENESIS
ENST00000622317.5 (up) = 0.517662386240785
ENST00000552951.7 (down) = -0.624807580243569



sampleType

LTL331_Cx_12w

LTL331_Cx_8w LTL331_Cx_R LTL331_PreCx PDX_longitudinal Gene = NOPCHAP1 Target = HALLMARK_ADIPOGENESIS ENST00000622317.5 (up) = 0.0201036028432354 ENST00000552951.7 (down) = -0.225872366383627

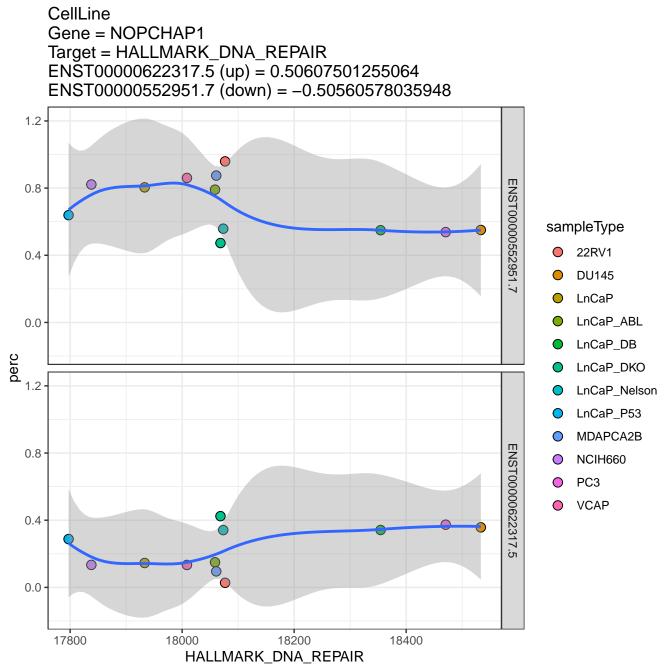


sampleType

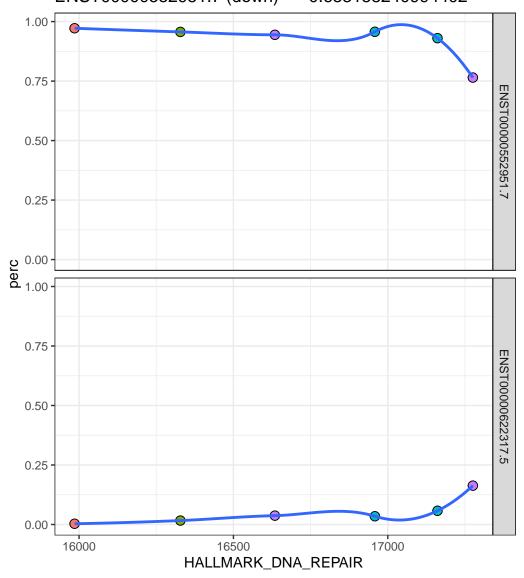
- LnCaP
- LnCaP_ABL

LnCaP_DKO_CX

- LnCaP_DKO_ENZA
- LnCaP_DKO_PreCX
- LuCaP_147LuCaP_147_CX
- ____



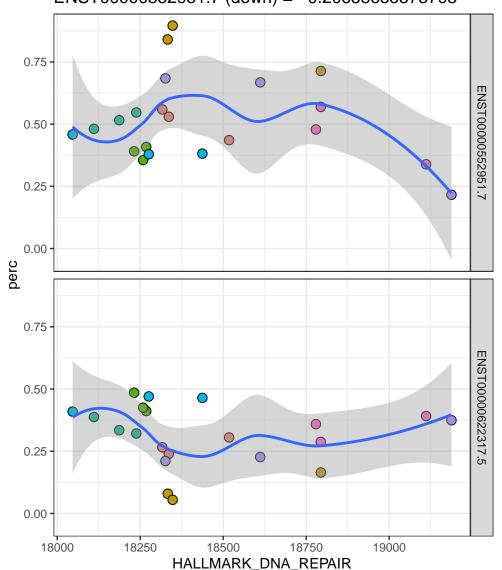
LTL_longitudinal
Gene = NOPCHAP1
Target = HALLMARK_DNA_REPAIR
ENST00000622317.5 (up) = 0.769371773536922
ENST00000552951.7 (down) = -0.653158240994402



sampleType

LTL331_Cx_12w

LTL331_Cx_8w LTL331_Cx_R LTL331_PreCx PDX_longitudinal Gene = NOPCHAP1 Target = HALLMARK_DNA_REPAIR ENST00000622317.5 (up) = -0.0772450315549626 ENST00000552951.7 (down) = -0.20658655575798



sampleType

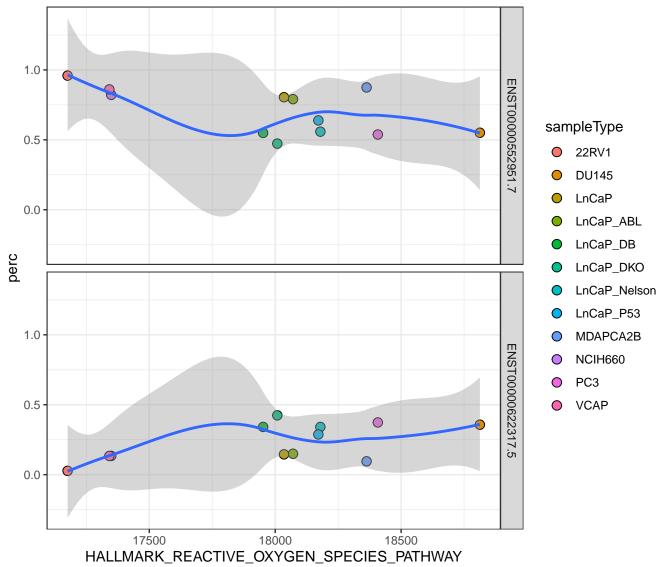
• LnCaP

LnCaP_ABL

LuCaP_147 LuCaP_147_CX

LnCaP_DKO_CX

LnCaP_DKO_ENZA LnCaP_DKO_PreCX CellLine
Gene = NOPCHAP1
Target = HALLMARK_REACTIVE_OXYGEN_SPECIES_PATHWAY
ENST00000622317.5 (up) = 0.601863692389627
ENST00000552951.7 (down) = -0.614917089376127



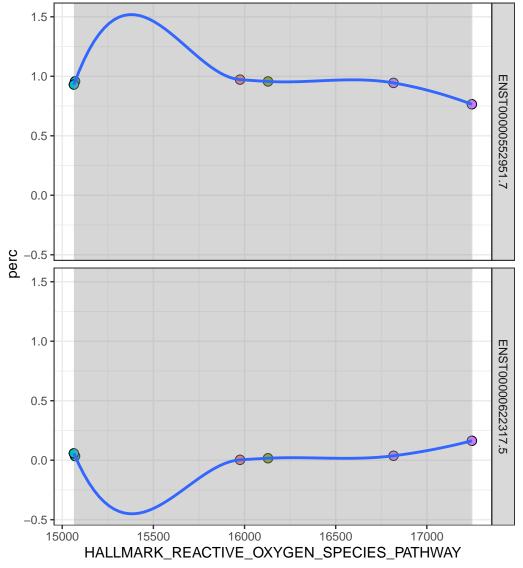
LTL_longitudinal
Gene = NOPCHAP1
Target = HALLMARK_REACTIVE_OXYGEN_SPECIES_PATHWAY
ENST00000622317.5 (up) = 0.523349996022222
ENST00000552951.7 (down) = -0.620186197953161

sampleType

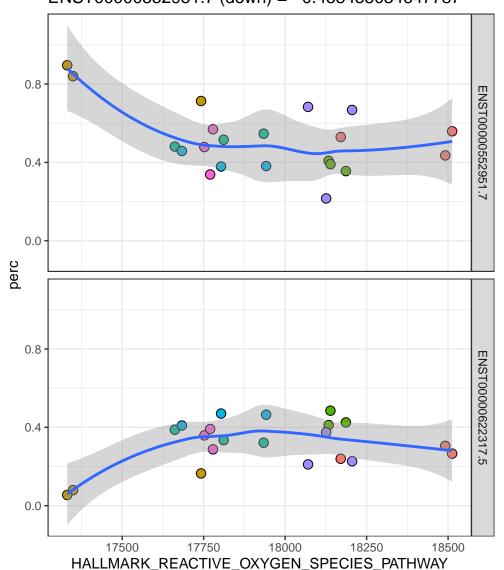
LTL331_Cx_12w

LTL331_Cx_8w

LTL331_Cx_R LTL331_PreCx



PDX_longitudinal
Gene = NOPCHAP1
Target = HALLMARK_REACTIVE_OXYGEN_SPECIES_PATHWAY
ENST00000622317.5 (up) = 0.321649722647872
ENST00000552951.7 (down) = -0.458455034647787

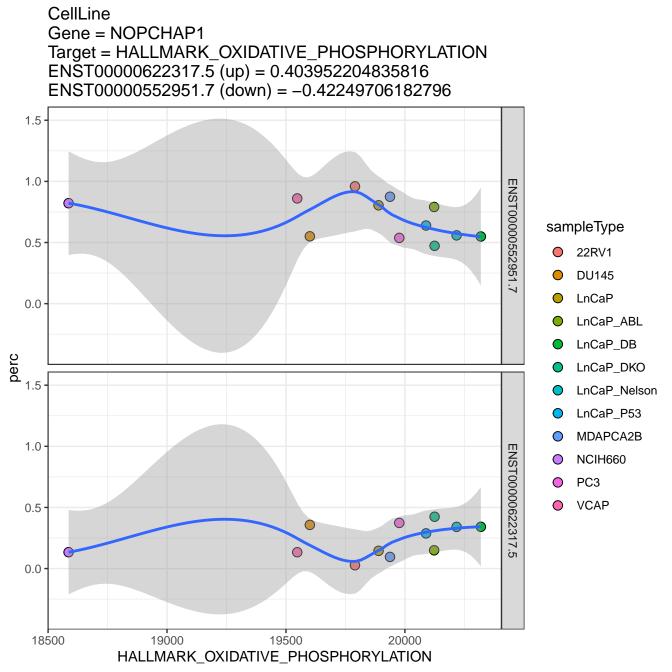


sampleType

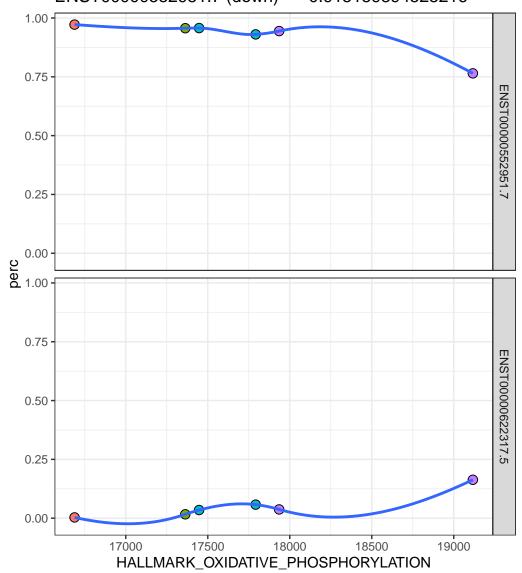
- LnCaP
- LnCaP_ABL

LnCaP_DKO_CX

- LnCaP_DKO_ENZA
- LnCaP_DKO_PreCX
- LuCaP_147
- LuCaP_147_CX



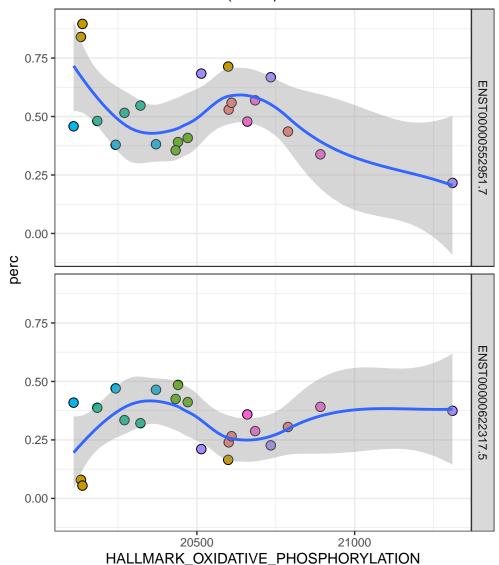
LTL_longitudinal
Gene = NOPCHAP1
Target = HALLMARK_OXIDATIVE_PHOSPHORYLATION
ENST00000622317.5 (up) = 0.946878584366554
ENST00000552951.7 (down) = -0.915159394525216



sampleType

LTL331_Cx_12w

LTL331_Cx_8w LTL331_Cx_R LTL331_PreCx PDX_longitudinal Gene = NOPCHAP1 Target = HALLMARK_OXIDATIVE_PHOSPHORYLATION ENST00000622317.5 (up) = 0.0879223184802229 ENST00000552951.7 (down) = -0.425472443067249



sampleType

- LnCaP
- LnCaP_ABL

LnCaP_DKO_CX

- LnCaP_DKO_ENZA
- LnCaP_DKO_PreCX
 - LuCaP_147
- LuCaP_147_CX

CellLine Gene = NOPCHAP1 Target = HALLMARK_FATTY_ACID_METABOLISM ENST00000622317.5 (up) = 0.106553686409925ENST00000552951.7 (down) = -0.1095458711918971.2 0 0.8 ENST00000552951.7 sampleType 0.4 22RV1 DU145 LnCaP 0.0 LnCaP_ABL LnCaP_DB perc LnCaP_DKO 1.2 LnCaP_Nelson LnCaP_P53 MDAPCA2B 8.0 ENST00000622317.5 NCIH660 PC3 **VCAP** 0.4 0.0

15250

15500

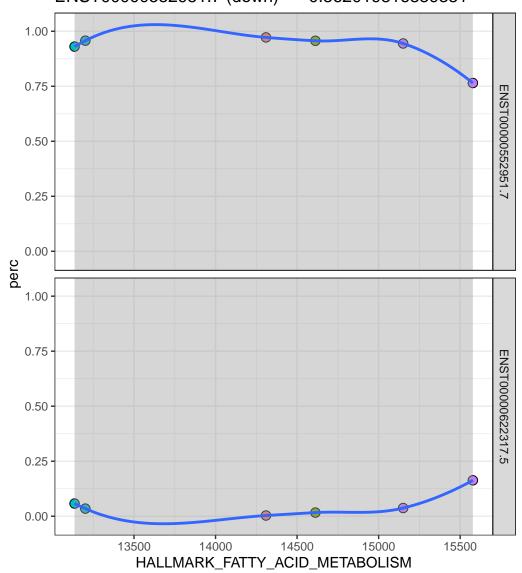
15750

HALLMARK_FATTY_ACID_METABOLISM

16000

16250

LTL_longitudinal
Gene = NOPCHAP1
Target = HALLMARK_FATTY_ACID_METABOLISM
ENST00000622317.5 (up) = 0.453143146887528
ENST00000552951.7 (down) = -0.562019816859381

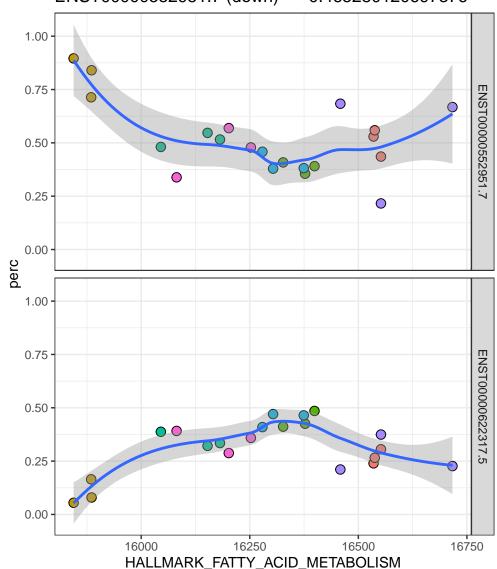


sampleType

LTL331_Cx_12w

LTL331_Cx_8w

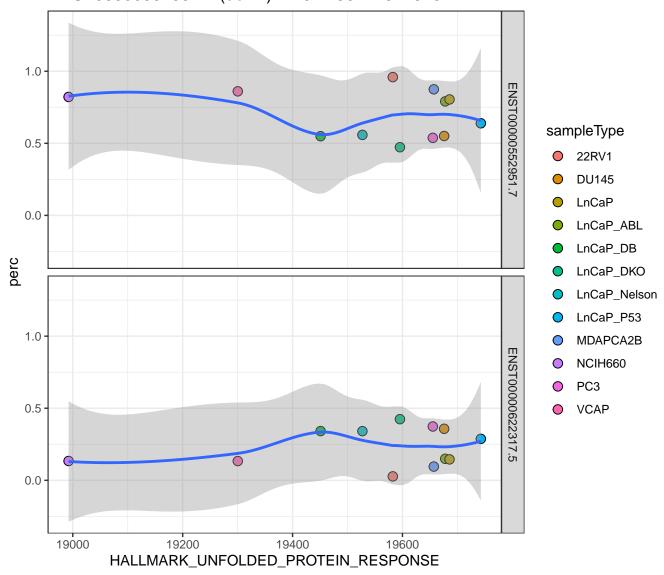
LTL331_Cx_R LTL331_PreCx PDX_longitudinal Gene = NOPCHAP1 Target = HALLMARK_FATTY_ACID_METABOLISM ENST00000622317.5 (up) = 0.35589362359285 ENST00000552951.7 (down) = -0.465280120697376



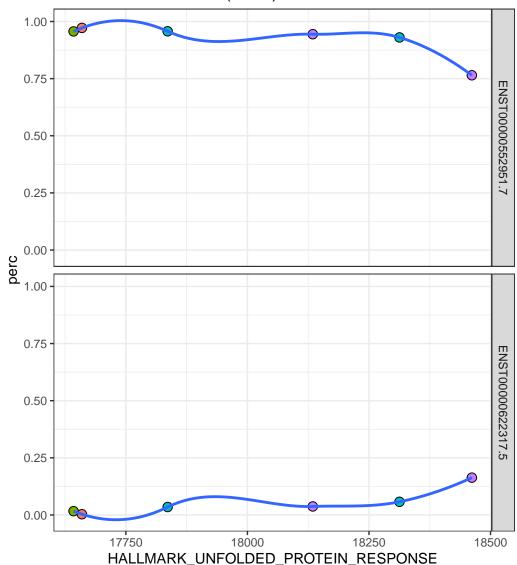
sampleType

- LnCaP
- LnCaP_ABL
- LnCaP_DKO_CXLnCaP_DKO_ENZA
- LnCaP_DKO_PreCX
- LuCaP_147
- LuCaP_147_CX

CellLine
Gene = NOPCHAP1
Target = HALLMARK_UNFOLDED_PROTEIN_RESPONSE
ENST00000622317.5 (up) = 0.23381022753098
ENST00000552951.7 (down) = -0.24361148415292



LTL_longitudinal
Gene = NOPCHAP1
Target = HALLMARK_UNFOLDED_PROTEIN_RESPONSE
ENST00000622317.5 (up) = 0.835605562732749
ENST00000552951.7 (down) = -0.758493479817635

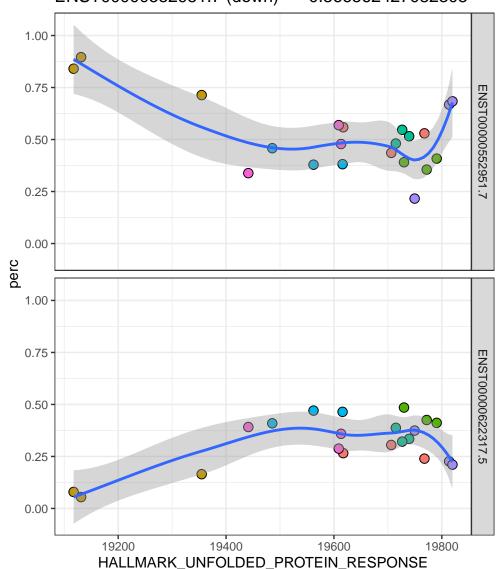


sampleType

LTL331_Cx_12w

LTL331_Cx_8w

LTL331_Cx_R LTL331_PreCx PDX_longitudinal
 Gene = NOPCHAP1
 Target = HALLMARK_UNFOLDED_PROTEIN_RESPONSE
 ENST00000622317.5 (up) = 0.541745393368909
 ENST00000552951.7 (down) = -0.566562427082303

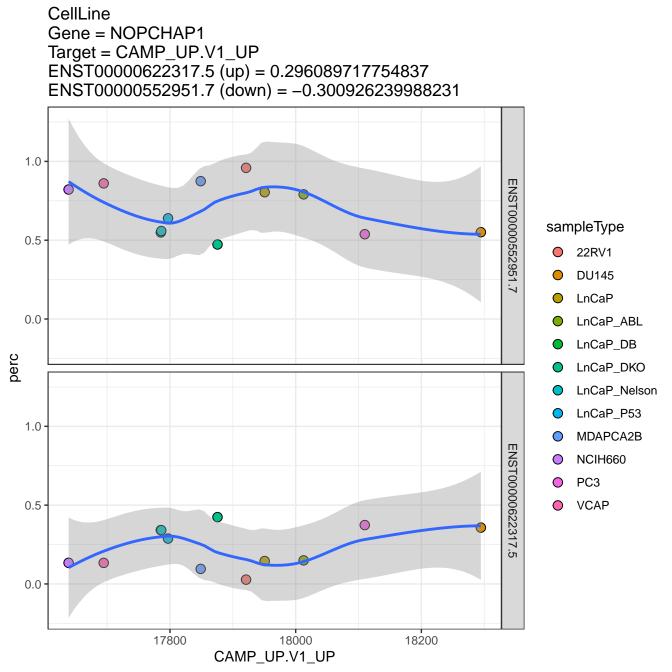


sampleType

- LnCaP
- LnCaP_ABL

LnCaP_DKO_CX

- LnCaP_DKO_ENZA
- LnCaP_DKO_PreCXLuCaP_147
- U LuCaP
- LuCaP_147_CX



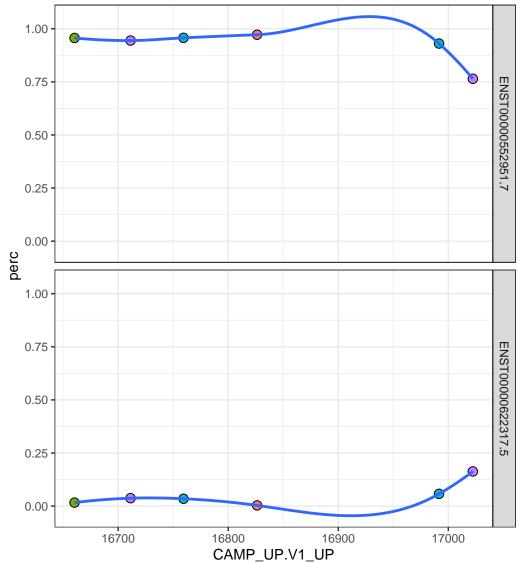
LTL_longitudinal
Gene = NOPCHAP1
Target = CAMP_UP.V1_UP
ENST00000622317.5 (up) = 0.734279027005523
ENST00000552951.7 (down) = -0.6927959940831

sampleType

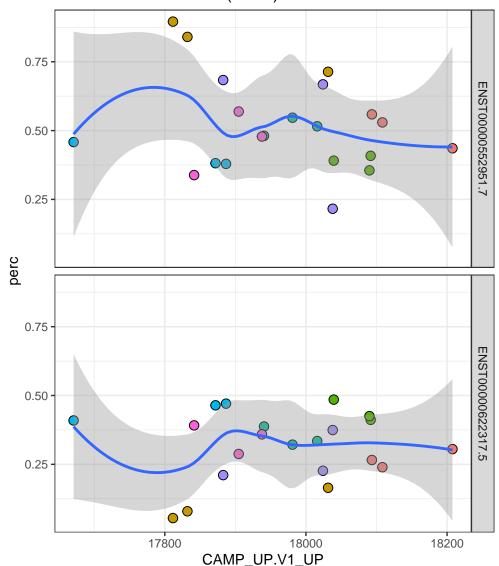
LTL331_Cx_12w

LTL331_Cx_8w LTL331_Cx_R

LTL331_PreCx

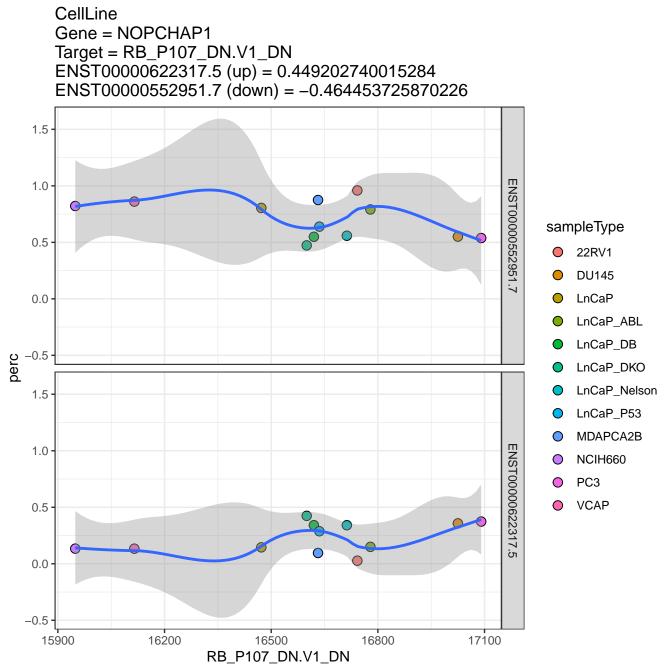


PDX_longitudinal
Gene = NOPCHAP1
Target = CAMP_UP.V1_UP
ENST00000622317.5 (up) = 0.0799379986308909
ENST00000552951.7 (down) = -0.255138525765996

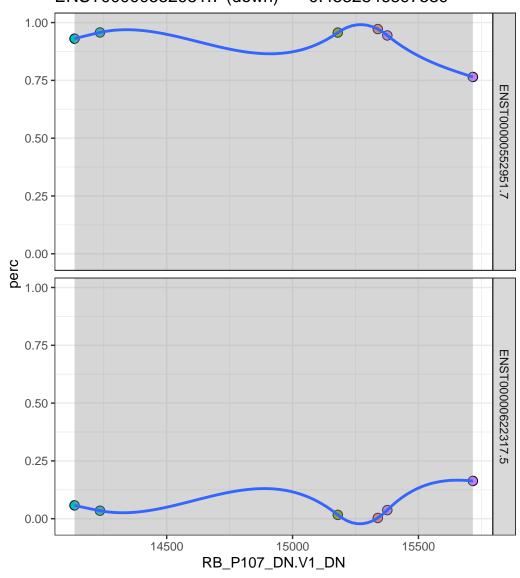


sampleType

- LnCaP
- LnCaP_ABL
- LnCaP_DKO_CXLnCaP_DKO_ENZA
- LnCaP_DKO_PreCX
- LuCaP_147
- LuCaP_147_CX



LTL_longitudinal Gene = NOPCHAP1 Target = RB_P107_DN.V1_DN ENST00000622317.5 (up) = 0.316883563846639 ENST00000552951.7 (down) = -0.4532846897539



sampleType

LTL331_Cx_12w

LTL331_Cx_8w

LTL331_Cx_R LTL331_PreCx

