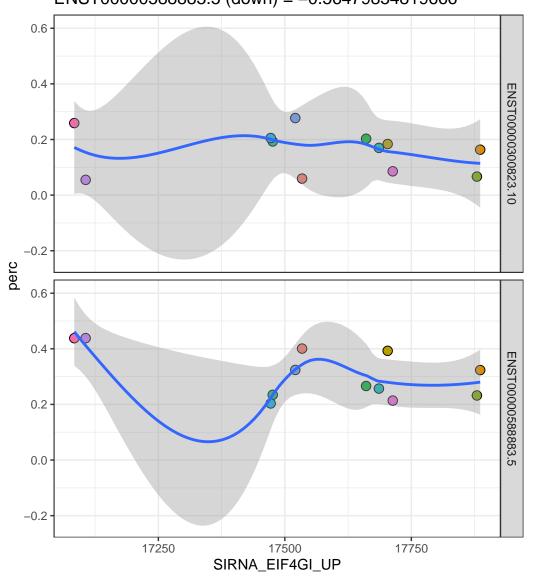
CellLine Gene = ZNF226 Target = SIRNA_EIF4GI_UP ENST00000300823.10 (up) = -0.181596023516573 ENST00000588883.5 (down) = -0.56479854819666

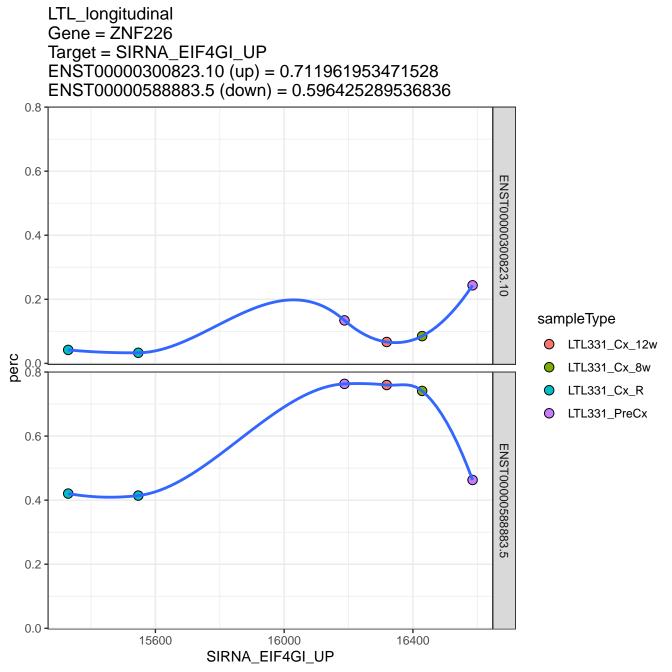


- 22RV1 DU145
- LnCaP
- LnCaP_ABL
- LnCaP_DB
- LnCaP_DKO

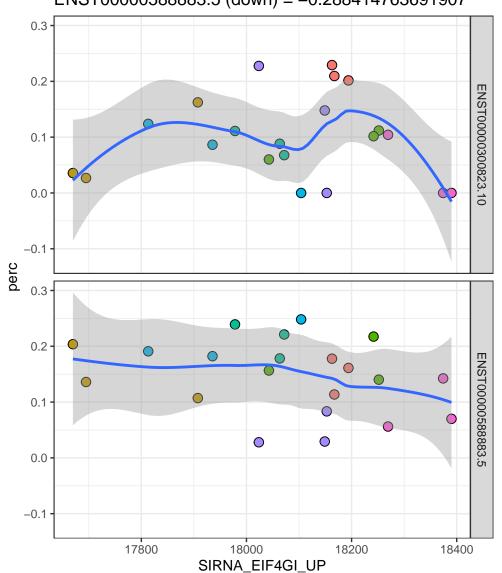
MDAPCA2B

- LnCaP_DKO

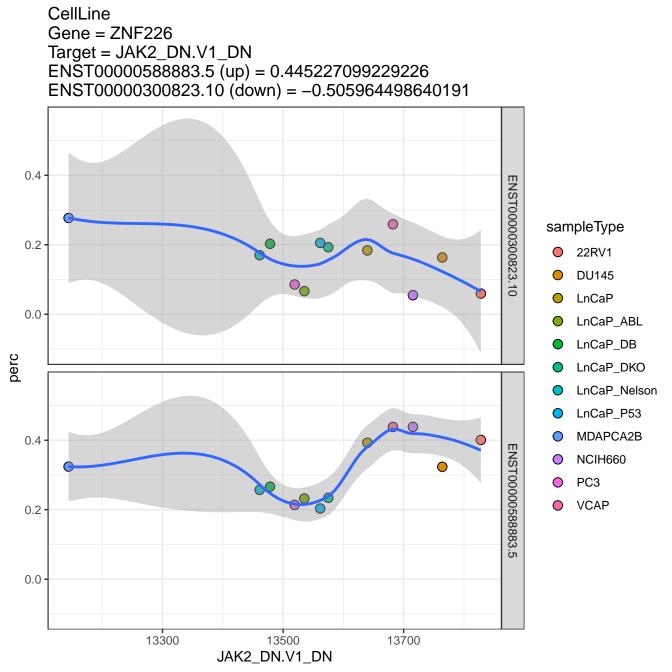
 LnCaP_Nelson
- _____ **O** LnCaP_P53
- O NCIH660
- F
- PC3 VCAP



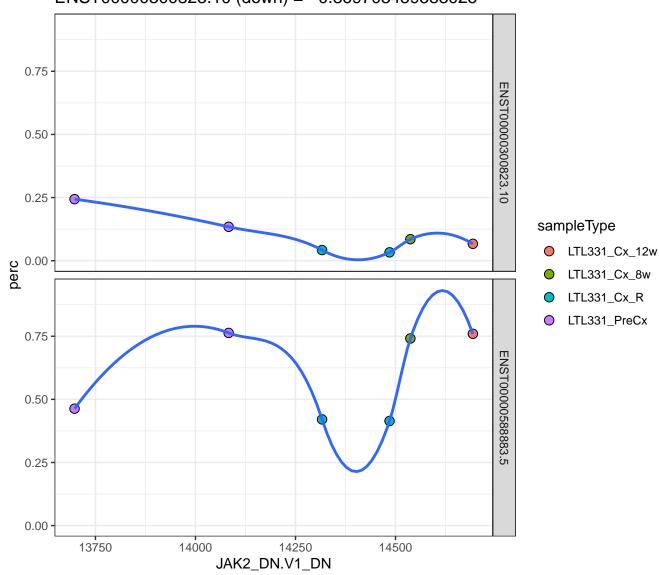
PDX_longitudinal Gene = ZNF226 Target = SIRNA_EIF4GI_UP ENST00000300823.10 (up) = 0.00402178532999899 ENST00000588883.5 (down) = -0.288414763691907

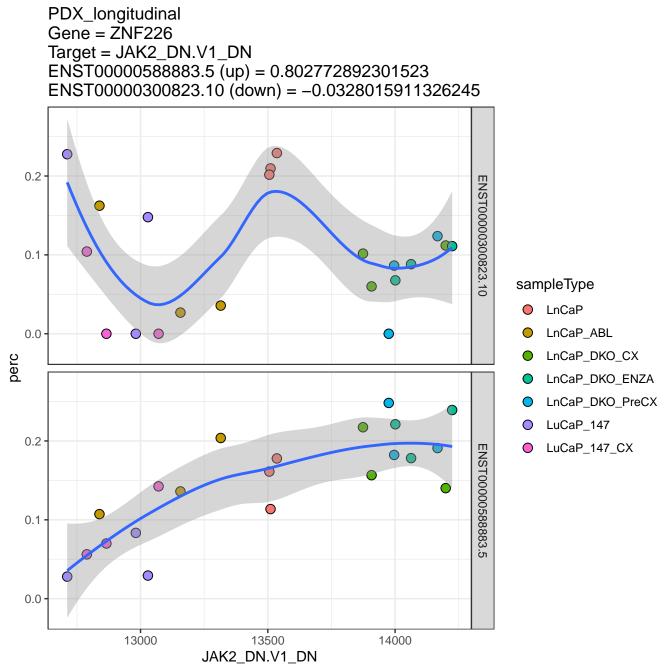


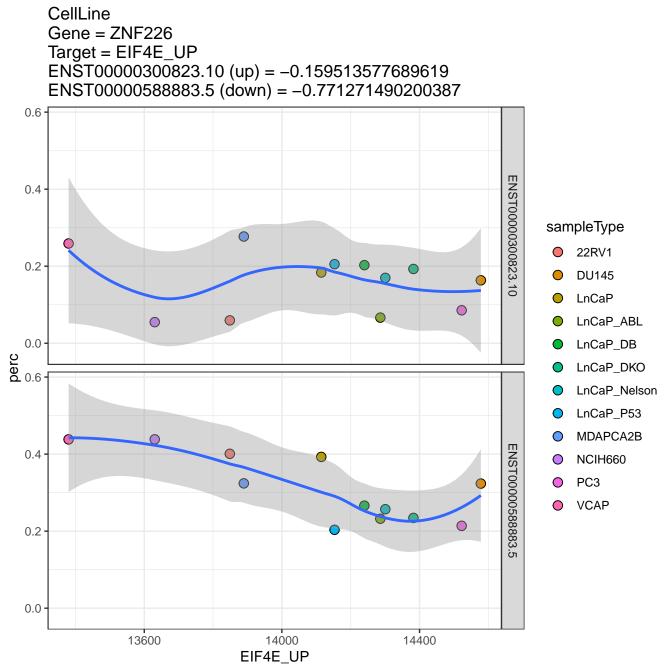
- LnCaP
- LnCaP_ABL
- LnCaP_DKO_CXLnCaP_DKO_ENZA
- LnCaP_DKO_PreCX
- LuCaP_147
- LuCaP_147_CX



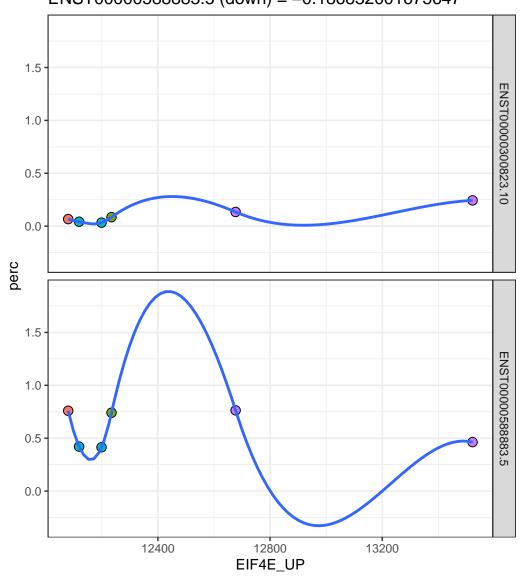
LTL_longitudinal
Gene = ZNF226
Target = JAK2_DN.V1_DN
ENST00000588883.5 (up) = 0.330408219967004
ENST00000300823.10 (down) = -0.869703459583025





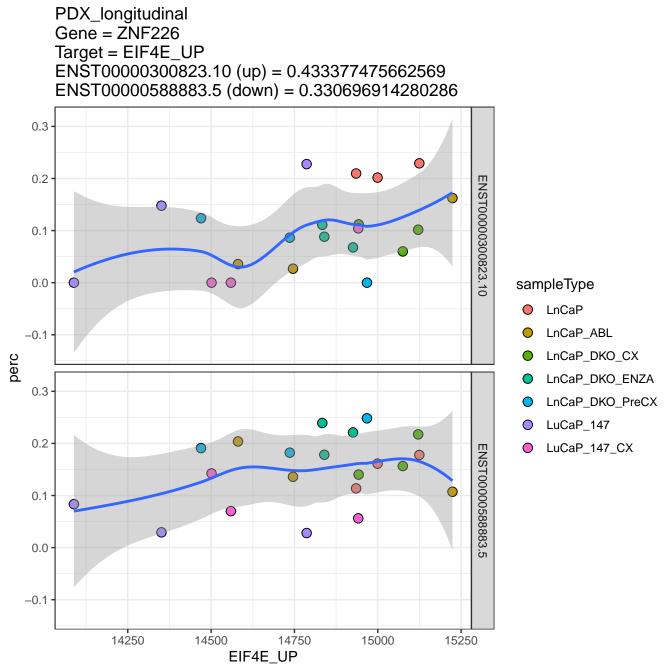


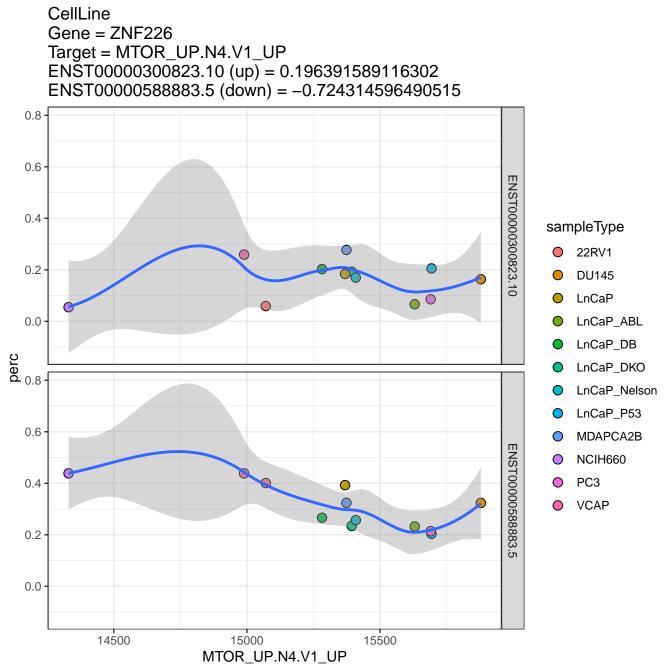
LTL_longitudinal
Gene = ZNF226
Target = EIF4E_UP
ENST00000300823.10 (up) = 0.97171696537653
ENST00000588883.5 (down) = -0.186832001675647



LTL331_Cx_12w

LTL331_Cx_8w LTL331_Cx_R LTL331_PreCx





LTL_longitudinal Gene = ZNF226 Target = MTOR_UP.N4.V1_UP ENST00000300823.10 (up) = 0.793212264071985 ENST00000588883.5 (down) = 0.455866769380865 0.8 ENST00000300823.10 0.6 0.4 0.2 sampleType LTL331_Cx_12w 0.0 perc LTL331_Cx_8w LTL331_Cx_R 8.0 LTL331_PreCx 0.6 ENST00000588883.5 0.4 0.2 0.0

13500

13750

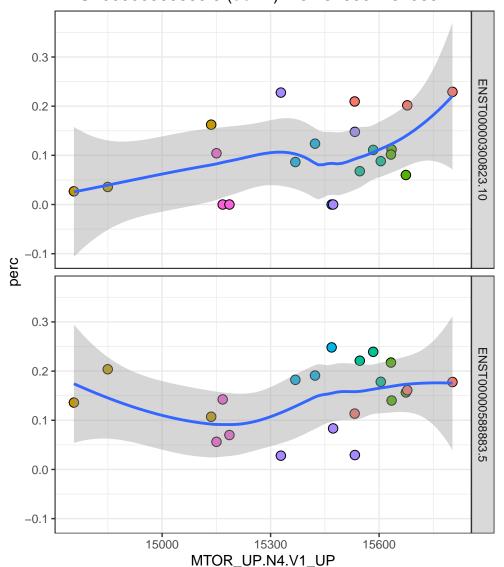
12750

13000

13250

MTOR_UP.N4.V1_UP

PDX_longitudinal Gene = ZNF226 Target = MTOR_UP.N4.V1_UP ENST00000300823.10 (up) = 0.420040444456134 ENST00000588883.5 (down) = 0.234353213158974



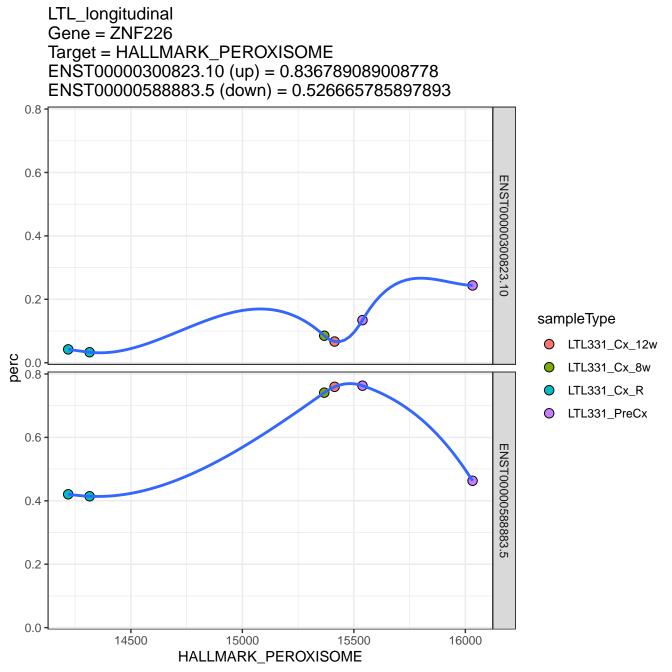
sampleType

- LnCaP
- LnCaP_ABL

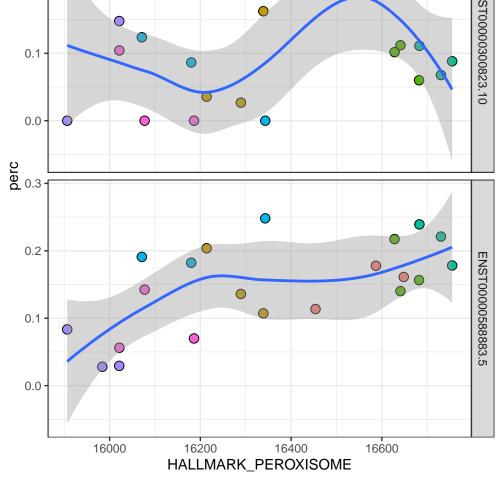
LnCaP_DKO_CX

- LnCaP_DKO_ENZA
- LnCaP_DKO_PreCX
- LuCaP_147
- LuCaP_147_CX

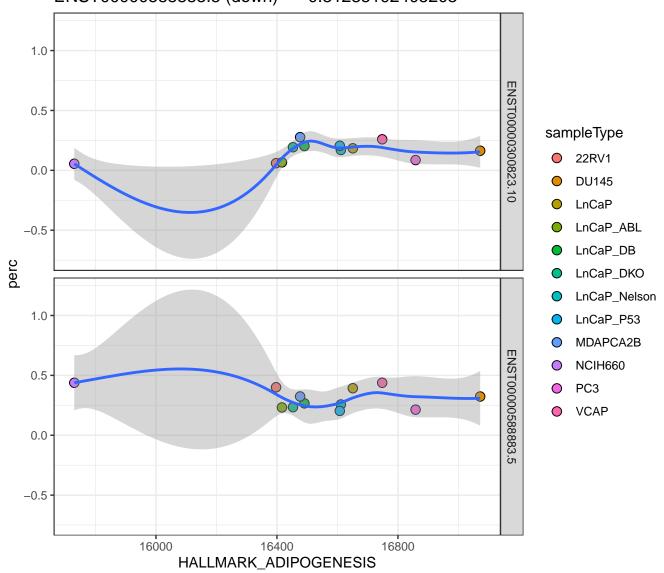
CellLine Gene = ZNF226 Target = HALLMARK_PEROXISOME ENST00000300823.10 (up) = 0.541290194927242 ENST00000588883.5 (down) = -0.0745329360525349 0.6 ENST00000300823.10 0.4 sampleType 0.2 22RV1 **DU145** LnCaP 0.0 LnCaP_ABL LnCaP_DB perc LnCaP_DKO LnCaP_Nelson 0.6 LnCaP_P53 MDAPCA2B ENST00000588883.5 0.4 NCIH660 PC3 **VCAP** 0.2 0.0 15500 16000 16500 HALLMARK_PEROXISOME

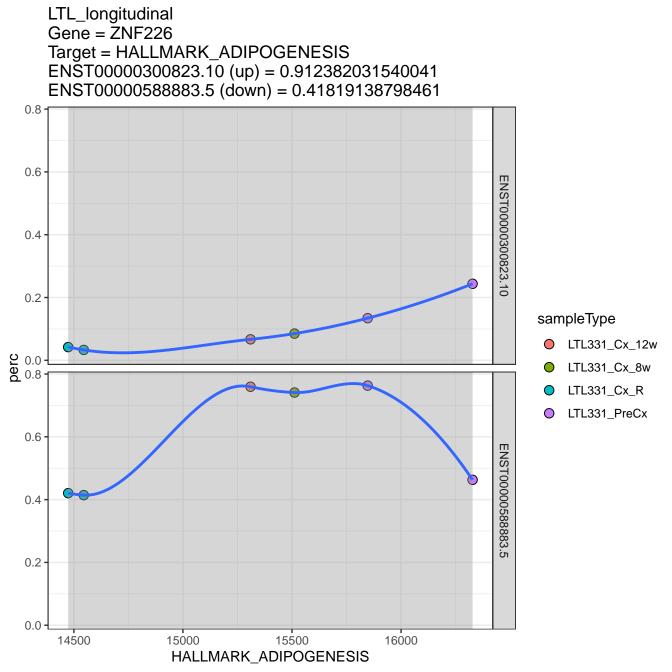


PDX_longitudinal Gene = ZNF226 Target = HALLMARK_PEROXISOME ENST00000300823.10 (up) = 0.191442193014702 ENST00000588883.5 (down) = 0.609270881531483 0.3 0 0.2 ENST00000300823.10 0.1 sampleType LnCaP 0 0.0 0 LnCaP_ABL LnCaP_DKO_CX LnCaP_DKO_ENZA LnCaP_DKO_PreCX LuCaP_147 0.2 LuCaP_147_CX

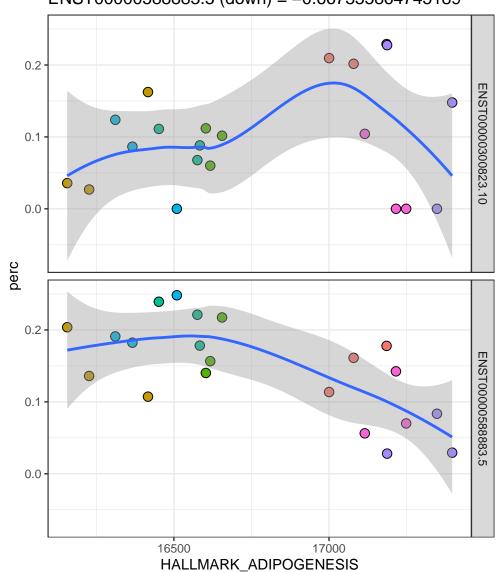


CellLine
Gene = ZNF226
Target = HALLMARK_ADIPOGENESIS
ENST00000300823.10 (up) = 0.380169089485977
ENST00000588883.5 (down) = -0.31235192405203





PDX_longitudinal Gene = ZNF226 Target = HALLMARK_ADIPOGENESIS ENST00000300823.10 (up) = 0.215376511014282 ENST00000588883.5 (down) = -0.667355864745189



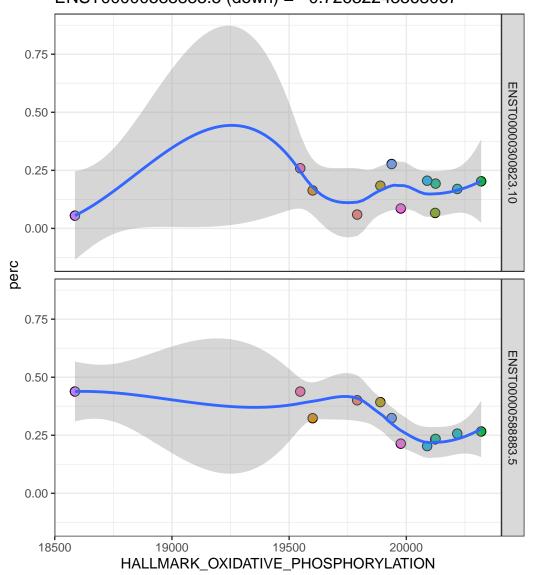
sampleType

- LnCaP
- LnCaP_ABL
- LnCaP_DKO_ENZA
- LnCaP_DKO_PreCX

LnCaP_DKO_CX

- LuCaP_147
 - LuCaP_147_CX

CellLine
Gene = ZNF226
Target = HALLMARK_OXIDATIVE_PHOSPHORYLATION
ENST00000300823.10 (up) = 0.329055813242178
ENST00000588883.5 (down) = -0.72632243368067



sampleType

22RV1

DU145 LnCaP

LnCaP_ABL LnCaP_DB

LnCaP_DKO
LnCaP_Nelson

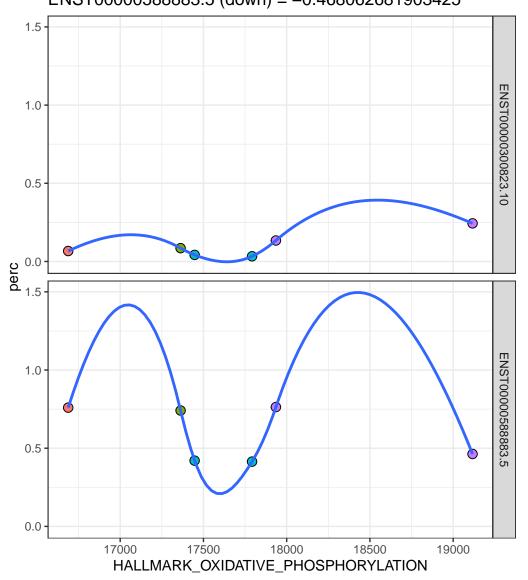
LnCaP_P53 MDAPCA2B

NCIH660

PC3

VCAP

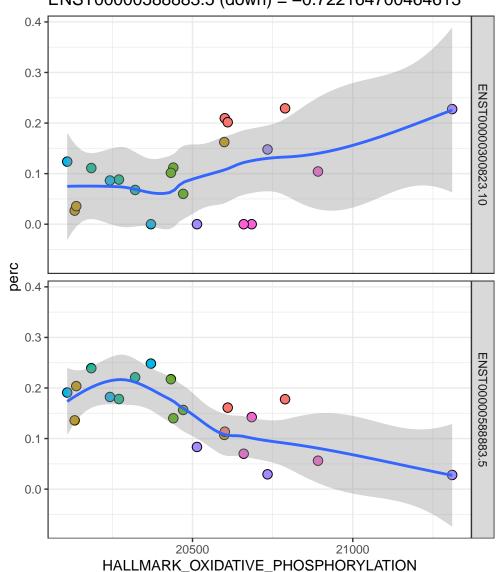
LTL_longitudinal
Gene = ZNF226
Target = HALLMARK_OXIDATIVE_PHOSPHORYLATION
ENST00000300823.10 (up) = 0.812634011009946
ENST00000588883.5 (down) = -0.468062681903425



LTL331_Cx_12w

LTL331_Cx_8w

LTL331_Cx_R LTL331_PreCx PDX_longitudinal Gene = ZNF226 Target = HALLMARK_OXIDATIVE_PHOSPHORYLATION ENST00000300823.10 (up) = 0.467806142451558 ENST00000588883.5 (down) = -0.722164700464613



sampleType

- LnCaP
- LnCaP_ABL

LnCaP_DKO_CX

- LnCaP_DKO_ENZA
- LnCaP_DKO_PreCX
- O LuCaP_147
- LuCaP_147_CX

CellLine Gene = ZNF226 Target = HALLMARK_FATTY_ACID_METABOLISM ENST00000300823.10 (up) = 0.61903992109056 ENST00000588883.5 (down) = -0.358956939823502 0.75 0.50 ENST00000300823.10 sampleType 0.25 -22RV1 **DU145** LnCaP 0.00 LnCaP_ABL LnCaP_DB ည 0.75 LnCaP_DKO LnCaP_Nelson LnCaP_P53 MDAPCA2B 0.50 ENST00000588883.5 NCIH660 PC3 **VCAP** 0.25 -0.00

15250

15500

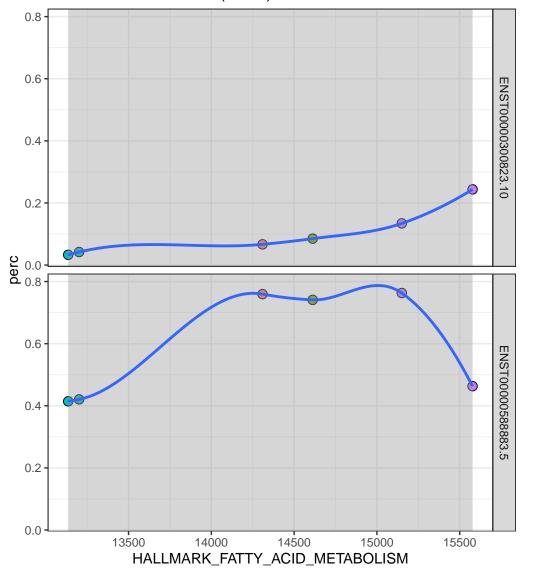
15750

HALLMARK_FATTY_ACID_METABOLISM

16000

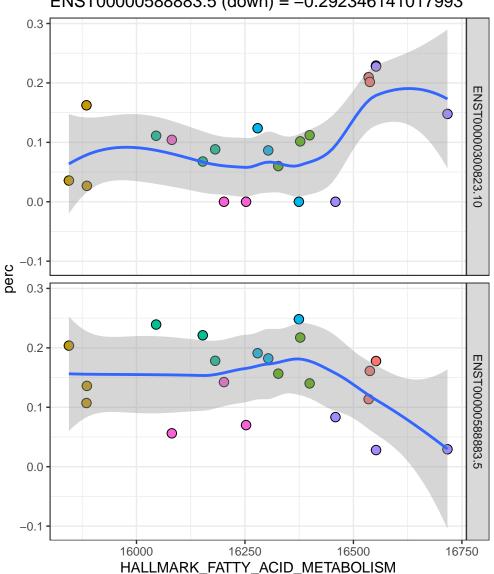
16250

LTL_longitudinal
Gene = ZNF226
Target = HALLMARK_FATTY_ACID_METABOLISM
ENST00000300823.10 (up) = 0.886390331670401
ENST00000588883.5 (down) = 0.477580265610543



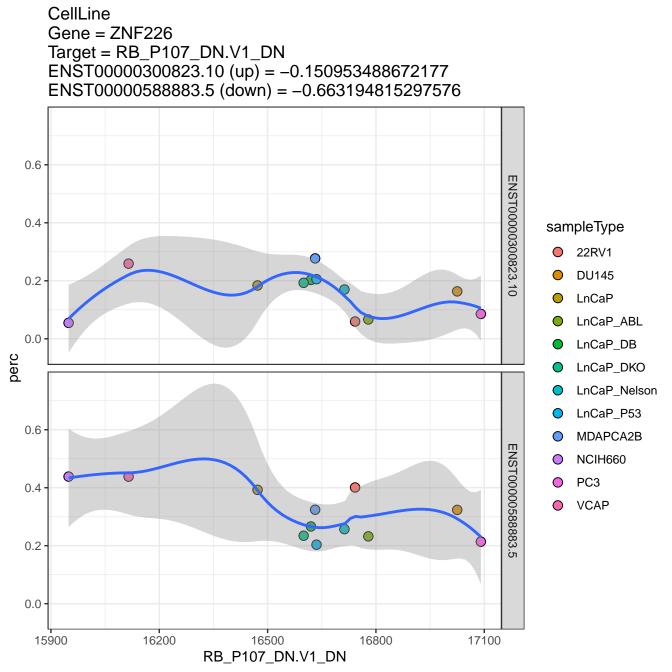
LTL331_Cx_12w

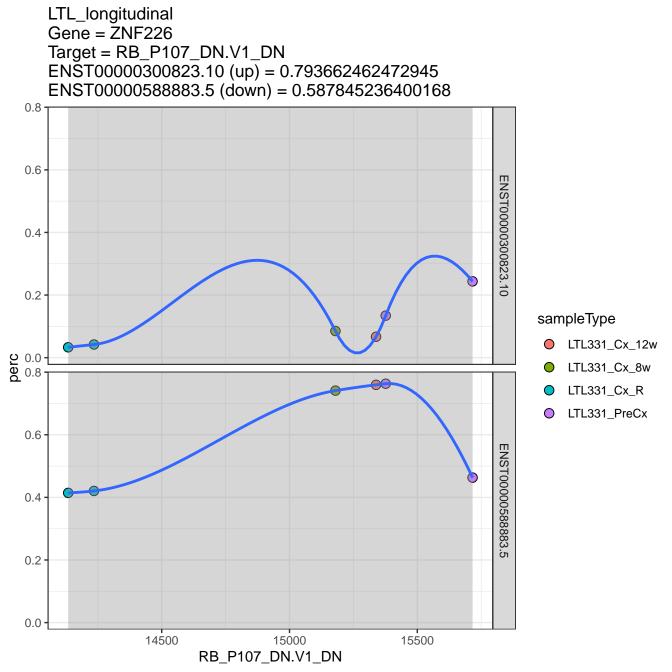
LTL331_Cx_8w LTL331_Cx_R LTL331_PreCx PDX_longitudinal Gene = ZNF226 Target = HALLMARK_FATTY_ACID_METABOLISM ENST00000300823.10 (up) = 0.441719233446378 ENST00000588883.5 (down) = -0.292346141017993



sampleType

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





PDX_longitudinal Gene = ZNF226 Target = RB_P107_DN.V1_DN ENST00000300823.10 (up) = 0.271531578587318 ENST00000588883.5 (down) = 0.0826238230725015 0.3 ENST00000300823.10 0.2 0.1 sampleType LnCaP 0.0 **O** LnCaP_ABL LnCaP_DKO_CX LnCaP_DKO_ENZA 0.3 LnCaP_DKO_PreCX LuCaP_147 LuCaP_147_CX 0.2 ENST00000588883.5 0 0.1 0 0.0 16800 17000 17200 RB_P107_DN.V1_DN