CellLine Gene = WNT2B Target = PID_HEDGEHOG_2PATHWAY ENST00000369684.5 (up) = 0.0800372790471527 ENST00000369686.9 (down) = 0.040796652834631 1 ENST00000369684.5 sampleType 22RV1 0 DU145 LnCaP LnCaP_ABL LnCaP_DB LnCaP_DKO LnCaP_Nelson LnCaP_P53 MDAPCA2B ENST00000369686.9 0 NCIH660

12000

PID_HEDGEHOG_2PATHWAY

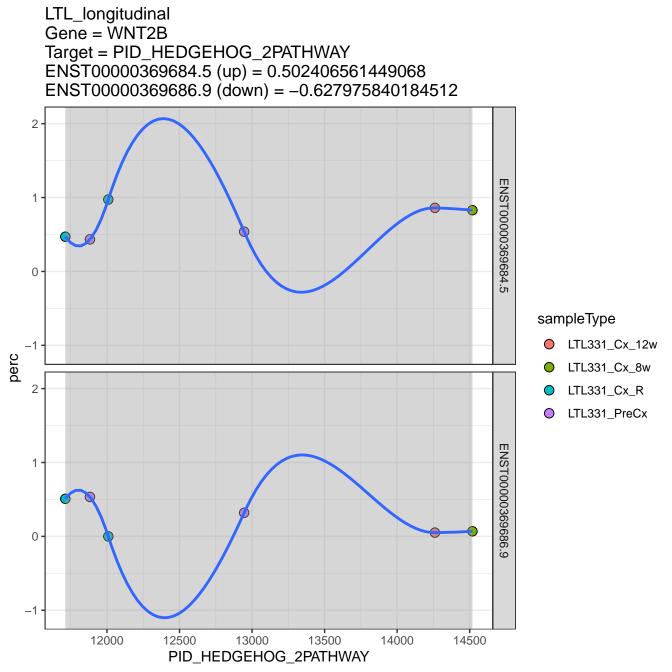
13000

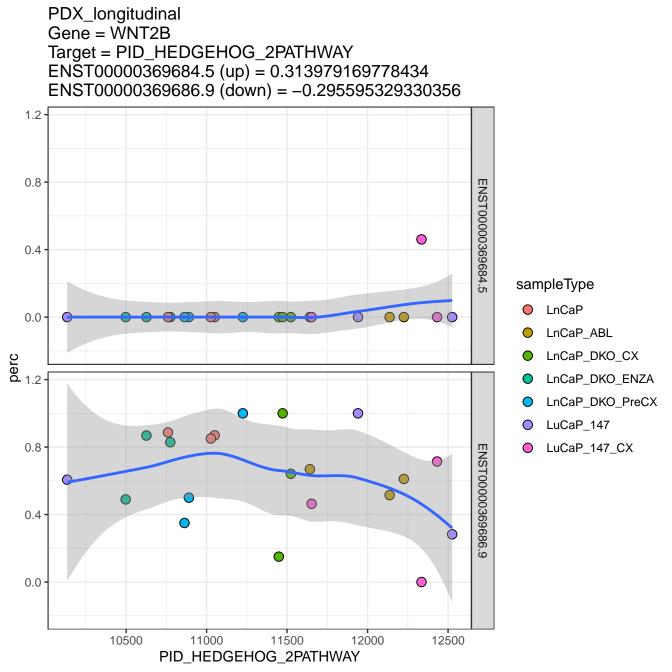
0

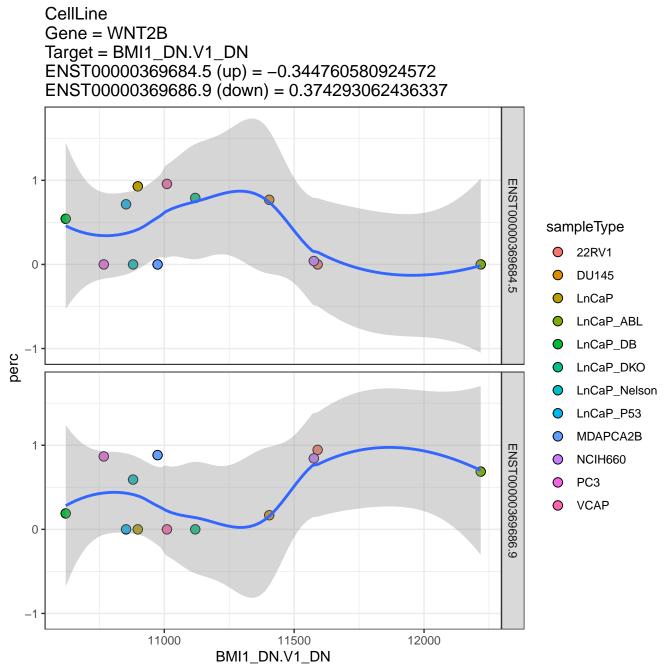
10000

11000

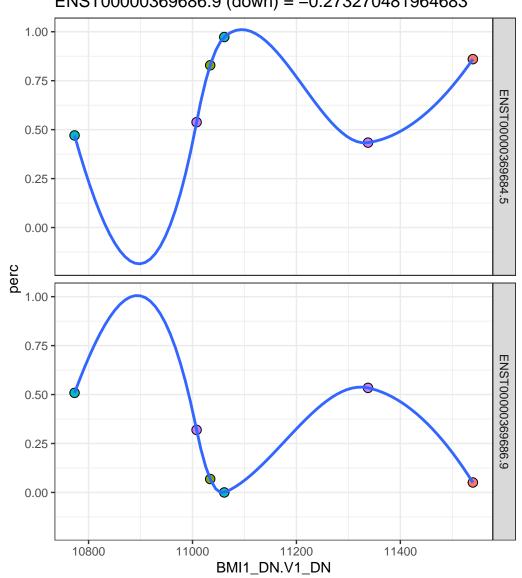
PC3 **VCAP**







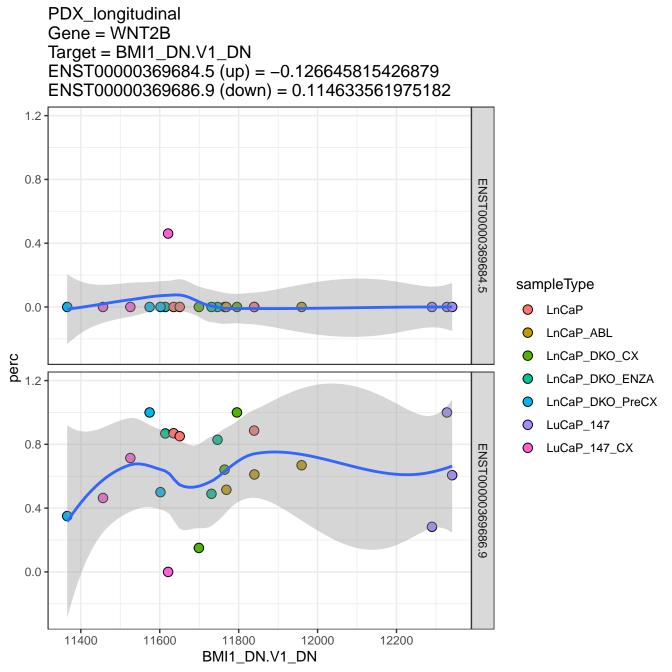
LTL_longitudinal Gene = WNT2B Target = BMI1_DN.V1_DN ENST00000369684.5 (up) = 0.257779824157854 ENST00000369686.9 (down) = -0.273270481964683

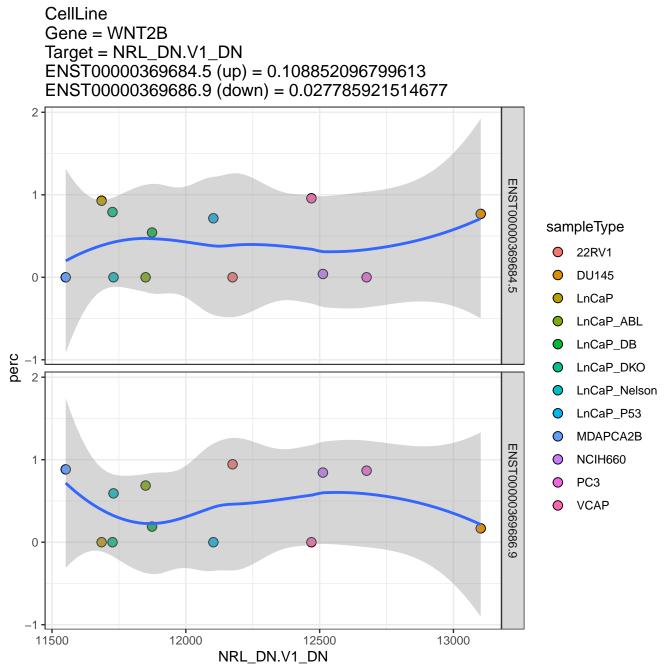


LTL331_Cx_12w

sampleType

- LTL331_Cx_8w
- - LTL331_Cx_R LTL331_PreCx





LTL_longitudinal Gene = WNT2B Target = NRL_DN.V1_DN ENST00000369684.5 (up) = -0.633550002775353ENST00000369686.9 (down) = 0.753968478712970.8 ENST00000369684.5 0.4 sampleType 0.0 LTL331_Cx_12w LTL331_Cx_8w LTL331_Cx_R LTL331_PreCx 8.0 ENST00000369686.9 0.4 0.0

perc

11600

11700

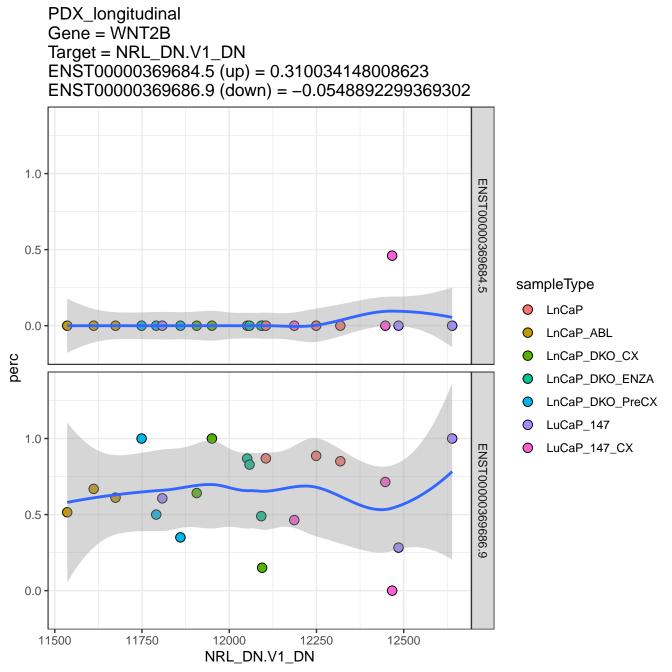
11800

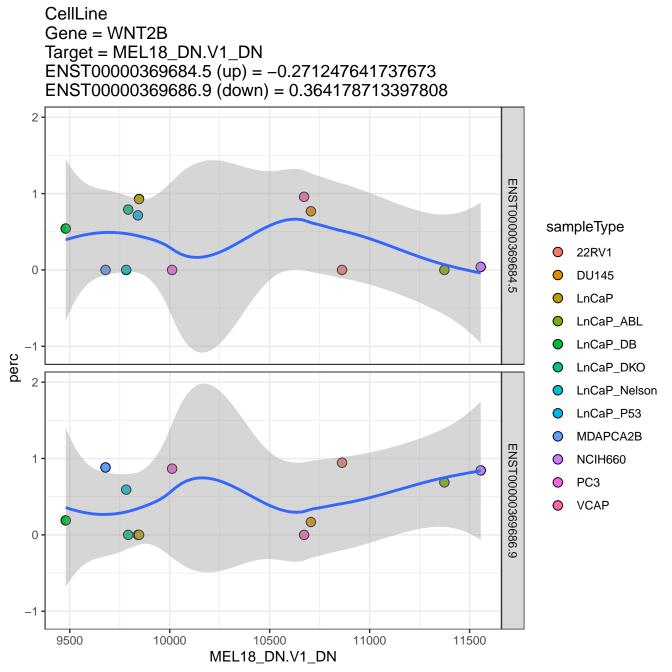
11900

NRL_DN.V1_DN

12000

12100





LTL_longitudinal Gene = WNT2B Target = MEL18_DN.V1_DN ENST00000369684.5 (up) = 0.143879905141259 ENST00000369686.9 (down) = -0.259573499506425 3 -2. ENST00000369684.5 1 0 -1 sampleType -2 **-**LTL331_Cx_12w perc LTL331_Cx_8w 3 -LTL331_Cx_R LTL331_PreCx 2. ENST00000369686.9 1 0 --1 -2 **-**

11100

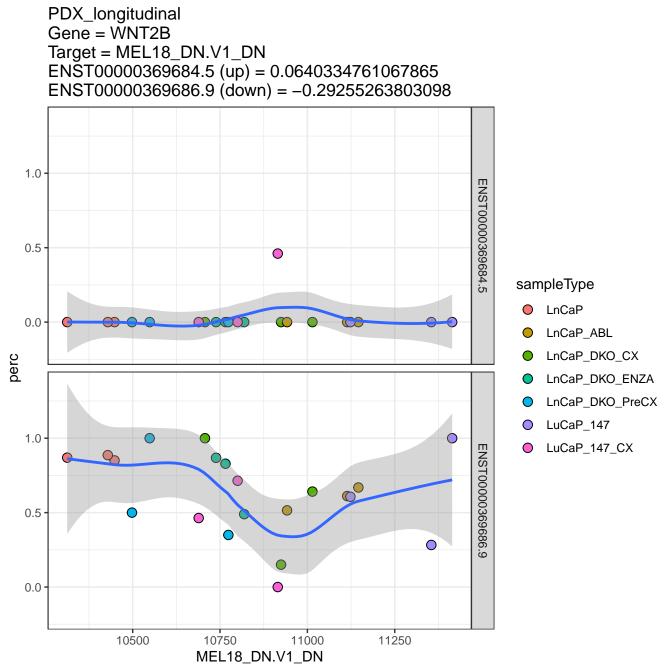
11400

10200

10500

10800

MEL18_DN.V1_DN



CellLine Gene = WNT2B Target = BMI1_DN_MEL18_DN.V1_DN ENST00000369684.5 (up) = -0.209029699111062ENST00000369686.9 (down) = 0.3857298078062361.5 1.0 0 ENST00000369684.5 0.5 sampleType 22RV1 0.0 00 DU145 LnCaP -0.5LnCaP_ABL LnCaP_DB LnCaP_DKO LnCaP_Nelson 1.5 LnCaP_P53 MDAPCA2B 1.0 0 00 ENST00000369686.9 NCIH660 0 PC3 0.5 -**VCAP** 0 0 0.0

11000

11500

perc

-0.5

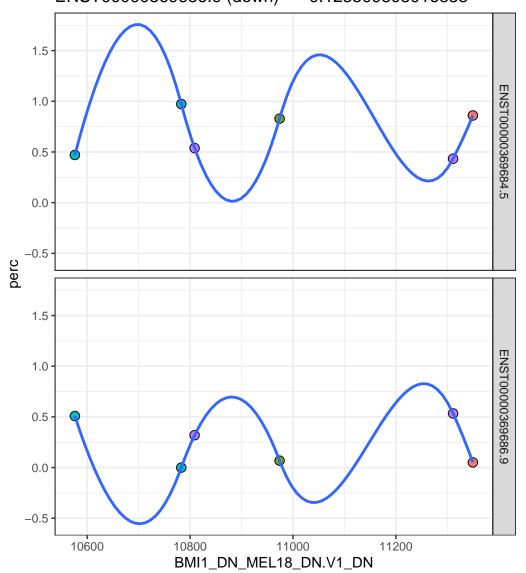
9500

10000

10500

BMI1_DN_MEL18_DN.V1_DN

LTL_longitudinal
Gene = WNT2B
Target = BMI1_DN_MEL18_DN.V1_DN
ENST00000369684.5 (up) = 0.100332940848864
ENST00000369686.9 (down) = -0.123509808016838



sampleType

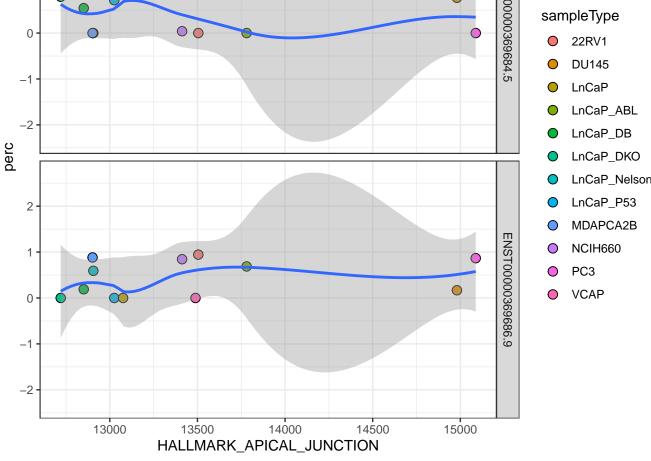
LTL331_Cx_12w

LTL331_Cx_8w LTL331_Cx_R

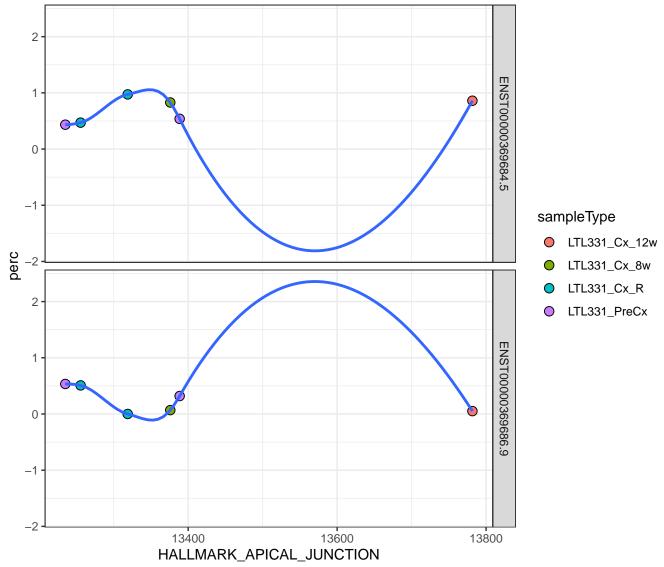
LTL331_PreCx

PDX_longitudinal Gene = WNT2B Target = BMI1_DN_MEL18_DN.V1_DN ENST00000369684.5 (up) = 0.235546152647538ENST00000369686.9 (down) = -0.2803511379776361.0 ENST00000369684.5 0.5 sampleType LnCaP 0.0 LnCaP_ABL LnCaP_DKO_CX perc LnCaP_DKO_ENZA LnCaP_DKO_PreCX LuCaP_147 1.0 LuCaP_147_CX 60 ENST00000369686.9 8 0.5 0.0 10400 10800 11200 11600 BMI1_DN_MEL18_DN.V1_DN

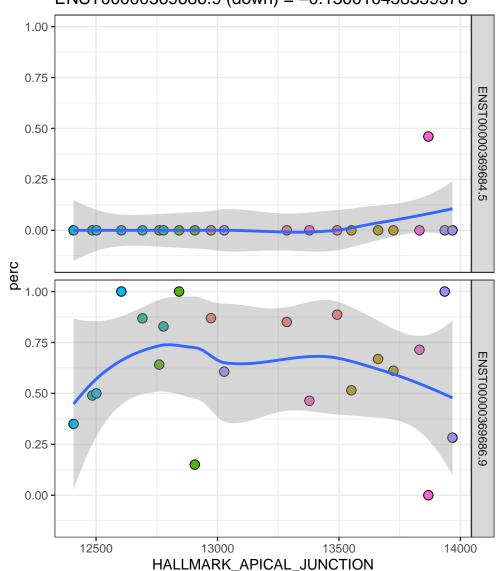
CellLine Gene = WNT2B Target = HALLMARK_APICAL_JUNCTION ENST00000369684.5 (up) = -0.129475867845981ENST00000369686.9 (down) = 0.2441187759589292 ENST00000369684.5 1 sampleType 0 22RV1 DU145 -1 LnCaP LnCaP_ABL -2 -LnCaP_DB LnCaP_DKO LnCaP_Nelson LnCaP_P53 2



LTL_longitudinal
Gene = WNT2B
Target = HALLMARK_APICAL_JUNCTION
ENST00000369684.5 (up) = 0.49459577105368
ENST00000369686.9 (down) = -0.568209550154603



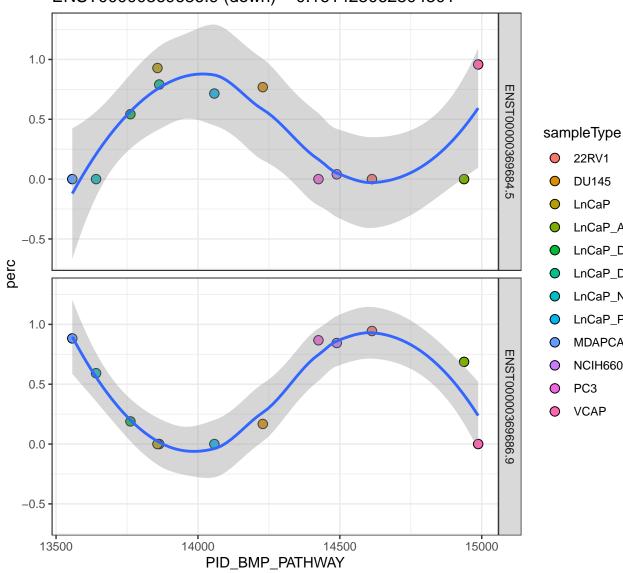
PDX_longitudinal
Gene = WNT2B
Target = HALLMARK_APICAL_JUNCTION
ENST00000369684.5 (up) = 0.30006481310348
ENST00000369686.9 (down) = -0.130010458359378



sampleType

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

CellLine Gene = WNT2B Target = PID_BMP_PATHWAY ENST00000369684.5 (up) = -0.0635604714798751ENST00000369686.9 (down) = 0.161425062804301



22RV1

DU145 LnCaP

LnCaP_ABL

LnCaP_DB

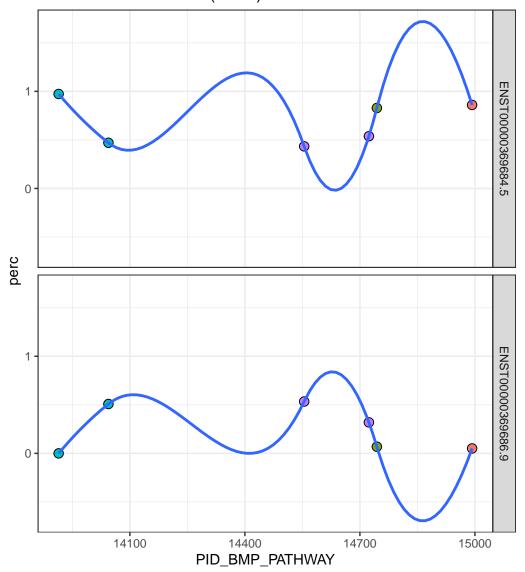
LnCaP_DKO LnCaP_Nelson LnCaP_P53

MDAPCA2B

NCIH660

PC3 **VCAP**

LTL_longitudinal
Gene = WNT2B
Target = PID_BMP_PATHWAY
ENST00000369684.5 (up) = 0.00815008844508385
ENST00000369686.9 (down) = -0.162389351329979

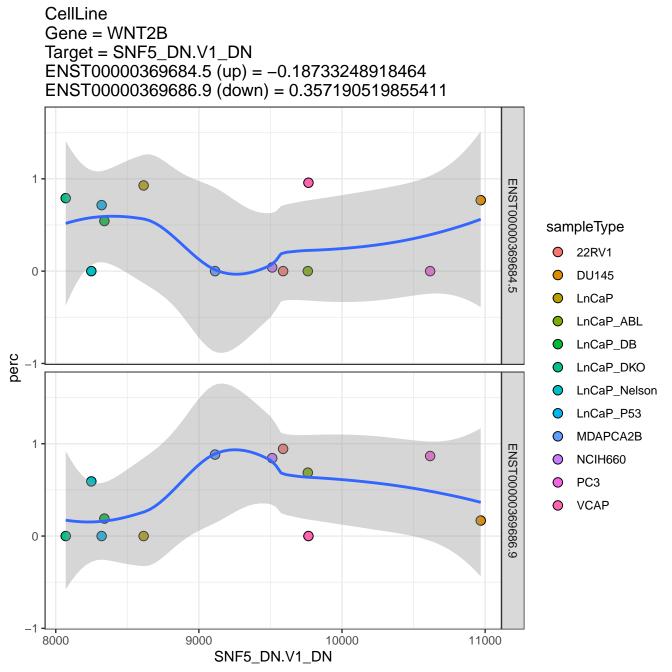


sampleType

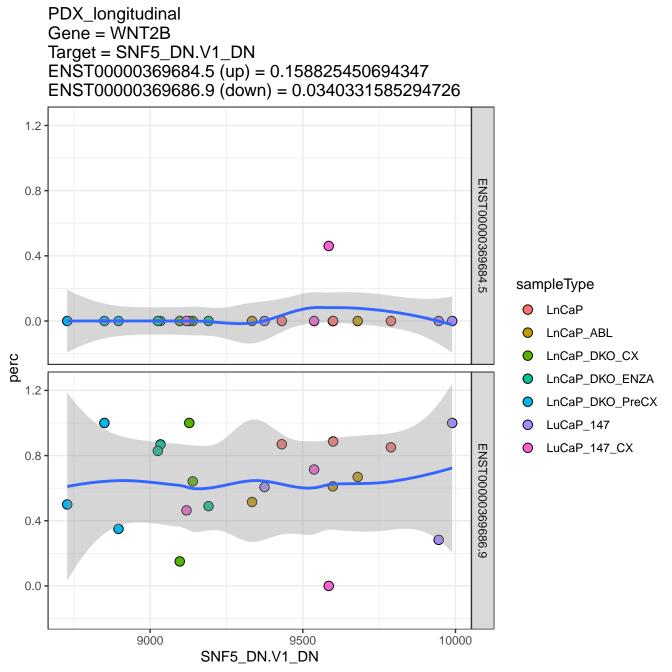
LTL331_Cx_12w

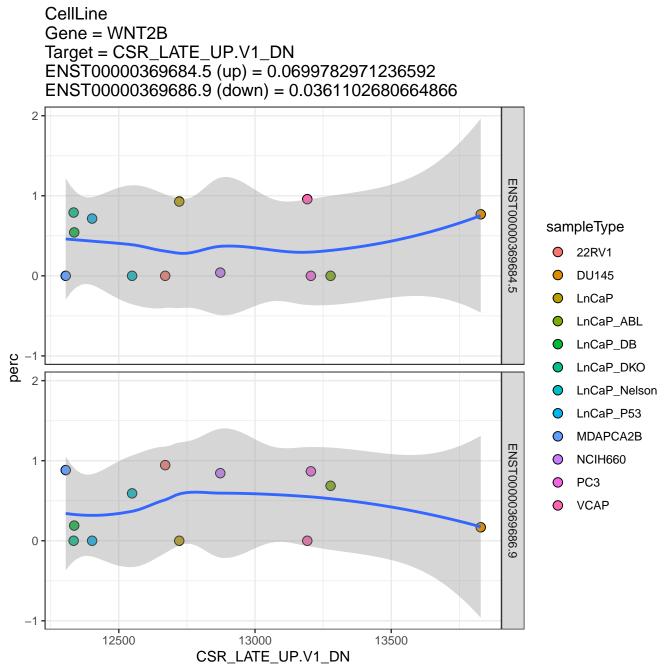
LTL331_Cx_8w LTL331_Cx_R LTL331_PreCx

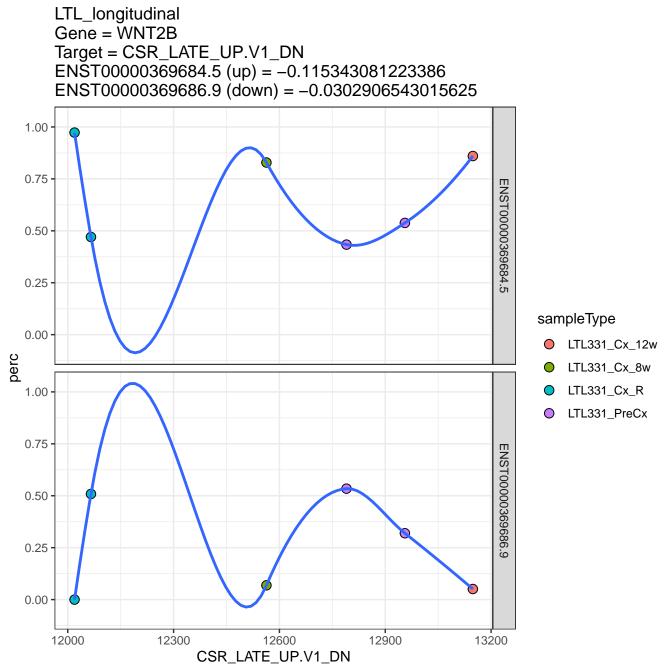
PDX_longitudinal Gene = WNT2B Target = PID_BMP_PATHWAY ENST00000369684.5 (up) = -0.0813082359634827ENST00000369686.9 (down) = -0.0483673786094721.0 ENST00000369684.5 0.5 0 sampleType LnCaP 0.0 LnCaP_ABL LnCaP_DKO_CX perc LnCaP_DKO_ENZA LnCaP_DKO_PreCX LuCaP_147 0 1.0 LuCaP_147_CX ENST00000369686.9 0.5 0 0.0 13000 13500 14000 14500 15000 PID_BMP_PATHWAY

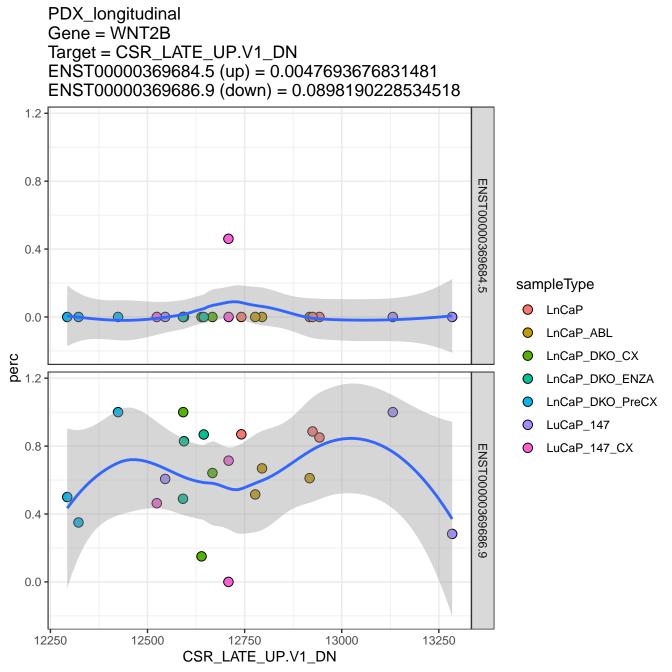


LTL_longitudinal Gene = WNT2B Target = SNF5_DN.V1_DN ENST00000369684.5 (up) = 0.0763746422953201 ENST00000369686.9 (down) = -0.191782728630605 ENST00000369684.5 0 sampleType LTL331_Cx_12w LTL331_Cx_8w LTL331_Cx_R LTL331_PreCx ENST00000369686.9 0 8000 8500 9000 9500 SNF5_DN.V1_DN





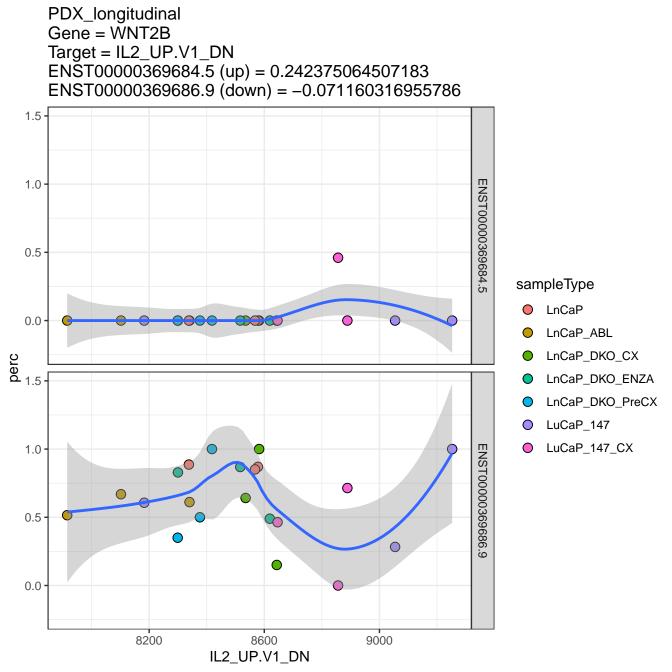


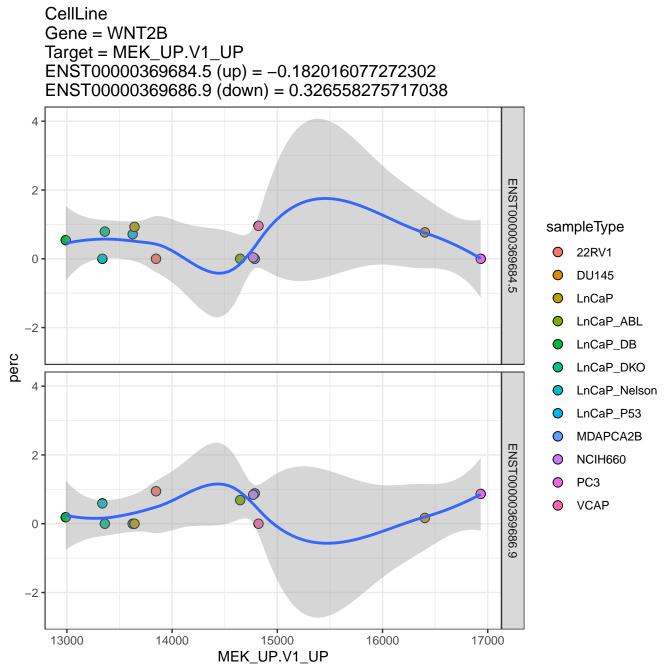


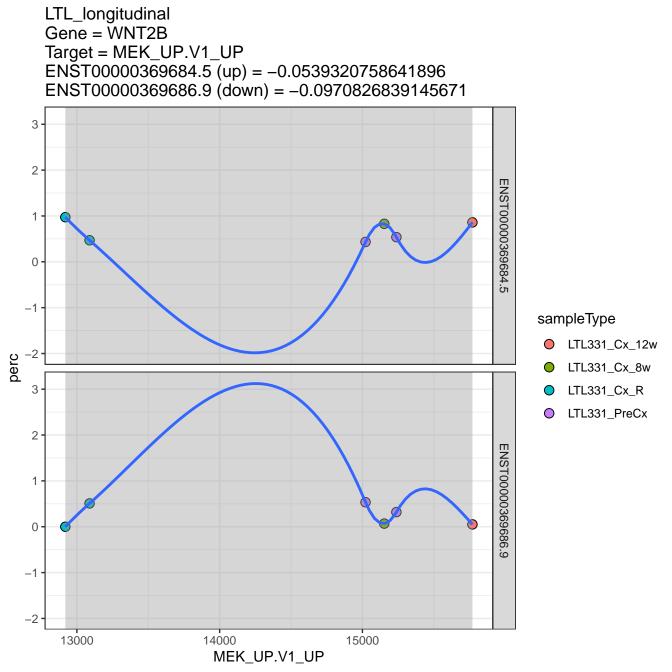
Gene = WNT2B Target = IL2_UP.V1_DN ENST00000369684.5 (up) = -0.198168072033492ENST00000369686.9 (down) = 0.309278790016973ENST00000369684.5 1 sampleType 22RV1 DU145 00 0 LnCaP LnCaP_ABL LnCaP_DB perc LnCaP_DKO LnCaP_Nelson LnCaP_P53 MDAPCA2B ENST00000369686.9 NCIH660 PC3 **VCAP** 0 8100 8400 8700 IL2_UP.V1_DN

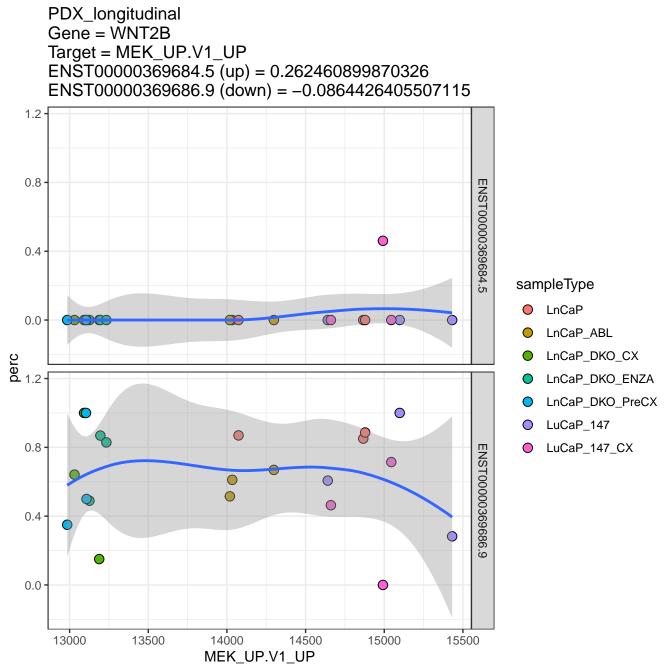
CellLine

LTL_longitudinal Gene = WNT2B Target = IL2_UP.V1_DN ENST00000369684.5 (up) = 0.417485689009327 ENST00000369686.9 (down) = -0.496580706358076 3 ENST00000369684.5 0 -3 sampleType LTL331_Cx_12w perc -6 LTL331_Cx_8w 6 LTL331_Cx_R LTL331_PreCx 3 -ENST00000369686.9 0 -3 8750 9000 9250 9500 9750 IL2_UP.V1_DN



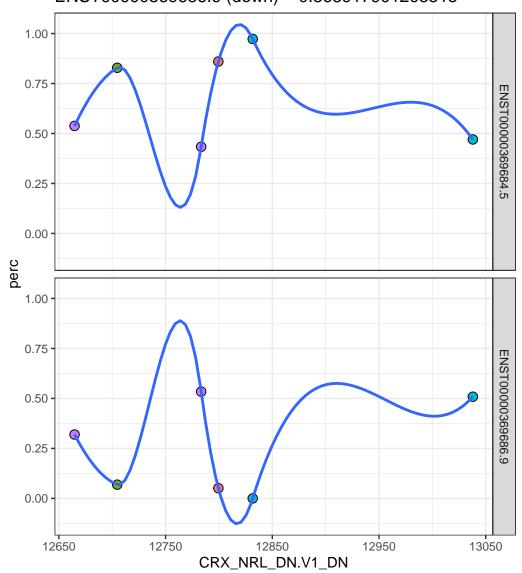






CellLine Gene = WNT2B Target = CRX_NRL_DN.V1_DN ENST00000369684.5 (up) = -0.239681299426219ENST00000369686.9 (down) = 0.347162174404545ENST00000369684.5 1 0 sampleType 22RV1 0 DU145 0 LnCaP LnCaP_ABL LnCaP_DB perc LnCaP_DKO LnCaP_Nelson LnCaP_P53 MDAPCA2B ENST00000369686.9 1 8 NCIH660 0 PC3 **VCAP** 0 0 12000 12500 13000 CRX_NRL_DN.V1_DN

LTL_longitudinal
Gene = WNT2B
Target = CRX_NRL_DN.V1_DN
ENST00000369684.5 (up) = -0.21078788060073
ENST00000369686.9 (down) = 0.363917901203515

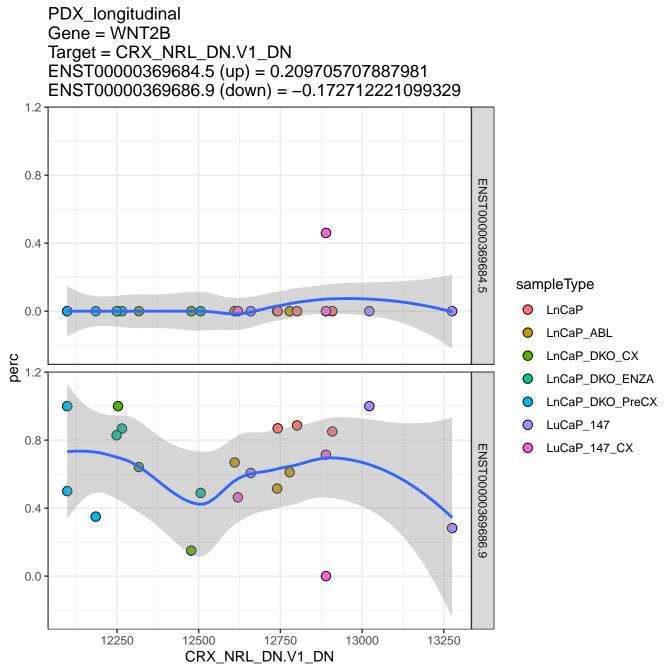


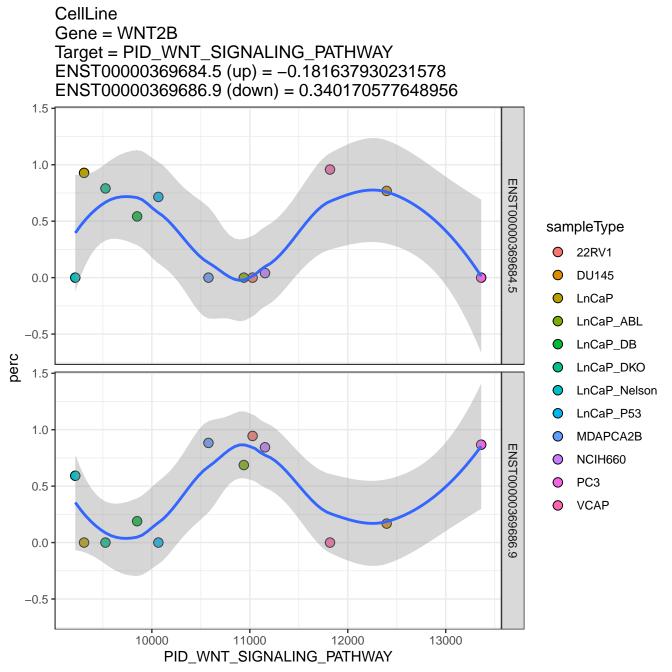
sampleType

LTL331_Cx_12w

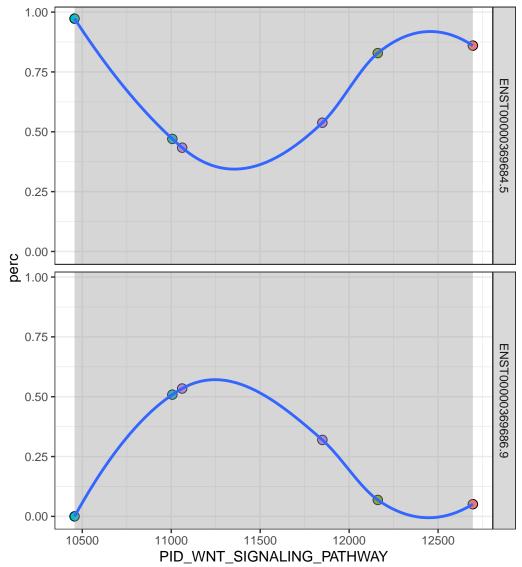
LTL331_Cx_8w LTL331_Cx_R

LTL331_PreCx





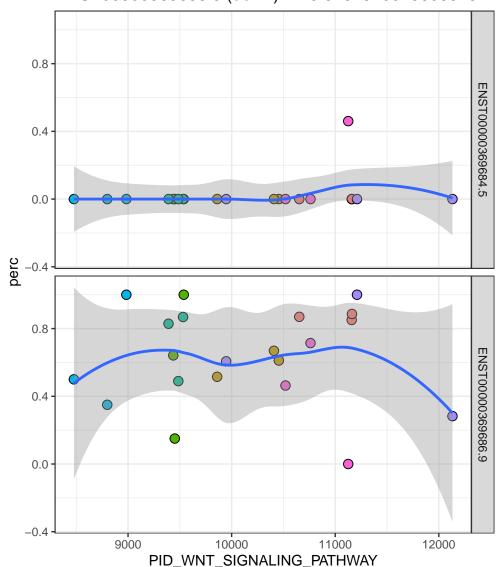
LTL_longitudinal
Gene = WNT2B
Target = PID_WNT_SIGNALING_PATHWAY
ENST00000369684.5 (up) = 0.175389497481883
ENST00000369686.9 (down) = -0.323207908121497



sampleType

LTL331_Cx_12w

LTL331_Cx_8w LTL331_Cx_R LTL331_PreCx PDX_longitudinal Gene = WNT2B Target = PID_WNT_SIGNALING_PATHWAY ENST00000369684.5 (up) = 0.24438725818419 ENST00000369686.9 (down) = -0.0401978079695515



sampleType

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

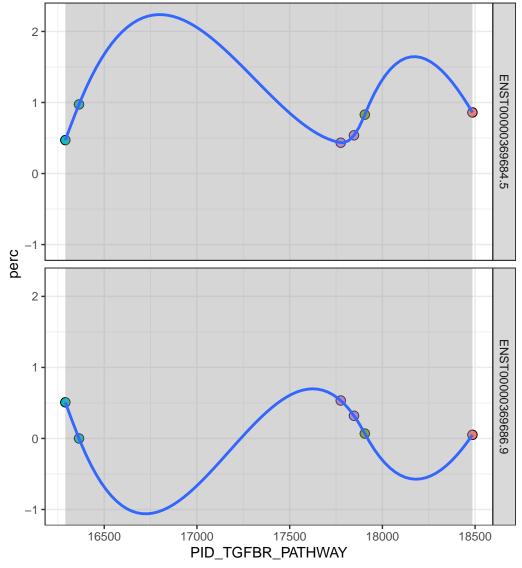
CellLine Gene = WNT2B Target = PID_TGFBR_PATHWAY ENST00000369684.5 (up) = -0.284892810066567ENST00000369686.9 (down) = 0.420329574688121.5 -1.0 0 ENST00000369684.5 0.5 sampleType 22RV1 0.0 **DU145** LnCaP LnCaP_ABL -0.5 -LnCaP_DB perc LnCaP_DKO 1.5 LnCaP_Nelson LnCaP_P53 1.0 MDAPCA2B 0 ENST00000369686.9 NCIH660 PC3 0.5 -**VCAP** 0.0 -0.5 -

18000

17000

PID_TGFBR_PATHWAY

LTL_longitudinal
Gene = WNT2B
Target = PID_TGFBR_PATHWAY
ENST00000369684.5 (up) = 0.0414188027274956
ENST00000369686.9 (down) = -0.182500046818596

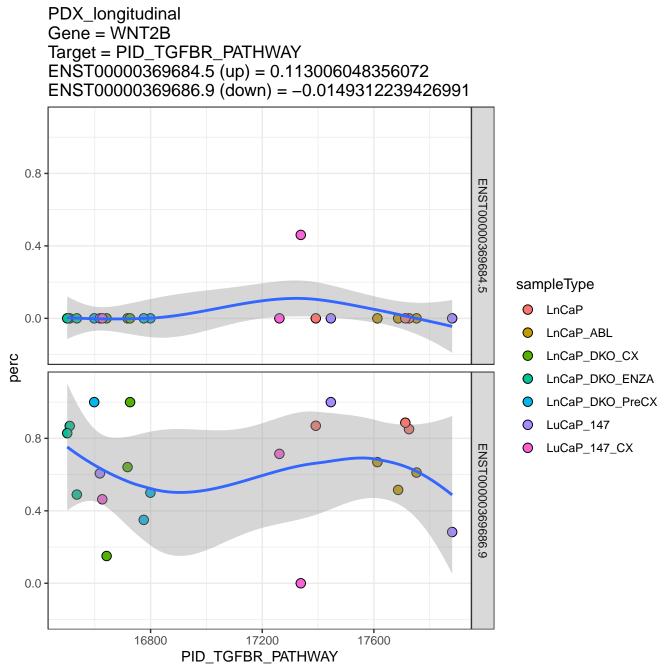


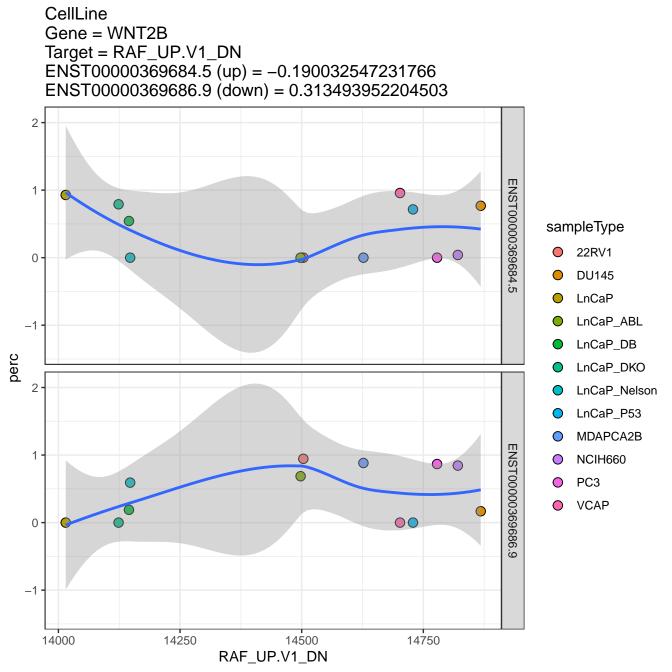
sampleType

LTL331_Cx_12w

LTL331_Cx_8w LTL331_Cx_R

LTL331_PreCx





LTL_longitudinal Gene = WNT2B Target = RAF_UP.V1_DN ENST00000369684.5 (up) = 0.0515708677444417ENST00000369686.9 (down) = -0.1844076891078721.5 1.0 ENST00000369684.5 0.5 0.0 sampleType -0.5 -LTL331_Cx_12w perc LTL331_Cx_8w 1.5 LTL331_Cx_R LTL331_PreCx 1.0 ENST00000369686.9 0.5

14500

RAF_UP.V1_DN

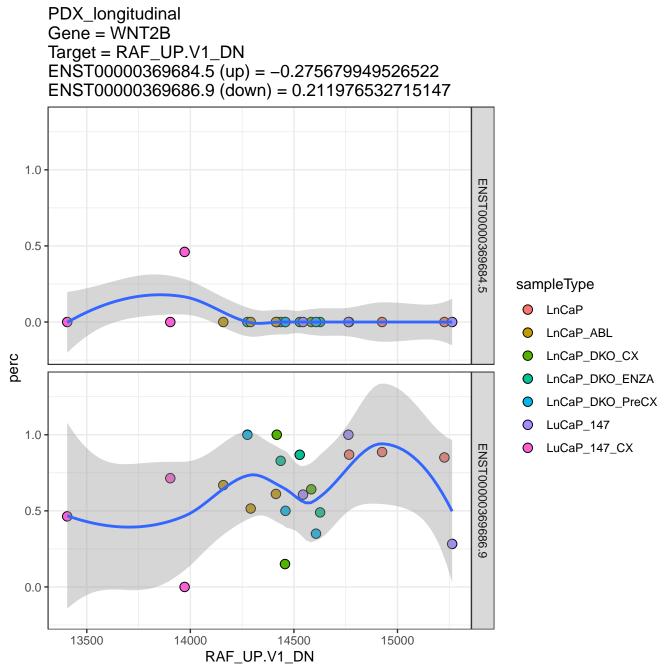
15000

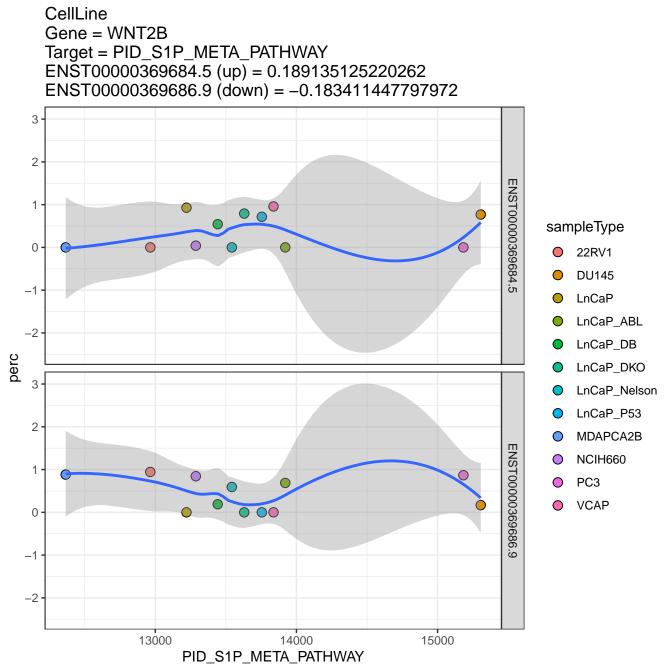
0.0

-0.5

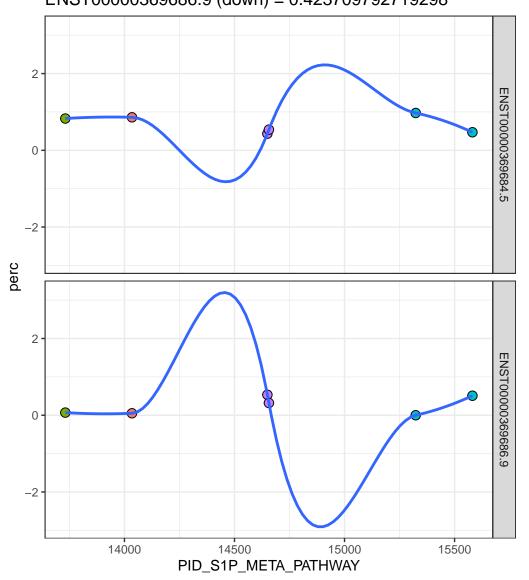
13500

14000





LTL_longitudinal
Gene = WNT2B
Target = PID_S1P_META_PATHWAY
ENST00000369684.5 (up) = -0.300495650777626
ENST00000369686.9 (down) = 0.423709792719298



sampleType

LTL331_Cx_12w

LTL331_Cx_8w LTL331_Cx_R LTL331_PreCx

