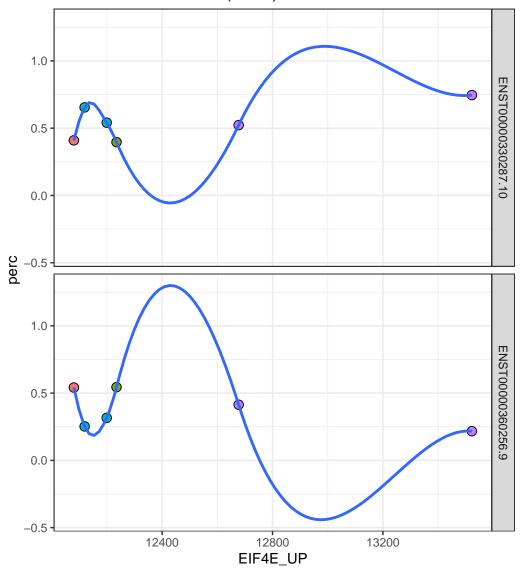
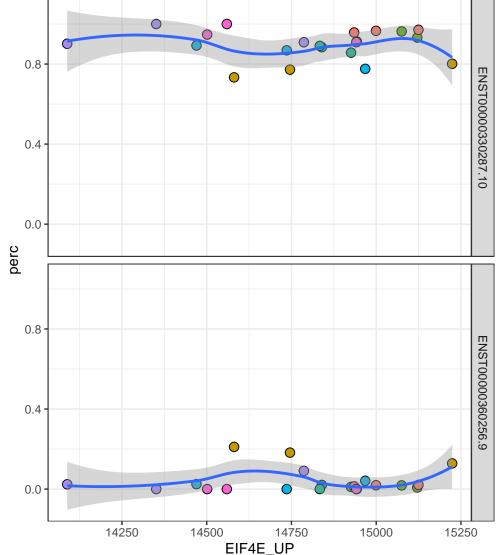


LTL_longitudinal
Gene = F8
Target = EIF4E_UP
ENST00000330287.10 (up) = 0.675705706650968
ENST00000360256.9 (down) = -0.514540677039531



sampleType

LTL331_Cx_12w LTL331_Cx_8w LTL331_Cx_R LTL331_PreCx PDX_longitudinal Gene = F8Target = EIF4E_UP ENST00000330287.10 (up) = -0.0713020047304767ENST00000360256.9 (down) = 0.008432884221408990 sampleType

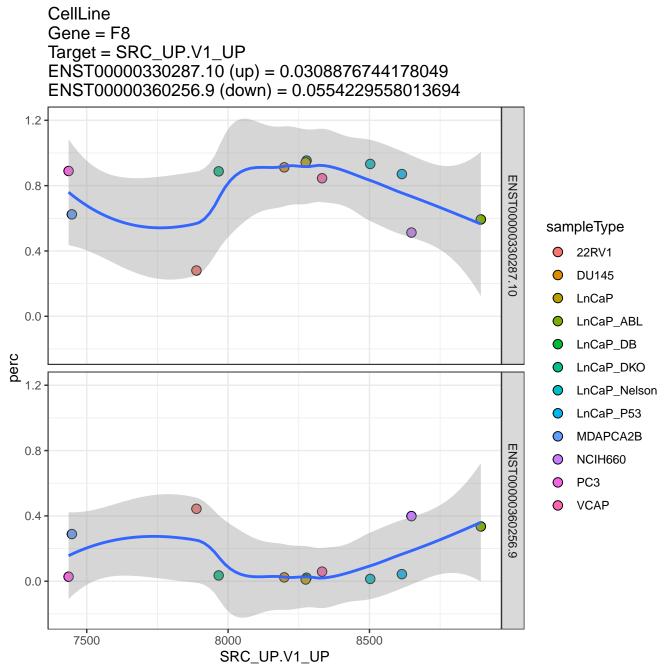


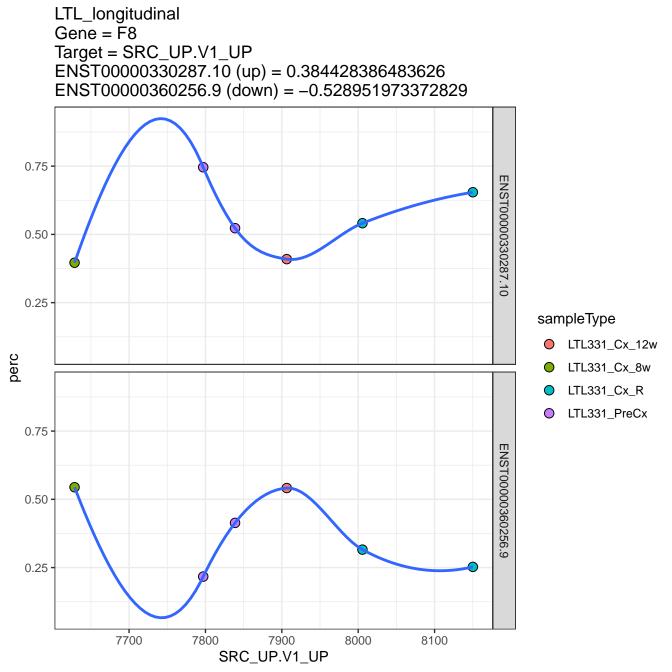
LnCaP

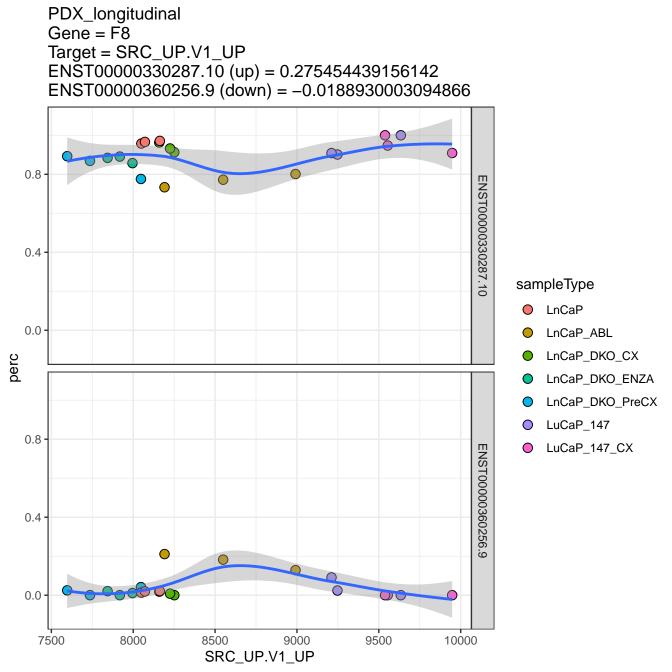
- LnCaP_ABL

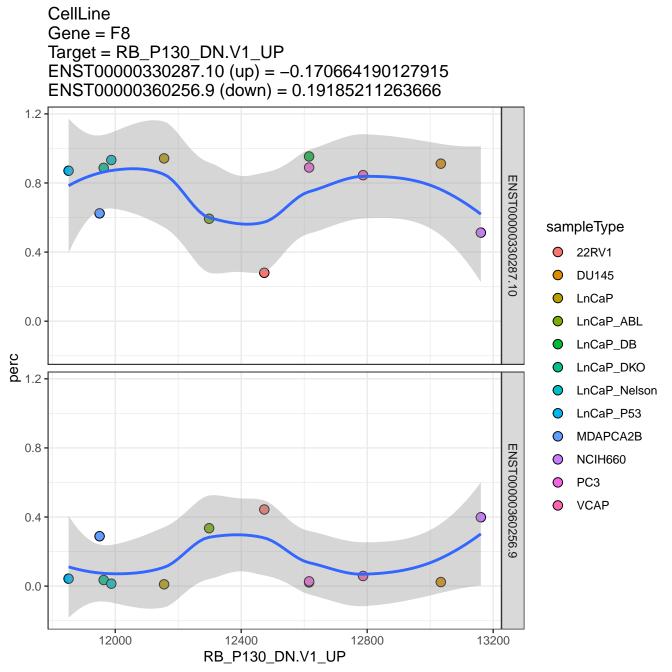
LnCaP_DKO_CX

- LnCaP_DKO_ENZA
- LnCaP_DKO_PreCX
- LuCaP_147
- LuCaP_147_CX

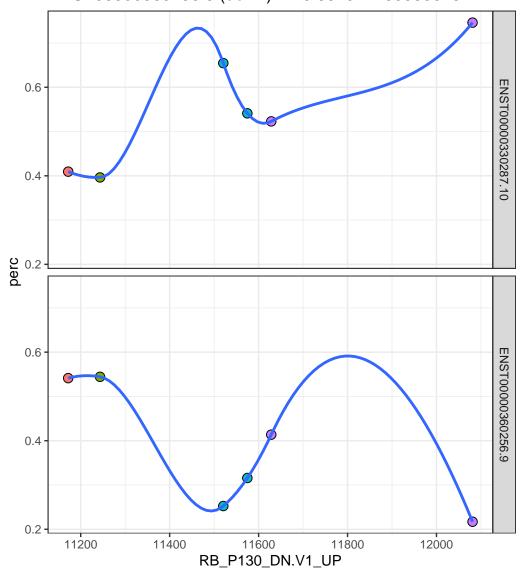






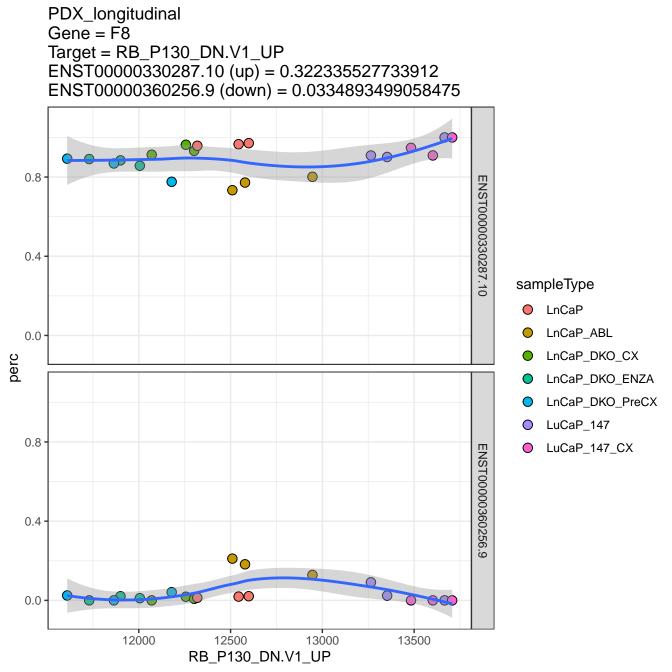


LTL_longitudinal
Gene = F8
Target = RB_P130_DN.V1_UP
ENST00000330287.10 (up) = 0.896215630646999
ENST00000360256.9 (down) = -0.837574400939878



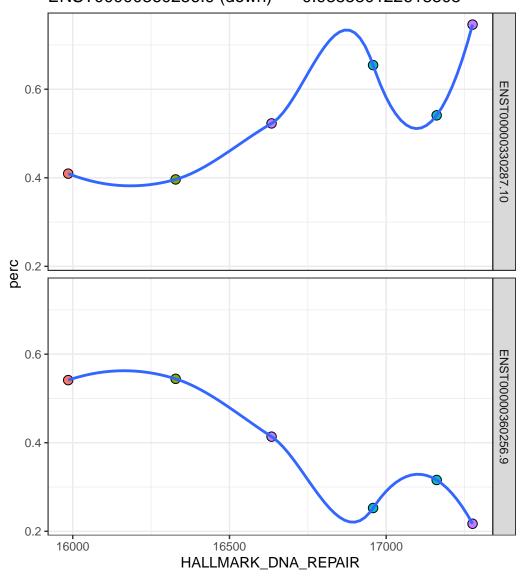
sampleType

LTL331_Cx_12w LTL331_Cx_8w LTL331_Cx_R LTL331_PreCx



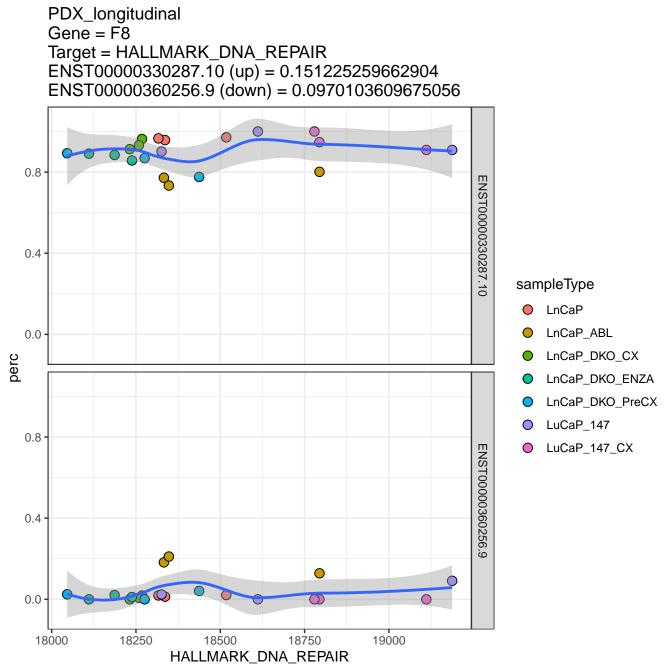
CellLine Gene = F8Target = HALLMARK_DNA_REPAIR ENST00000330287.10 (up) = 0.308952111138586ENST00000360256.9 (down) = -0.3362865652305391.5 -1.0 -ENST00000330287.10 sampleType 0.5 22RV1 **DU145** LnCaP 0.0 LnCaP_ABL LnCaP_DB perc -0.5LnCaP_DKO 1.5 -LnCaP_Nelson LnCaP_P53 MDAPCA2B 1.0 ENST00000360256.9 NCIH660 PC3 0.5 **VCAP** 0.0 -0.517800 18000 18200 18400 HALLMARK_DNA_REPAIR

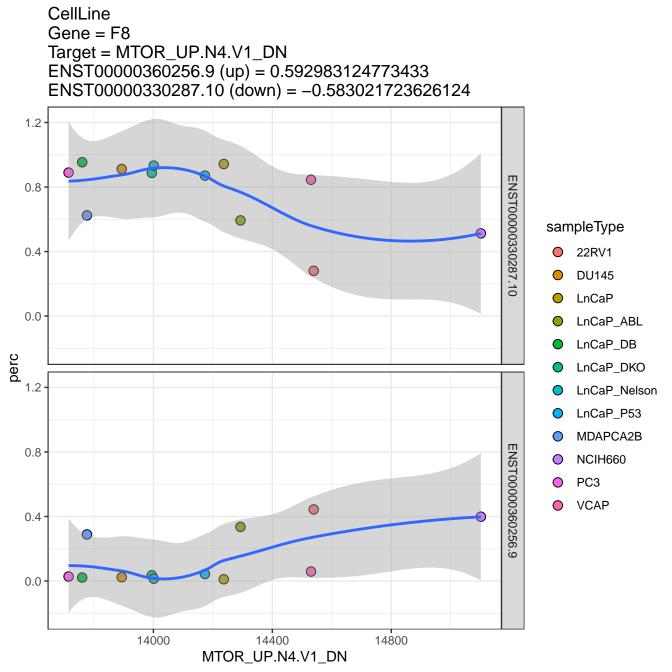
LTL_longitudinal
Gene = F8
Target = HALLMARK_DNA_REPAIR
ENST00000330287.10 (up) = 0.862123168403344
ENST00000360256.9 (down) = -0.938580122618505



sampleType

LTL331_Cx_12w LTL331_Cx_8w LTL331_Cx_R LTL331_PreCx





LTL_longitudinal Gene = F8Target = MTOR_UP.N4.V1_DN ENST00000360256.9 (up) = 0.803063548333023ENST00000330287.10 (down) = -0.8406454822421280.75 ENST00000330287.10 0.50 0.25 sampleType LTL331_Cx_12w perc LTL331_Cx_8w LTL331_Cx_R LTL331_PreCx 0.75 ENST00000360256.9 0.50

15100

15300

0.25

14700

14900

MTOR_UP.N4.V1_DN

