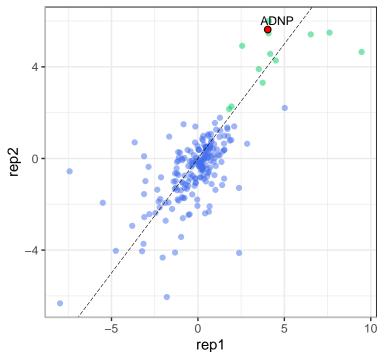
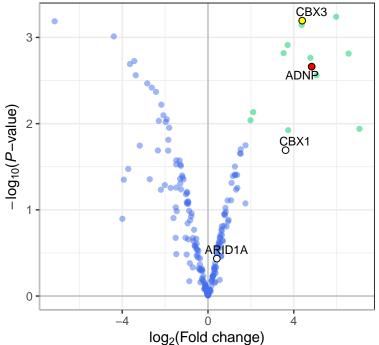
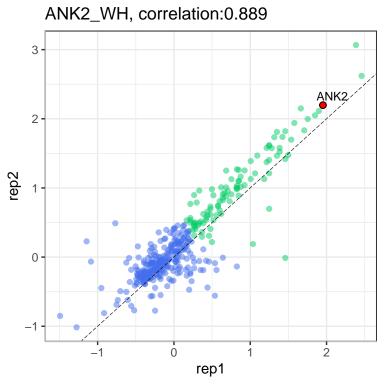
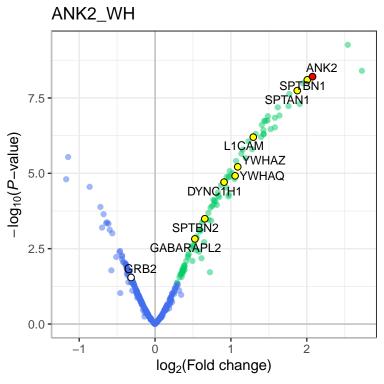
ADNP_CNCR1, correlation:0.736



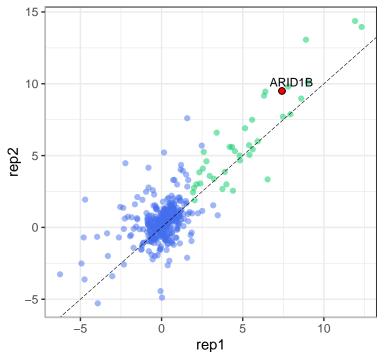
ADNP_CNCR1 3 ·



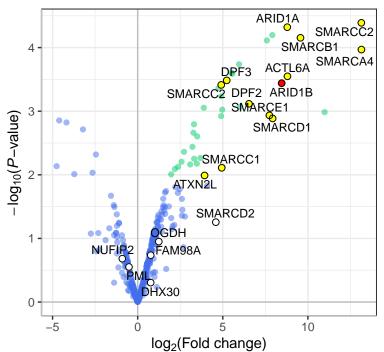




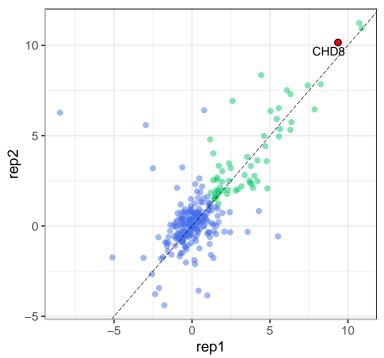
ARID1B_CNCR1, correlation:0.804



ARID1B_CNCR1

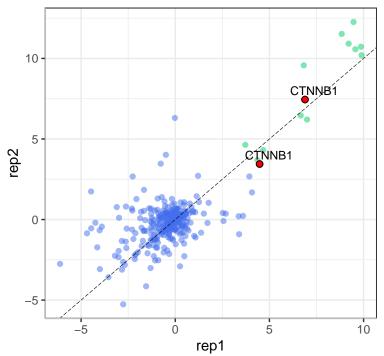


CHD8_CNCR2, correlation:0.721

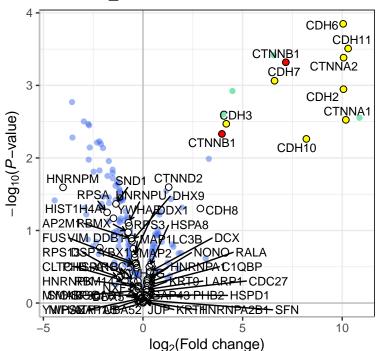


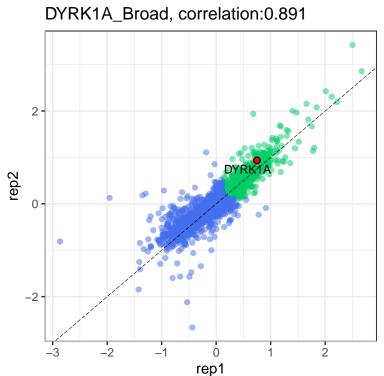
CHD8_CNCR2 6 -OGDH CHD8 OGDH -log₁₀(P-value) SPTBN1 CTNNB1 -4 log₂(Fold change)

CTNNB1_CNCR1, correlation:0.789

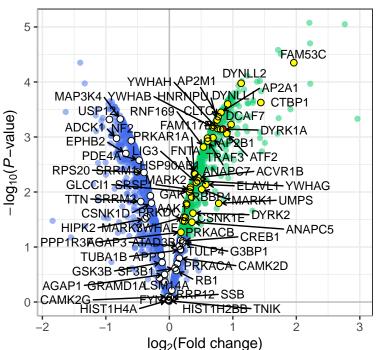


CTNNB1_CNCR1

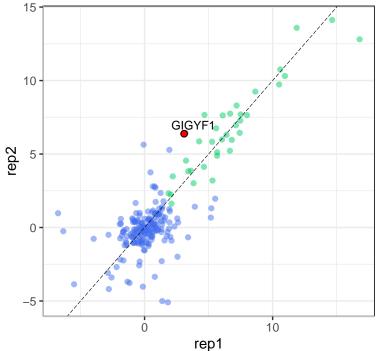




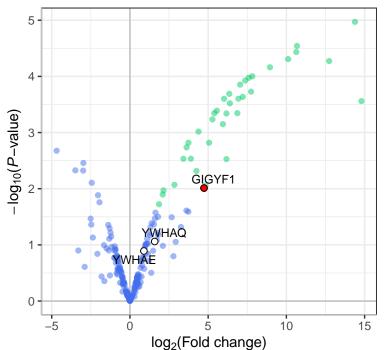
DYRK1A_Broad



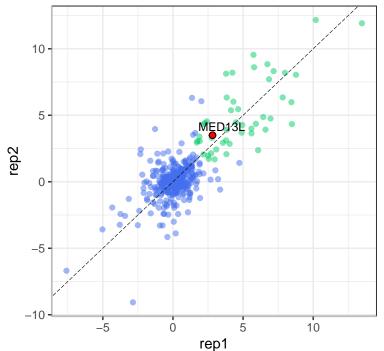




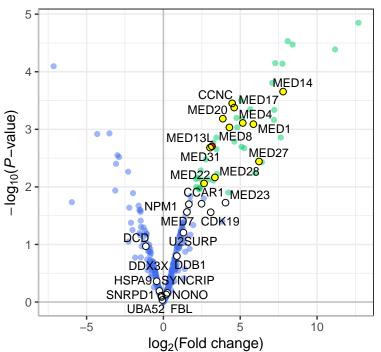
GIGYF1_CNCR2



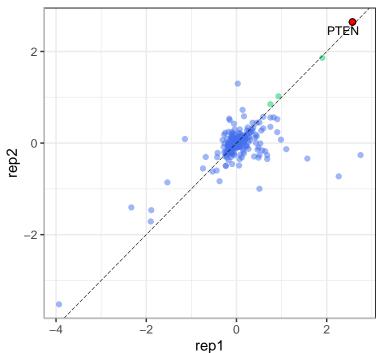
MED13L_CNCR1, correlation:0.791



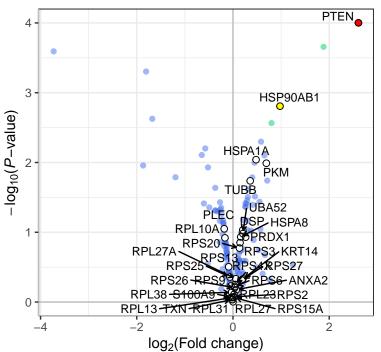
MED13L_CNCR1

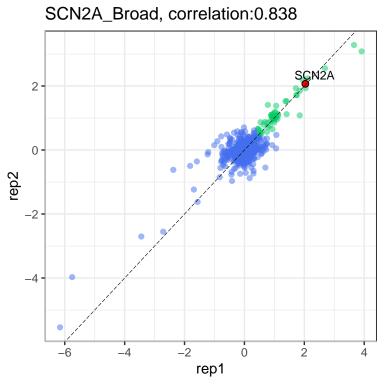


PTEN_WH1, correlation:0.653

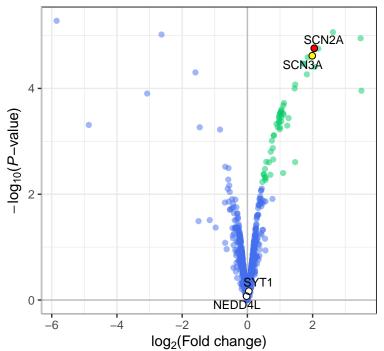


PTEN_WH1

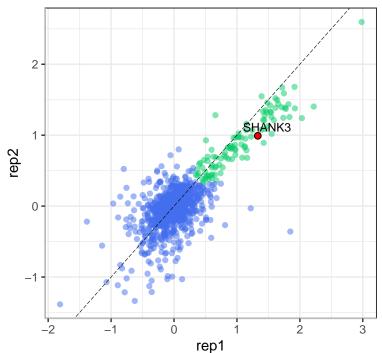




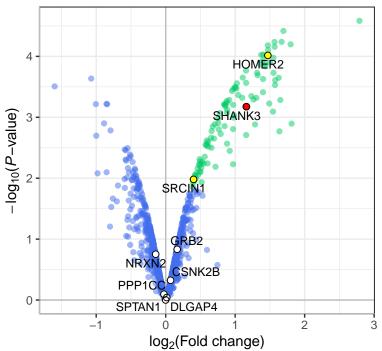
SCN2A_Broad



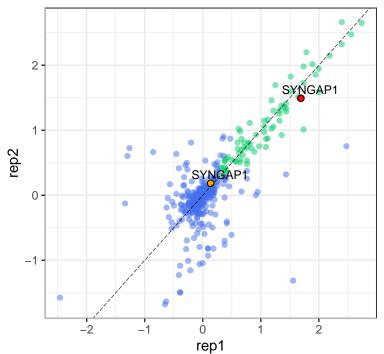
SHANK3_Broad, correlation:0.785



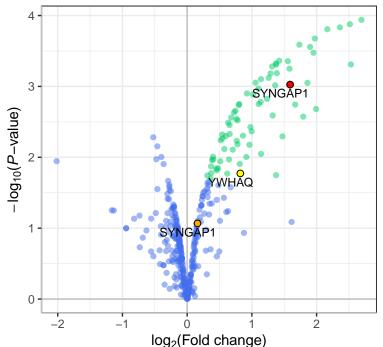
SHANK3_Broad



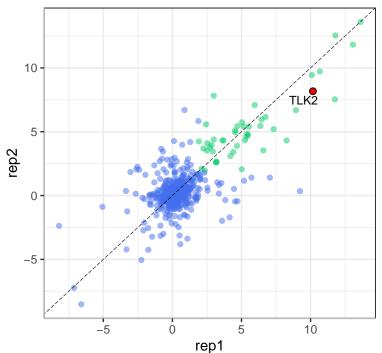
SYNGAP1_WH1, correlation:0.785



SYNGAP1_WH1



TLK2_CNCR1, correlation:0.733



TLK2_CNCR1 TLK2 -log₁₀(P-value) YWHAB HAZ 0 10 -5 log₂(Fold change)