Full_Dataset_protein_coding_interaction_Driver_EGFRvV_vs_PDGFRA

Full_Dataset_protein_coding_interaction

Legend: NS ● Log$_2$ FC ● p − value and log$_2$ FC

y-axis: $-Log_{10} P$ (values: 0, 30, 60, 90, 120)

x-axis: Log$_2$ fold change (values: −15, −10, −5, 0, 5, 10)

total = 13689 variables

Labeled points (right side, upregulated): Egfr, Yes1, Bag3, Cavin1, Smo, Gcnt4, Tmem98, Mxra8, Aprt, Ntn4, Dusp6, Smad3, Angpt2, Sdc4, Erbb2, Serpinh1, Ifitm2, Clic5, Wwtr1, Icosl, Areg, Tm9sf5, Tubb6, Synj2, Bdkrb2, Spry2, Egr2, Liph, Phlda1, Pygm, Smim43, Tslp, Acsf2, Trim16, Acy3, Chl1, Dgkk, Notum, Galns, Sox1, Otx2, Gria1, Retsat, Ugt1a6a, Mtap, Oacyl, Thbs1, Mmp16, Bmp1, Wnt11, Flt3, Rpl36a

Labeled points (left side, downregulated): Pdgfra, Myt1l, Fbxl7, St6gal2, Vstm2a, Flrt3, Aff3, Grin3a, Adal, Dach1, Sult4a1, Slit2, Tmem119, Sh3bp4, Cadps, Nyap2, Ntpcr, Gsap, Lcmt2, Snca, Prmt8, Ptprk, Flrt1, Lrch3, Alx4, Ptprd, Id4, Byes, Evc, Dop6, Rhof, Rab29, Rtp4, Gnpda1, Gm43302, Rhod, Vps37b