

Gene	PR		VP		Description
	Fold change	Std dev	Fold change	Std dev	
VAV2	-1.0	0.02	2.3	0.29	Rho-associated, coiled-coil containing protein 2 (synaptobrevin 2)
ROCK2	1.6	0.02	2.8	0.29	Rho-associated, coiled-coil containing protein kinase 1
SH3GLB2	-1.2	0.15	2.8	0.58	SH3-domain GRB2-like endophilin B2
EPS15L	0.1	1.62	2.7	0.48	Epidermal growth factor receptor pathway substrate 15
RAB3C	1.5	0.11	2.6	0.59	RAB3C, member RAS oncogene family
AP2M1	1.8	0.05	2.6	0.56	Adaptor-related protein complex 2, alpha 1 subunit
DAPH1	-1.5	0.01	2.6	0.45	Diaphanous-related protein 1
RAB29	-1.1	0.01	2.6	0.59	RAB29, member RAS oncogene family
MAPK8IP1	-1.2	0.15	2.6	0.42	Mitogen-activated protein kinase 8 interacting protein 1
ROCK2	-1.8	0.27	2.5	0.45	Rho-associated, coiled-coil containing protein kinase 2
AMPH	1.1	0.05	2.5	0.50	Amphiphysin
RAB3D	1.3	0.03	2.5	0.39	RAB3D, member RAS oncogene family
GRB2	-1.1	0.03	2.5	0.63	Growth factor receptor-bound protein 2
DNMS	1.1	0.05	2.5	0.38	ADP-ribosylation factor 6
ARF6	-1.4	0.1	2.5	0.32	WAS protein family, member 1
WASF1	-0.3	1.91	2.5	0.35	RAS oncogene family member A
RHOA	1.3	0.08	2.5	0.32	RHOA, member RHO
SYT1	-1.1	0.12	2.5	0.32	Actin related protein 3 homolog (yeast)
ACTR3	-1.1	0.07	2.4	0.53	ARP1 actin-related protein 3 homolog (yeast)
CYTH3	1.2	0.11	2.4	0.46	Cythromodulin 3
TSG101	1.2	0.10	2.4	0.47	Tumor susceptibility 101
ATG16L	0.0	1.49	2.4	0.38	Autophagy related 16-like
ARF10	-2.1	0.01	1.60	0.24	ADP-ribosylation factor 10
RAB17B	1.3	0.1	2.4	0.48	RAB17B, member RAS oncogene family
RAB11B	-3.2	0.24	2.3	0.41	RAB11B, member RAS oncogene family
ARPC1B	-1.4	0.12	2.3	0.31	Actin related protein 2/3 complex, subunit 1B, 41kDa
CB3	-1.2	0.09	2.3	0.43	Calmodulin and integrin binding family member 3
PI4KA	0.0	1.64	2.3	0.37	PI4KA, catalytic subunit
PI4KC2	-1.3	0.03	2.3	0.37	Phosphatidylinositol-4-kinase type 2 alpha
EEA1	-1.1	0.06	2.3	0.40	Early endosome antigen 1
VAMP1	-1.7	0.29	2.2	0.48	Vesicle-associated membrane protein 1 (synaptobrevin 1)
VAPA	-1.1	0.12	2.2	0.10	VAMP (vesicle-associated membrane protein)-associated protein A, 33kDa
SNX2	0.1	1.64	2.2	0.07	Sorting nexin 2
RAB3B	-1.2	0.08	2.2	0.30	RAB3B, member RAS oncogene family
ATP9B041	-1.8	0.20	2.2	0.50	ATPase, H+ transporting, lysosomal V0 subunit a1
BECN1	-1.2	0.11	2.1	0.41	Bet61, autophagy related
CLCT1L	1.5	0.07	2.1	0.22	Clathrin, heavy chain-like 1
PI4KA	-1.3	0.01	2.1	0.27	Phosphatidylinositol-4-kinase, catalytic, alpha
PI4KB	-2.4	0.36	2.1	0.56	Phosphatidylinositol-4-kinase, catalytic, beta
NEED1	1.1	0.1	2.1	0.32	Neuroendocrine cell expressed, developmentally down-regulated 4, E3 ubiquitin protein ligase
CDC42	-2.1	0.04	2.1	0.43	Cell division cycle 42
SEC13	-1.1	0.07	2.0	0.48	SEC13 homolog (S. cerevisiae)
RAB4A	-1.3	0.09	2.0	0.10	RAB4A, member RAS oncogene family
ACACA	-1.3	0.09	2.0	0.29	Acetyl-CoA carboxylase alpha
PCDHLM	1.3	0.07	2.0	0.29	Protein-coding domain containing clathrin assembly protein
WAS	-1.5	0.04	2.0	0.30	WAS, protein involved in Arthrogryposis syndrome
AP1B1	-1.9	0.09	2.0	0.54	Adaptor-related protein complex 1, beta 1 subunit
MAP4K2	-1.4	0.07	2.0	0.55	Mitogen-activated protein kinase kinase kinase kinase 2
AP1M1	1.0	0.02	1.9	0.30	Adaptor-related protein complex 1, mu 2 subunit
STAU1	1.2	0.01	1.9	0.04	Stau1, neuronally-expressed, developmentally down-regulated 4, E3 ubiquitin protein ligase
NSP	0.3	1.46	1.9	0.22	Neuropilin-associated protein
VAPP	-1.4	0.05	1.9	0.10	VAMP (vesicle-associated membrane protein)-associated protein B and C
ARPC2	-1.0	0.05	1.9	0.27	Actin related protein 2/3 complex, subunit 4, 20kDa
RAB5B	-1.8	0.29	1.9	0.17	RAB5B, member RAS oncogene family
IPRK3	-1.7	0.18	1.8	0.17	Irk3, rodent homolog
EZK	-1.2	0.2	1.8	0.69	Ezrin
PTRP9M	-1.9	0.12	1.8	0.24	Phosphatidylinositol transfer protein, membrane-associated 1
NEOD4L	-1.3	0.01	1.8	0.19	Neurod4-like precursor cell expressed, developmentally down-regulated 4-like, E3 ubiquitin protein ligase
TNK	-1.7	0.26	1.8	0.25	TNF receptor-associated factor
CAV1	-1.1	0.01	1.8	0.20	Caveolin 1, caveolae protein, 22kDa
ARPC5	-1.0	0.00	1.8	0.54	Actin related protein 2/3 complex, subunit 5, 16kDa
MFH1CLC3A	0.0	0.12	1.8	0.34	Mfh1, membrane-associated protein 1 light chain 3 alpha
LAMP1	1.1	0.05	1.7	0.34	LAMP1, membrane-associated protein 1
RAB5A	-1.1	0.03	1.6	0.25	RAB5A, member RAS oncogene family
PAK1	-2.8	0.05	1.6	0.04	p21 proto-oncogene protein
DNMS	-0.2	1.65	1.6	0.28	Dynamin 2
CAV2	-1.2	0.06	1.6	0.43	Caveolin 2
MAP4K8P3	1.0	0.14	1.6	0.34	Mitogen-activated protein kinase 8 interacting protein 3
AP2M1	0.0	1.46	1.6	0.18	Adaptor-related protein complex 3, delta 1 subunit
ARPC3	-2.4	0.02	1.6	0.12	Actin related protein 2/3 complex, subunit 3, 16kDa
CAV2C	-2.0	0.20	1.6	0.27	Caveolin 2, isoform 2
PCDHIN1	-1.1	0.03	1.6	0.59	Hepatocyte growth factor-regulated tyrosine kinase substrate
CTBP1	-2.4	1.01	1.6	0.15	Intersectin 2
HGS	-1.7	0.03	1.6	0.51	Intersectin 2
AP2A2	-2.0	0.04	1.6	0.14	Actin-related protein complex 2, alpha 2 subunit
SYNU1	-1.4	0.14	1.3	0.13	Synapsin 1
GIT1	1.3	0.03	1.3	0.02	G protein-coupled receptor kinase interacting ArfGAP 1
POCD6P1	-1.5	0.15	1.3	0.21	Programmed cell death 6 interacting protein
SREBF1	-1.3	0.31	1.3	0.05	Sterol regulatory element binding transcription factor 1
CD63	-1.1	0.01	1.3	0.01	Cd63, prob-oncogene C, E3 ubiquitin protein ligase
RAB9A	-1.2	0.06	1.2	0.06	RAB9A, member RAS oncogene family
EPN3	-1.3	0.01	1.2	0.24	Epnr 3
GORASP1	1.1	0.02	1.2	0.24	Golgi reassembly stacking protein 1, 65kDa
AP4E1	-2.3	0.02	1.2	0.21	Adaptor-related protein complex 4, epsilon 1 subunit
CB1	-2.2	0.20	1.1	0.27	Calmodulin-binding protein 1 (calmyn)
HIP1	-2.0	0.16	1.1	0.31	Interactin strength interacting protein 1
EPS15L1	-2.4	0.19	1.1	0.17	Epidermal growth factor receptor pathway substrate 15-like 1
PKD2L1	-2.3	0.84	1.1	0.70	Protein kinase D1
AP2M2	1.1	0.03	1.0	0.20	Adaptor-related protein interacting protein 2
FASN	-2.5	1.49	1.0	0.47	Fatty acid synthase
CFL1	-1.4	0.42	0.9	0.73	Cofilin 1 (non-muscle)
SYT1	1.1	0.04	0.9	0.67	Syntaptotagmin 1
VA2V	-1.5	0.12	0.9	0.49	VAV 2 guanine nucleotide exchange factor
CLTA	-1.4	0.17	-0.1	1.65	Clathrin, light chain A
PCDHIN3	-2.7	1.18	-0.1	2.65	Membrane kinase C and casein kinase substrate in neurons 3
RAB3A	1.1	0.04	-0.1	1.63	RAB3A, member RAS oncogene family
RAB29	-1.0	0.02	-0.1	1.62	RAB29, member RAS oncogene family
ARF1	-1.3	0.02	-0.2	1.71	ADP-ribosylation factor 1
DNMT1	-1.6	0.04	-0.4	1.97	Dynamine 1
MAP4K8P2	-1.8	0.48	-0.6	2.67	Mitogen-activated protein kinase 8 interacting protein 2
ADM1M10	1.2	0.19	-1.0	2.68	ADM1, metastaplasia domain 10
ARF5	-1.2	0.10	-1.1	2.01	ADP-ribosylation factor 5
CD81	-0.1	1.60	-1.3	0.15	CD81 molecule
EFS	-1.4	0.16	-1.3	0.34	Embryonal Fyn-associated substrate
RAB9B	-1.3	0.17	-1.3	0.36	Embryonal Fyn-associated substrate
CLTC	0.0	1.47	-1.3	0.38	Calmodulin, heavy chain (hc)
RAB11FIP1 [#]	1.2	0.04	-1.5	0.18	RAB11 family interacting protein 5 (class I)
RAF1A	-1.7	0.02	-1.5	0.10	RAF1A, member RAS oncogene family
CAMK1	-1.3	0.05	-1.5	0.34	Calmodulin/calmodulin-dependent protein kinase I
WASF2	-1.7	0.10	-1.5	0.67	WAS protein family, member 2
GNB1L1	-1.0	0.01	-1.5	0.36	Guanine nucleotide binding protein (G protein), beta polypeptide 2-like 1
RAB9C	-1.3	0.05	-1.6	0.38	RAB9C, member RAS oncogene family
EPN1	-1.5	0.16	-1.6	0.68	Epnr 1
RAB6B	-2.4	1.60	-1.7	0.34	RAB6B, member RAS oncogene family
AP2A1	-2.5	0.00	-1.7	0.19	Adaptor-related protein complex 2, alpha 1 subunit
ARF3	-1.9	0.07	-1.7	0.50	ADP-ribosylation factor 3
CET	-1.0	1.52	-1.7	0.57	Cetuin, light chain B
PPK91A	-1.6	0.03	-1.9	0.54	Phosphatidylinositol-4-phosphate 5-kinase, type I, alpha
EPN2	-1.6	0.57	-2.1	1.26	Epnr 2
SYNU2	-1.2	0.06	-2.2	0.69	Synapsin 2
SNAP91	-3.6	0.13	-2.5	0.27	Synaptosomal-associated protein, 91kDa
SART1	-1.1	0.01	-2.5	0.29	Sarco(endo)plasmic reticulum calcium release channel protein 1A
RAB9A	-1.2	0.08	-2.5	1.09	RAB9A, member RAS oncogene family
SH3GLB1 [#]	-1.9	0.08	-2.6	0.91	SH3-domain GRB2-like endophilin B1
PKC3G	-5.3	0.4	-2.8	0.94	Phosphatidylinositol-4-kinase
ARRB2	0.0	1.57	-2.9	1.60	Arsenitin, beta 2
RAF1	-1.4	0.24	-3.3	2.01	RAF1, member C3 boomerang toxic substrate 1 (Rho family, small GTP binding protein Rac1)
RAB11A	-1.7	0.04	-3.4	1.44	Rab11A, member RhoS oncogene family
CBLL	-3.1	0.17	-3.6	1.76	Cbl proto-oncogene B, E3 ubiquitin protein ligase
ATG5	-1.6	0.16	-3.6	0.95	Autophagy related 5
AP1M1	-1.2	0.10	-4.2	1.61	Adaptor-related protein complex 1, mu 1 subunit
CLNT1	-1.4	0.13	-4.8	0.67	Clathrin, light chain 1
BMY	-1.1	0.08	-4.8	0.72	Bruchpflinge integral 1
ATM	-1.5	0.20	-5.7	0.82	ATM serine/threonine kinase
WASF3	-2.0	0.55	-6.1	6.81	WAS protein family, member 3
VPS4A	1.1	0.01	-6.4	0.71	Vacuolar protein sorting 4 homolog A (S. cerevisiae)
DAB2	-3.1	0.38	-6.6	2.48	Death mitogen-responsive phosphoprotein, homolog 2 (Drosophila)
ITSN1	-1.4	0.27	-6.9	2.09	Intersectin 1 (SH3-domain protein)
RAB9B	-1.3	0.57	-7.1	2.79	RAB9B, RAB40 gene family
FYN	-4.1	1.98	-7.9	4.79	FYN proto-oncogene, Src family tyrosine kinase
ASAP2	-1.6	0.37	-9.3	0.42	ArfGAP with SH3 domain, ankyrin repeat and PH domain
PKC3G2	-2.0	0.83	-9.4	3.34	Phosphatidylinositol-4-phosphate 3-kinase, catalytic subunit type 2 gamma
VPS4D	-3.0	0.59	-10.4	0.18	Vacuolar protein sorting 4 homolog D (S. cerevisiae)
ERCI	0.0	1.50	-11.0	0.44	LETS/Rab9-interacting-S-CAST family member 1
COPA	-1.6	0.07	-12.2	0.44	Copart protein complex, subunit alpha
ATP9J/C	-1.5	0.10	-12.3	0.34	ATPase, H+ transporting, lysosomal 16kDa, V0 subunit c
COPB2	-2.0	0.25	-12.3	0.30	Copart protein complex, subunit beta 2 (beta prime)
VCP	-4.3	0.7	-12.4	0.49	Vimentin containing protein