

SYMBOL	FC	P.VALUE	ADJ.P.VAL*	NAME
<i>ARL16</i>	-1.69805	6.49E-06	0.00084	ADP ribosylation factor like GTPase 16
<i>KLHL28</i>	-1.54792	1.99E-07	0.000152	kelch like family member 28
<i>SUCLA2</i>	-1.52808	3.23E-07	0.000178	succinate-CoA ligase ADP-forming beta subunit
<i>ATP6V0C</i>	-1.4514	2.01E-06	0.000444	ATPase H ⁺ transporting V0 subunit c
<i>DNAJC12</i>	-1.43653	1.88E-07	0.000152	DnaJ heat shock protein family (Hsp40) member C12
<i>SLC35E1</i>	-1.42346	1.33E-07	0.000152	solute carrier family 35 member E1
<i>UHMK1</i>	-1.39353	1.82E-06	0.000443	U2AF homology motif kinase 1
<i>POLR3A</i>	-1.37816	1.91E-06	0.000443	RNA polymerase III subunit A
<i>AACS</i>	-1.37704	1.75E-06	0.000443	acetoacetyl-CoA synthetase
<i>ISCU</i>	-1.37026	3.91E-06	0.000687	iron-sulfur cluster assembly enzyme
<i>SYAP1</i>	-1.3691	5.82E-06	0.000799	synapse associated protein 1
<i>AP2S1</i>	-1.36662	6.39E-06	0.00084	adaptor related protein complex 2 sigma 1 subunit
<i>RAN</i>	-1.36345	4.76E-09	1.09E-05	RAN member RAS oncogene family
<i>UCHL1</i>	-1.35504	4.01E-06	0.000687	ubiquitin C-terminal hydrolase L1
<i>GSKIP</i>	-1.35441	5.50E-07	0.00023	GSK3B interacting protein
<i>HSP90AA1</i>	-1.34292	3.58E-06	0.000664	heat shock protein 90 alpha family class A member 1
<i>ISG20L2</i>	-1.31859	1.10E-06	0.000397	interferon stimulated exonuclease gene 20 like 2
<i>MAP2K4</i>	-1.30635	4.37E-06	0.000691	mitogen-activated protein kinase kinase 4
<i>ACTR10</i>	-1.29453	6.87E-06	0.000873	actin related protein 10 homolog
<i>SMS</i>	-1.28592	9.13E-08	0.000125	spermine synthase
<i>CNOT7</i>	-1.28154	1.30E-06	0.000424	CCR4-NOT transcription complex subunit 7
<i>HEATR5B</i>	-1.26606	2.74E-06	0.000554	HEAT repeat containing 5B
<i>ISCA1</i>	-1.24402	1.40E-06	0.000436	iron-sulfur cluster assembly 1
<i>GTF2H3</i>	-1.24333	5.55E-06	0.000777	general transcription factor IIH subunit 3
<i>COPS8</i>	-1.22979	5.10E-06	0.00073	COP9 signalosome subunit 8
<i>KRIT1</i>	-1.21943	5.94E-06	0.000799	KRIT1 ankyrin repeat containing
<i>TROVE2</i>	1.220267	3.66E-07	0.000178	TROVE domain family member 2
<i>ZNF22</i>	1.228388	2.91E-07	0.000178	zinc finger protein 22
<i>GPBP1L1</i>	1.245112	5.02E-06	0.00073	GC-rich promoter binding protein 1 like 1
<i>SLC48A1</i>	1.25577	1.98E-07	0.000152	solute carrier family 48 member 1
<i>TRAM1</i>	1.275035	2.55E-06	0.00053	translocation associated membrane protein 1
<i>LSM14A</i>	1.30744	3.94E-06	0.000687	LSM14A mRNA processing body assembly factor
<i>PTAR1</i>	1.323362	1.65E-06	0.000443	protein prenyltransferase alpha subunit repeat containing 1
<i>HMGB1</i>	1.330248	4.52E-06	0.000691	high mobility group box 1
<i>PHLDB1</i>	1.332512	4.86E-06	0.000725	pleckstrin homology like domain family B member 1
<i>PRRG1</i>	1.337767	1.83E-06	0.000443	proline rich and Gla domain 1
<i>PTPN11</i>	1.346854	4.14E-06	0.000691	protein tyrosine phosphatase
<i>Sep 02</i>	1.362179	1.55E-06	0.000443	septin 2
<i>PACSL1</i>	1.382089	3.29E-06	0.000628	protein kinase C and casein kinase substrate in neurons 3
<i>TRAK2</i>	1.404963	4.53E-06	0.000691	trafficking kinesin protein 2
<i>CDK19</i>	1.412032	4.16E-08	7.14E-05	cyclin dependent kinase 19
<i>FNBP1</i>	1.41257	9.09E-10	6.24E-06	formin binding protein 1
<i>NDRG1</i>	1.415055	1.67E-06	0.000443	N-myc downstream regulated 1
<i>PCBP4</i>	1.430148	3.12E-06	0.000611	poly(rC) binding protein 4
<i>COL9A2</i>	1.432579	2.10E-06	0.00045	collagen type IX alpha 2 chain
<i>RASSF2</i>	1.442064	7.34E-07	0.00028	Ras association domain family member 2
<i>YES1</i>	1.443003	2.38E-07	0.000164	YES proto-oncogene 1
<i>MARCKSL1</i>	1.462477	4.50E-06	0.000691	MARCKS like 1
<i>CNP</i>	1.501119	2.31E-09	7.94E-06	2'3'-cyclic nucleotide 3' phosphodiesterase

[illegible]

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LOC285074	-1.50537	5.94E-09	4.08E-05	anaphase promoting complex subunit 1 pseudogene