

**Supplemental Table 13. Intersection with Neural Stem Cell Associated Genes.**

<b><u>Probe</u></b>	<b><u>Symbol</u></b>	<b><u>Description</u></b>	<b><u>P-value</u></b>	<b><u>Fold-Change</u></b>
ADXEC.31458.C1_at	POLN	polymerase (DNA directed) nu (POLN) gene complete cds.	0.0244	4.91
ADXEC.31458.C1_x_at	POLN	polymerase (DNA directed) nu (POLN) gene complete cds.	0.0268	3.63
ADXEC.7503.C1_s_at	DPP3	dipeptidyl-peptidase 3 (DPP3) transcript variant 1	0.0214	3.50
ADXEC.4766.C1-a_s_at	LAS1L	LAS1-like (S. cerevisiae) (LAS1L) transcript variant	0.0233	3.37
ADXECNTDJ.3576_s_at	VCP	valosin-containing protein (VCP) mRNA.	0.0118	3.24
ADXEC.652.C3-a_s_at	RFC5	replication factor C (activator 1) 5 36.5kDa (RFC5)	0.0022	3.03
ADXEC.5487.C2_s_at	PNKD	paroxysmal nonkinesigenic dyskinesia (PNKD) nuclear	0.05	3.01
ADXEC.34747.C1_s_at	SRR	Serine racemase	0.0247	2.91
ADXEC.9076.C1_s_at	PKM2	pyruvate kinase muscle (PKM2) gene complete cds.	0.0431	2.89
ADXECAD.7889_at	GATM	glycine amidinotransferase (L-arginine:glycine	0.0227	2.86
ADXEC.23283.C1_s_at	EZH2	enhancer of zeste homolog 2 (Drosophila) (EZH2)	0.0213	2.77
ADXECADA.18168_at	CHAF1A	SH3-containing protein EEN (EEN) and chromatin assemblyfactor-I p15	0.0114	2.75
ADXEC.1467.C1_at	UTP18	UTP18 small subunit (SSU) processome component homolog (yeast)	0.0421	2.71
ADXEC.16051.C2_s_at	PTPRJ	protein tyrosine phosphatase receptor type J	0.02	2.64
ADXEC.27079.C1_at	RAD1	RAD1 homolog (S. pombe) (RAD1) transcript variant 1	0.0102	2.63
ADXEC.6850.C1-a_s_at	DNAJB6	DnaJ (Hsp40) homolog subfamily B member 6 (DNAJB6)	0.0423	2.63
ADXPCEC.11099.C1_at	GSPT1	Human chromosome 16p13.1 BAC clone CIT987SK-551G9 complete sequence.	0.0034	2.62
ADXECADA.18979_s_at	USP38	ubiquitin specific peptidase 38 (USP38) mRNA.	0.05	2.58
ADXEC.6333.C1_at	RPUSD4	RNA pseudouridylate synthase domain containing 4	0.0012	2.53
ADXEC.14462.C1_at	PCYT1A	phosphate cytidyltransferase 1 choline alpha	0.05	2.48
ADXEC.12118.C1_s_at	BCL2L13	BCL2-like 13 (apoptosis facilitator) (BCL2L13)	0.0417	2.44
ADXEC.6876.C1_at	SRPRB	signal recognition particle receptor B subunit	0.05	2.42
ADXEC.12903.C1_s_at	TRIP6	thyroid hormone receptor interactor 6 (TRIP6) mRNA.	0.0047	2.38
ADXECNTDJ.3859_at	TMEM165	Transmembrane protein 165	0.0097	2.37
ADXECAD.7010_at	PARL	presenilin associated rhomboid-like (PARL) nuclear	0.05	2.35
ADXECNTDJ.3106_at	TNC	tenascin C (TNC) mRNA.	0.05	2.35
ADXEC.14462.C1_x_at	PCYT1A	phosphate cytidyltransferase 1 choline alpha	0.0416	2.34
ADXEC.6557.C2_s_at	HTATIP2	HIV-1 Tat interactive protein 2 30kDa (HTATIP2)	0.0231	2.33
ADXECAD.12804_x_at	PTPN1	Protein tyrosine phosphatase non-receptor type 1	0.0279	2.33
ADXEC.487.C1-a_s_at	RNF14	ring finger protein 14 (RNF14) transcript variant 2	0.0152	2.3

ADXECNTDJ.1237_at	KPNA1	karyopherin alpha 1 (importin alpha 5) (KPNA1)	0.05	2.28
ADXEC.10647.C1-a_s_at	ZNF644	zinc finger protein 644 (ZNF644) transcript variant	0.0355	2.28
ADXEC.1226.C2_x_at	TYMS	gene for thymidylate synthase complete cds.	0.017	2.27
ADXEC.5973.C1_at	RMI1	RMI1 RecQ mediated genome instability 1 homolog (S.	0.05	2.24
ADXEC.1226.C2_at	TYMS	gene for thymidylate synthase complete cds.	0.0134	2.24
ADXECAD.17716_x_at	PTPN1	Protein tyrosine phosphatase non-receptor type 1	0.0087	2.21
Adx-200054-up_s_at	ZNF259	zinc finger protein 259 (ZNF259) mRNA.	0.0413	2.21
ADXEC.15673.C1_s_at	RAB8B	RAB8B member RAS oncogene family (RAB8B) mRNA.	0.0389	2.19
ADXEC.4103.C1_at	CSNK2A2	casein kinase 2 alpha prime polypeptide (CSNK2A2)	0.0122	2.19
ADXECAD.28846_s_at	CASP2	caspase 2 apoptosis-related cysteine peptidase	0.0117	2.19
ADXECRS.39826_at	PGK1	phosphoglycerate kinase 1 (PGK1) mRNA.	0.05	2.16
ADXEC.20111.C1_x_at	SFXN1	sideroflexin 1 (SFXN1) mRNA.	0.0338	2.15
ADXECRS.6440_s_at	STRBP	spermatid perinuclear RNA binding protein (STRBP)	0.0132	2.14
ADXEC.24458.C1-a_s_at	DTYMK	Deoxythymidylate kinase (thymidylate kinase)	0.0154	2.12
ADXECAD.16766_x_at	MRPL9	mitochondrial ribosomal protein L9 (MRPL9) nuclear	0.0034	2.11
ADXECAD.24402_s_at	SELT	selenoprotein T (SELT) mRNA.	0.05	2.11
ADXECNTDJ.2671_at	KCNAB3	potassium voltage-gated channel shaker-related subfamily beta member 3	0.05	2.11
ADXECNTDJ.6300_at	AP1G1	adaptor-related protein complex 1 gamma 1 subunit	0.029	2.10
ADXEC.12903.C3_s_at	TRIP6	thyroid hormone receptor interactor 6 (TRIP6) mRNA.	0.0138	2.09
ADXECRS.27012_at	STX7	syntaxin 7 (STX7) mRNA.	0.0073	2.07
ADXEC.28173.C1-a_s_at	C14orf106	chromosome 14 open reading frame 106 (C14orf106)	0.0089	2.05
ADXECAD.1657_at	SAE1	SUMO1 activating enzyme subunit 1 (SAE1) transcript	0.0375	2.05
ADXEC.7491.C1_at	DNAJC9	DnaJ (Hsp40) homolog subfamily C member 9	0.05	2.05
ADXEC.2392.C9_x_at	CHAF1A	SH3-containing protein EEN (EEN) and chromatin assemblyfactor-I p15	0.05	2.02
ADXECAD.20979_s_at	ADAM10	ADAM metallopeptidase domain 10 (ADAM10) mRNA.	0.0011	2.00
ADXEC.11266.C1_at	PELI1	Pellino homolog 1 (Drosophila)	0.0226	1.99
ADXEC.111.C99_s_at	USP22	ubiquitin specific peptidase 22 (USP22) mRNA.	0.04	1.98
ADXECAD.11485_s_at	RANBP9	RAN binding protein 9 (RANBP9) mRNA.	0.0053	1.98
ADXEC.2920.C1_at	CHD7	Chromodomain helicase DNA binding protein 7	0.05	1.97
ADXECAD.17716_at	PTPN1	Protein tyrosine phosphatase non-receptor type 1	0.0163	1.97
ADXECNTDJ.1492_at	TAF15	TAF15 RNA polymerase II TATA box binding protein(TBP)- associated fa	0.05	1.96
ADXECADA.20527_s_at	ABCF2	ATP-binding cassette sub-family F (GCN20) member 2	0.0135	1.96
ADXECRS.13593_at	PDAP1	PDGFA associated protein 1 (PDAP1) mRNA.	0.038	1.93
ADXECAD.26276_s_at	GNA13	guanine nucleotide binding protein (G protein) alpha	0.0102	1.92

ADXEC.3840.C1_at	MFN1	mitofusin 1 (MFN1) nuclear gene encoding	0.028	1.92
ADXEC.33954.C1_at	TBRG4	mRNA for KIAA0948 protein partial cds.	0.0221	1.92
ADXECAD.18989_at	CHD7	chromodomain helicase DNA binding protein 7 (CHD7)	0.0346	1.92
ADXEC.9451.C1_s_at	BCL2L13	BCL2-like 13 (apoptosis facilitator) (BCL2L13)	0.0005	1.90
ADXEC.11501.C1_s_at	NUP93	nucleoporin 93kDa (NUP93) mRNA.	0.0166	1.89
ADXECADA.22882_s_at	PDCD2	programmed cell death 2 (PDCD2) transcript variant 1	0.0375	1.89
ADXEC.1909.C1_at	ETF1	eukaryotic translation termination factor 1 (ETF1)	0.0027	1.88
ADXECAD.19332_s_at	CFL2	cofilin 2 (muscle) (CFL2) transcript variant 1 mRNA.	0.052	1.88
ADXECNTDJ.4966_s_at	SPATA5	spermatogenesis associated 5 (SPATA5) mRNA.	0.0411	1.88
ADXPCEC.11965.C1_at	TMEM165	Transmembrane protein 165	0.0092	1.87
ADXECAD.16766_at	MRPL9	mitochondrial ribosomal protein L9 (MRPL9) nuclear	0.0051	1.87
ADXEC.16224.C1_at	SCPEP1	Serine carboxypeptidase 1	0.0347	1.87
ADXEC.3279.C1-a_s_at	ARID1A	AT rich interactive domain 1A (SWI-like) (ARID1A)	0.0213	1.86
ADXEC.16104.C2_s_at	NUBP1	nucleotide binding protein 1 (MinD homolog E. coli)	0.0384	1.84
ADXECNTDJ.5398_s_at	DUSP16	Dual specificity phosphatase 16	0.05	1.83
ADXECRS.8012_at	DGAT1	diacylglycerol O-acyltransferase homolog 1 (mouse) mRNA (cDNAclone)	0.0424	1.82
ADXECAD.22713_s_at	GAB1	GRB2-associated binding protein 1 (GAB1) transcript	0.0427	1.82
ADXEC.3751.C2_at	TBC1D1	TBC1 (tre-2/USP6 BUB2 cdc16) domain family member 1	0.0157	1.81
ADXEC.409.C32_at	PTP4A2	Transcribed locus	0.0357	1.80
ADXECAD.22145_s_at	GSR	glutathione reductase (GSR) mRNA.	0.0363	1.79
ADXEC.3211.C2_s_at	RBM39	RNA binding motif protein 39	0.0417	1.79
ADXEC.11161.C1-a_s_at	KIF5B	kinesin family member 5B (KIF5B) mRNA.	0.0217	1.79
ADXECRS.6844_at	ADAM10	ADAM metallopeptidase domain 10 (ADAM10) mRNA.	0.037	1.78
ADXEC.409.C56_s_at	PTP4A2	Transcribed locus	0.0398	1.77
ADXEC.6728.C2_at	PTP4A1	protein tyrosine phosphatase type IVA member 1	0.0382	1.75
ADXEC.3642.C2_s_at	MLF2	myeloid leukemia factor 2 (MLF2) transcript variant	0.032	1.74
ADXEC.7693.C1_x_at	MBTD1	mbt domain containing 1 (MBTD1) mRNA.	0.0341	1.74
ADXEC.14337.C2_at	PPP1R15B	protein phosphatase 1 regulatory (inhibitor) subunit cDNA FLJ58787 complete cds highly similar to Cleavagestimulation fa	0.0107	1.73
ADXEC.14780.C2_s_at	CSTF2	polymerase (DNA directed) epsilon 3 (p17 subunit)	0.0214	1.72
ADXEC.1048.C1_x_at	POLE3	polymerase (DNA directed) epsilon 3 (p17 subunit)	0.0238	1.72
ADXEC.9082.C1-a_s_at	SRP68	signal recognition particle 68kDa (SRP68) mRNA.	0.05	1.72
ADXEC.7383.C1_x_at	ZNF639	zinc finger protein 639 (ZNF639) mRNA.	0.0033	1.71
ADXEC.33954.C1_x_at	TBRG4	mRNA for KIAA0948 protein partial cds.	0.0078	1.71
ADXEC.7383.C1_at	ZNF639	zinc finger protein 639 (ZNF639) mRNA.	0.0264	1.70

ADXEC.9347.C1_s_at	UFC1	Ubiquitin-fold modifier conjugating enzyme 1	0.05	1.70
ADXECAD.11135_at	SFRS3	Splicing factor arginine/serine-rich 3	0.0168	1.69
ADXEC.14372.C2_s_at	WEE1	WEE1 homolog (S. pombe) (WEE1) transcript variant 1	0.05	1.69
ADXEC.2954.C1_at	SERP1	stress-associated endoplasmic reticulum protein 1	0.0081	1.68
ADXEC.3268.C1_at	ASF1B	ASF1 anti-silencing function 1 homolog B	0.0233	1.67
ADXEC.7693.C1_at	MBTD1	mbt domain containing 1 (MBTD1) mRNA.	0.05	1.67
ADXEC.22227.C1_s_at	FNBP1	Formin binding protein 1	0.0132	1.66
ADXEC.9171.C1_at	USP34	Ubiquitin specific peptidase 34	0.05	1.65
ADXEC.29585.C1_at	C11orf58	Chromosome 11 open reading frame 58	0.0424	1.64
ADXEC.6014.C1_at	CAND1	Cullin-associated and neddylation-dissociated 1	0.0317	1.64
ADXECS.19560_s_at	BZW1	basic leucine zipper and W2 domains 1 (BZW1) mRNA.	0.0221	1.64
ADXECADA.18789_at	CENPK	centromere protein K (CENPK) mRNA.	0.0365	1.64
ADXEC.14865.C2_s_at	SS18	synovial sarcoma translocation chromosome 18 (SS18)	0.0327	1.64
ADXPCEC.16122.C1_at	HIATL1	hippocampus abundant transcript-like 1	0.05	1.63
ADXECAD.14311_at	UFM1	Ubiquitin-fold modifier 1	0.0084	1.61
ADXPCEC.11965.C1_x_at	TMEM165	Transmembrane protein 165	0.0363	1.61
ADXEC.23157.C1_s_at	GNAI3	guanine nucleotide binding protein (G protein) alpha inhibiting activity polypep	0.0316	1.60
ADXEC.7681.C1_s_at	SMC2	structural maintenance of chromosomes 2 (SMC2)	0.0051	1.60
ADXEC.4544.C1_at	RGS19	regulator of G-protein signaling 19 (RGS19)	0.0436	1.60
ADXEC.7696.C1-a_s_at	STX12	syntaxin 12 (STX12) mRNA.	0.05	1.57
ADXECAD.16985_s_at	PA2G4	proliferation-associated 2G4 38kDa (PA2G4) mRNA.	0.05	1.57
ADXEC.1690.C5_s_at	GNB1	guanine nucleotide binding protein (G protein) beta	0.0021	1.56
ADXECS.4234_s_at	CAP1	CAP adenylate cyclase-associated protein 1 (yeast)	0.0265	1.56
ADXECAD.9888_x_at	ROCK1	Rho-associated coiled-coil containing protein kinase	0.042	1.56
ADXECEMUTR.1821_s_at	PVRL3	poliovirus receptor-related 3 mRNA (cDNA cloneIMAGE:4066485) comple	0.05	1.56
ADXEC.5984.C1_at	ARF4	ADP-ribosylation factor 4 (ARF4) mRNA.	0.05	1.55
ADXEC.2886.C2-a_s_at	GOLPH3	golgi phosphoprotein 3 (coat-protein) (GOLPH3) mRNA.	0.0146	1.55
ADXECAD.23957_s_at	BACH1	BTB and CNC homology 1 basic leucine zipper	0.0088	1.54
ADXEC.2959.C3_s_at	COPG	coatomer protein complex subunit gamma (COPG) mRNA.	0.05	1.54
ADXEC.16562.C1_s_at	CCNJ	cyclin J (CCNJ) transcript variant 3 mRNA.	0.0251	1.54
ADXEC.6599.C1_s_at	FAM50A	family with sequence similarity 50 member A (FAM50A)	0.05	1.54
ADXEC.6728.C2_x_at	PTP4A1	protein tyrosine phosphatase type IVA member 1	0.05	1.53
ADXECAD.9650_at	MAT2A	methionine adenosyltransferase II alpha (MAT2A)	0.0263	1.53
ADXECAD.9888_at	ROCK1	Rho-associated coiled-coil containing protein kinase	0.0043	1.53

ADXECADA.13461_at	PPIE	Peptidylprolyl isomerase E (cyclophilin E)	0.0285	1.53
ADXEC.2744.C2_s_at	CTTN	cortactin (CTTN) transcript variant 1 mRNA.	0.0451	1.53
ADXEC.719.C1_s_at	LMNB2	lamin B2 (LMNB2) mRNA.	0.0343	1.52
ADXEC.1114.C2-a_s_at	CAPZA1	capping protein (actin filament) muscle Z-line alpha	0.0212	1.52
ADXEC.2206.C2_at	NDUFC1	NADH dehydrogenase (ubiquinone) 1 subcomplex unknown 1 6kDa	0.032	1.52
ADXECAD.1571_s_at	ASAH1	N-acylsphingosine amidohydrolase (acid ceramidase) 1	0.037	1.52
ADXECADA.16206_at	ARFGEF1	ADP-ribosylation factor guanine nucleotide-exchange factor 1(brefeldin A-inhibit	0.0338	1.52
ADXECADA.24708_s_at	TNKS2	tankyrase TRF1-interacting ankyrin-related ADP-ribose	0.05	1.52
ADXPCEC.15490.C1_s_at	SMAD2	SMAD family member 2 (SMAD2) transcript variant 1	0.0206	1.51
ADXOCEC.2903.C1_s_at	UBE2D2	ubiquitin-conjugating enzyme E2D 2 (UBC4/5 homolog	0.0138	1.51
ADXEC.1260.C8_s_at	PSMB4	Proteasome (prosome macropain) subunit beta type 4	0.0297	1.51
ADXEC.2291.C1_x_at	NDUFV2	NDUFV2 gene for 24-kDa subunit of complex I partial cds.	0.0189	1.51
ADXEC.11953.C2_s_at	APAF1	apoptotic peptidase activating factor 1 (APAF1)	0.0069	1.51
ADXECAD.26276_at	GNA13	guanine nucleotide binding protein (G protein) alpha	0.0273	1.51
ADXECADA.253_s_at	KIAA1524	KIAA1524 (KIAA1524) mRNA.	0.0123	1.50
ADXECEMUTR.410_at	WDR26	WD repeat domain 26 (WDR26) transcript variant 1	0.0473	1.49
ADXEC.7491.C1_x_at	DNAJC9	DnaJ (Hsp40) homolog subfamily C member 9	0.05	1.48
ADXECAD.1_at	SCARB1	Scavenger receptor class B member 1	0.0261	1.48
ADXECADA.21050_at	SLC4A7	solute carrier family 4 sodium bicarbonate	0.0227	1.47
ADXECADA.17436_s_at	FOXK2	forkhead box K2 (FOXK2) mRNA.	0.0252	1.47
ADXECAD.23807_s_at	ING1	inhibitor of growth family member 1 (ING1)	0.0342	1.46
ADXECNTDJ.676_at	FIP1L1	FIP1 like 1 (S. cerevisiae) (FIP1L1) transcript	0.0061	1.46
ADXEC.2577.C1-a_s_at	ABCE1	ATP-binding cassette sub-family E (OABP) member 1	0.0432	1.46
ADXEC.1945.C5_s_at	IPO7	importin 7 (IPO7) mRNA.	0.023	1.46
ADXEC.121.C1_s_at	ARL5A	ADP-ribosylation factor-like 5A (ARL5A) transcript	0.0123	1.46
ADXEC.505.C3_x_at	MPHOSPH10	M-phase phosphoprotein 10 (U3 small nucleolar	0.0251	1.45
ADXEC.11961.C1_at	AQR	aquarius homolog (mouse) (AQR) mRNA.	0.018	1.44
ADXECAD.4765_x_at	LIMS1	LIM and senescent cell antigen-like domains 1	0.0267	1.44
ADXECADA.1528_s_at	CCNJ	cyclin J (CCNJ) transcript variant 3 mRNA.	0.0207	1.44
ADXEC.15900.C1-a_s_at	FAM60A	family with sequence similarity 60 member A (FAM60A)	0.0263	1.43
ADXECAD.20409_s_at	IVNS1ABP	influenza virus NS1A binding protein (IVNS1ABP) mRNA.	0.0422	1.43
ADXECAD.20345_s_at	RAD23B	RAD23 homolog B (S. cerevisiae) (RAD23B) mRNA.	0.0065	1.42
ADXECNTDJ.4261_s_at	ANAPC4	anaphase promoting complex subunit 4 (ANAPC4) mRNA.	0.0335	1.42
ADXEC.1119.C2_s_at	HMGB2	high-mobility group box 2 (HMGB2) transcript variant	0.0455	1.40

ADXEC.20377.C1-a_s_at	SOAT1	sterol O-acyltransferase 1 (SOAT1) transcript variant	0.0241	1.40
ADXECAD.5485_at	INTS7	Integrator complex subunit 7	0.0376	1.40
ADXEC.5198.C2_s_at	HSPA4	heat shock 70kDa protein 4 (HSPA4) mRNA.	0.0442	1.40
ADXEC.19665.C1-a_s_at	NNT	nicotinamide nucleotide transhydrogenase (NNT)	0.05	1.39
ADXEC.25009.C1_x_at	ZRANB2	Zinc finger RAN-binding domain containing 2	0.0074	1.39
ADXEC.7650.C2_at	TNPO1	transportin 1 (TNPO1) transcript variant 2 mRNA.	0.036	1.38
ADXEC.913.C1_at	SLTM	SAFB-like transcription modulator (SLTM) transcript	0.0306	1.37
ADXEC.33040.C1_x_at	KIAA1161	KIAA1161 (KIAA1161) mRNA.	0.0202	1.36
ADXEC.5035.C3_x_at	PABPC1	poly(A) binding protein cytoplasmic 1 (PABPC1) mRNA.	0.0417	1.35
ADXEC.1104.C1_at	PSMD12	proteasome (prosome macropain) 26S subunit	0.0389	1.34
ADXEC.6634.C2_at	PAPOLA	Poly(A) polymerase alpha	0.0113	1.34
ADXEC.8227.C2_s_at	ARF6	ADP-ribosylation factor 6 (ARF6) mRNA.	0.0159	1.34
ADXEC.9641.C1_s_at	ETF1	eukaryotic translation termination factor 1 (ETF1)	0.0209	1.34
ADXECAD.9838_s_at	RRBP1	Ribosome binding protein 1 homolog 180kDa (dog)	0.0054	1.34
ADXEC.7837.C1_at	ANP32E	acidic (leucine-rich) nuclear phosphoprotein 32	0.0079	1.33
ADXEC.2424.C1_s_at	SERP1	stress-associated endoplasmic reticulum protein 1	0.0141	1.32
ADXEC.3894.C1_at	SH3GLB1	SH3-domain GRB2-like endophilin B1 (SH3GLB1) mRNA.	0.0238	1.32
ADXEC.12233.C1_at	PYCR2	pyrroline-5-carboxylate reductase family member 2	0.0418	1.31
ADXEC.6911.C2_s_at	PKN2	protein kinase N2 (PKN2) mRNA.	0.035	1.31
ADXEC.7849.C1_at	KIF5B	kinesin family member 5B (KIF5B) mRNA.	0.0283	1.31
ADXEC.3334.C1-a_s_at	STX7	syntaxin 7 (STX7) mRNA.	0.0354	1.29
ADXECAD.19045_at	TM9SF3	transmembrane 9 superfamily member 3 (TM9SF3) mRNA.	0.0027	1.29
ADXEC.11850.C1_at	RSU1	Ras suppressor protein 1 (RSU1) transcript variant 1	0.05	1.28
ADXEC.2886.C1_at	GOLPH3	golgi phosphoprotein 3 (coat-protein) (GOLPH3) mRNA.	0.05	1.26
ADXEC.21117.C1_at	ROD1	ROD1 regulator of differentiation 1 (S. pombe) (ROD1)	0.05	1.26
ADXEC.3576.C1_x_at	LAMP2	lysosomal-associated membrane protein 2 (LAMP2)	0.0137	1.25
ADXECRS.13131_at	BIRC5	baculoviral IAP repeat-containing 5 (BIRC5)	0.05	1.25
ADXECAD.12808_at	PTPN1	Protein tyrosine phosphatase non-receptor type 1	0.0239	1.24
ADXECADA.10529_s_at	ARF6	ADP-ribosylation factor 6 (ARF6) mRNA.	0.05	1.23
ADXECRS.13131_x_at	BIRC5	baculoviral IAP repeat-containing 5 (BIRC5)	0.0444	1.22
ADXEC.21002.C1_at	EIF4G2	eukaryotic translation initiation factor 4 gamma 2	0.036	1.21
ADXEC.242.C1_x_at	UBE2D3	ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog	0.0232	1.20
ADXEC.10563.C1_at	SMARCA4	SWI/SNF related matrix associated actin dependent regulator of chromatin	0.0277	1.19
ADXEC.8227.C1-a_s_at	ARF6	ADP-ribosylation factor 6 (ARF6) mRNA.	0.0313	1.17
ADXEC.3746.C2-a_s_at	KIF2C	kinesin family member 2C (KIF2C) mRNA.	0.0353	1.16

ADXEC.3261.C2\_at

TM7SF2

putative sterol reductase SR-1 (TM7SF2) gene complete cds. 0.0264

1.15

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