

Supplemental Figure 1. SNAI2 dependent gene expression changes in UM-SCC-38 cells.

P-value	FDR	Fold-change (Slug/Neo Control)	Probe set	Gene symbol	Description
4.00E-07	3.77E-05	125.764	<a href="#">215076_s_at</a>	<a href="#">COL3A1</a>	collagen, type III, alpha 1
6.50E-06	8.10E-05	77.033	<a href="#">212464_s_at</a>	<a href="#">FN1</a>	fibronectin 1
4.00E-07	3.77E-05	73.860	<a href="#">211719_x_at</a>	<a href="#">FN1</a>	fibronectin 1
1.80E-06	5.65E-05	68.125	<a href="#">201852_x_at</a>	<a href="#">COL3A1</a>	collagen, type III, alpha 1
9.00E-07	5.36E-05	60.721	<a href="#">216442_x_at</a>	<a href="#">FN1</a>	fibronectin 1
5.30E-06	7.44E-05	52.226	<a href="#">211161_s_at</a>	<a href="#">COL3A1</a>	collagen, type III, alpha 1
1.50E-06	5.65E-05	50.446	<a href="#">210495_x_at</a>	<a href="#">FN1</a>	fibronectin 1
3.00E-07	3.57E-05	34.892	<a href="#">225242_s_at</a>	<a href="#">CCDC80</a>	coiled-coil domain containing 80
1.00E-07	3.57E-05	31.631	<a href="#">212764_at</a>	<a href="#">ZEB1</a>	zinc finger E-box binding homeobox 1
1.70E-06	5.65E-05	31.440	<a href="#">201280_s_at</a>	<a href="#">DAB2</a>	disabled homolog 2, mitogen-responsive phosphoprotein (Drosophila)
6.80E-06	8.13E-05	29.978	<a href="#">202627_s_at</a>	<a href="#">SERPINE1</a>	serpin peptidase inhibitor, clade E (nexin, plasminogen activator inhibitor type 1), member 1
3.58E-05	0.0001783	29.051	<a href="#">235236_at</a>	<a href="#">LOC100131897</a>	Uncharacterized protein LOC100131897
7.60E-06	8.38E-05	28.914	<a href="#">202628_s_at</a>	<a href="#">SERPINE1</a>	serpin peptidase inhibitor, clade E (nexin, plasminogen activator inhibitor type 1), member 1
1.20E-06	5.65E-05	28.279	<a href="#">201278_at</a>	<a href="#">DAB2</a>	disabled homolog 2, mitogen-responsive phosphoprotein (Drosophila)
3.00E-07	3.57E-05	27.903	<a href="#">202998_s_at</a>	<a href="#">LOXL2</a>	lysyl oxidase-like 2
5.58E-05	0.0002302	25.733	<a href="#">203700_s_at</a>	<a href="#">DIO2</a>	deiodinase, iodothyronine, type II
1.89E-05	0.0001291	25.347	<a href="#">226218_at</a>	<a href="#">IL7R</a>	interleukin 7 receptor
0.0001374	0.0003977	24.015	<a href="#">231240_at</a>	<a href="#">DIO2</a>	deiodinase, iodothyronine, type II
2.70E-06	6.21E-05	23.453	<a href="#">218332_at</a>	<a href="#">BEX1</a>	brain expressed, X-linked 1
1.90E-06	5.65E-05	23.422	<a href="#">201426_s_at</a>	<a href="#">VIM</a>	vimentin
1.90E-06	5.65E-05	22.666	<a href="#">215446_s_at</a>	<a href="#">LOX</a>	lysyl oxidase
3.00E-07	3.57E-05	20.409	<a href="#">210757_x_at</a>	<a href="#">DAB2</a>	disabled homolog 2, mitogen-responsive phosphoprotein (Drosophila)
2.97E-05	0.0001658	19.681	<a href="#">235004_at</a>	<a href="#">RBM24</a>	RNA binding motif protein 24
3.40E-06	6.74E-05	18.980	<a href="#">203699_s_at</a>	<a href="#">DIO2</a>	deiodinase, iodothyronine, type II
4.00E-07	3.77E-05	17.842	<a href="#">201279_s_at</a>	<a href="#">DAB2</a>	disabled homolog 2, mitogen-responsive phosphoprotein (Drosophila)
1.80E-06	5.65E-05	17.442	<a href="#">239218_at</a>	<a href="#">PDE1C</a>	phosphodiesterase 1C, calmodulin-dependent 70kDa
8.49E-05	0.0003004	16.772	<a href="#">219935_at</a>	<a href="#">ADAMTSS5</a>	ADAM metallopeptidase with thrombospondin type 1 motif, 5
1.00E-07	3.57E-05	16.169	<a href="#">220014_at</a>	<a href="#">PRR16</a>	proline rich 16
3.12E-05	0.0001676	16.004	<a href="#">215767_at</a>	<a href="#">ZNF804A</a>	zinc finger protein 804A
9.70E-06	9.41E-05	15.927	<a href="#">231867_at</a>	<a href="#">ODZ2</a>	odz, odd Oz/ten-m homolog 2 (Drosophila)
1.90E-06	5.65E-05	15.657	<a href="#">222862_s_at</a>	<a href="#">AK5</a>	adenylate kinase 5
7.10E-06	8.26E-05	15.590	<a href="#">224823_at</a>	<a href="#">MYLK</a>	myosin light chain kinase
4.70E-06	7.04E-05	13.947	<a href="#">201645_at</a>	<a href="#">TNC</a>	tenascin C
0.0001379	0.0003977	13.742	<a href="#">1555673_at</a>	<a href="#">KRTAP2-1</a>	keratin associated protein 2-1
9.70E-06	9.41E-05	13.477	<a href="#">204584_at</a>	<a href="#">L1CAM</a>	L1 cell adhesion molecule
1.40E-06	5.65E-05	13.437	<a href="#">211340_s_at</a>	<a href="#">MCAM</a>	melanoma cell adhesion molecule
1.00E-07	3.57E-05	13.347	<a href="#">203060_s_at</a>	<a href="#">PAPSS2</a>	3-adenosine 5-phosphosulfate synthase 2
2.46E-05	0.0001487	13.265	<a href="#">211959_at</a>	<a href="#">IGFBP5</a>	insulin-like growth factor binding protein 5
8.70E-06	9.19E-05	12.976	<a href="#">236344_at</a>	<a href="#">PDE1C</a>	phosphodiesterase 1C, calmodulin-dependent 70kDa
4.69E-05	0.0002039	12.950	<a href="#">209101_at</a>	<a href="#">CTGF</a>	connective tissue growth factor

0.0002393	0.0005894	12.705	<a href="#">1570155_at</a>	<a href="#">NA</a>	NA
1.30E-06	5.65E-05	12.693	<a href="#">225793_at</a>	<a href="#">LIX1L</a>	Lix1 homolog (mouse)-like
9.00E-07	5.36E-05	12.676	<a href="#">202766_s_at</a>	<a href="#">FBN1</a>	fibrillin 1
2.00E-06	5.72E-05	12.521	<a href="#">230147_at</a>	<a href="#">F2RL2</a>	coagulation factor II (thrombin) receptor-like 2
7.80E-06	8.44E-05	12.282	<a href="#">203440_at</a>	<a href="#">CDH2</a>	cadherin 2, type 1, N-cadherin (neuronal)
3.30E-06	6.60E-05	12.076	<a href="#">229461_x_at</a>	<a href="#">NEGR1</a>	neuronal growth regulator 1
8.52E-05	0.0003009	11.808	<a href="#">229228_at</a>	<a href="#">CREB5</a>	cAMP responsive element binding protein 5
1.40E-06	5.65E-05	11.572	<a href="#">204472_at</a>	<a href="#">GEM</a>	GTP binding protein overexpressed in skeletal muscle
9.80E-06	9.43E-05	11.556	<a href="#">212077_at</a>	<a href="#">CALD1</a>	caldesmon 1
3.05E-05	0.0001676	11.404	<a href="#">205798_at</a>	<a href="#">IL7R</a>	interleukin 7 receptor
3.60E-06	6.75E-05	11.159	<a href="#">200982_s_at</a>	<a href="#">ANXA6</a>	annexin A6
1.90E-06	5.65E-05	10.776	<a href="#">228335_at</a>	<a href="#">CLDN11</a>	claudin 11
1.20E-06	5.65E-05	10.623	<a href="#">206969_at</a>	<a href="#">KRT34</a>	keratin 34
1.70E-05	0.000122	10.574	<a href="#">229357_at</a>	<a href="#">ADAMTS5</a>	ADAM metalloproteinase with thrombospondin type 1 motif, 5
5.90E-06	7.80E-05	10.247	<a href="#">243818_at</a>	<a href="#">SFTA1P</a>	surfactant associated 1 (pseudogene)
2.00E-06	5.72E-05	10.037	<a href="#">213790_at</a>	<a href="#">ADAM12</a>	ADAM metalloproteinase domain 12
0.0002596	0.0006198	9.684	<a href="#">235518_at</a>	<a href="#">SLC8A1</a>	solute carrier family 8 (sodium/calcium exchanger), member 1
3.59E-05	0.0001784	9.477	<a href="#">235417_at</a>	<a href="#">SPOCD1</a>	SPOC domain containing 1
1.19E-05	0.0001035	9.354	<a href="#">226237_at</a>	<a href="#">NA</a>	NA
0.0001207	0.0003668	9.262	<a href="#">232914_s_at</a>	<a href="#">SYTL2</a>	synaptotagmin-like 2
1.78E-05	0.0001254	9.051	<a href="#">202237_at</a>	<a href="#">NNMT</a>	nicotinamide N-methyltransferase
1.95E-05	0.0001304	8.832	<a href="#">200665_s_at</a>	<a href="#">SPARC</a>	secreted protein, acidic, cysteine-rich (osteonectin)
2.04E-05	0.0001345	8.757	<a href="#">201616_s_at</a>	<a href="#">CALD1</a>	caldesmon 1
4.33E-05	0.0001946	8.719	<a href="#">238919_at</a>	<a href="#">NA</a>	NA
4.00E-07	3.77E-05	8.686	<a href="#">235368_at</a>	<a href="#">ADAMTS5</a>	ADAM metalloproteinase with thrombospondin type 1 motif, 5
9.00E-07	5.36E-05	8.506	<a href="#">221584_s_at</a>	<a href="#">KCNMA1</a>	potassium large conductance calcium-activated channel, subfamily M, alpha member 1
1.20E-05	0.0001036	8.390	<a href="#">232235_at</a>	<a href="#">DSEL</a>	dermatan sulfate epimerase-like
6.00E-07	4.84E-05	8.370	<a href="#">209087_x_at</a>	<a href="#">MCAM</a>	melanoma cell adhesion molecule
4.96E-05	0.000214	8.347	<a href="#">204298_s_at</a>	<a href="#">LOX</a>	lysyl oxidase
0.0001568	0.0004313	8.264	<a href="#">213416_at</a>	<a href="#">ITGA4</a>	integrin, alpha 4 (antigen CD49D, alpha 4 subunit of VLA-4 receptor)
1.16E-05	0.0001021	8.204	<a href="#">203058_s_at</a>	<a href="#">PAPSS2</a>	3-adenosine 5-phosphosulfate synthase 2
1.53E-05	0.0001177	7.969	<a href="#">227061_at</a>	<a href="#">NA</a>	NA
2.68E-05	0.0001554	7.880	<a href="#">212607_at</a>	<a href="#">AKT3</a>	v-akt murine thymoma viral oncogene homolog 3 (protein kinase B, gamma)
0.0002017	0.0005147	7.880	<a href="#">227123_at</a>	<a href="#">RAB3B</a>	RAB3B, member RAS oncogene family
6.10E-06	7.93E-05	7.877	<a href="#">226777_at</a>	<a href="#">NA</a>	NA
1.21E-05	0.0001036	7.766	<a href="#">229004_at</a>	<a href="#">NA</a>	NA
3.90E-06	6.75E-05	7.530	<a href="#">227099_s_at</a>	<a href="#">LOC387763</a>	hypothetical protein LOC387763
0.0001203	0.0003667	7.479	<a href="#">202007_at</a>	<a href="#">NID1</a>	nidogen 1
3.00E-06	6.40E-05	7.403	<a href="#">210869_s_at</a>	<a href="#">MCAM</a>	melanoma cell adhesion molecule
5.45E-05	0.0002265	7.253	<a href="#">215743_at</a>	<a href="#">NMT2</a>	N-myristoyltransferase 2
0.00012	0.0003667	7.142	<a href="#">238447_at</a>	<a href="#">RBMS3</a>	RNA binding motif, single stranded interacting protein
0.0001378	0.0003977	7.131	<a href="#">205525_at</a>	<a href="#">CALD1</a>	caldesmon 1
0.0007644	0.0013652	7.113	<a href="#">226136_at</a>	<a href="#">GLIPR1</a>	GLI pathogenesis-related 1

0.0001117	0.0003517	7.105	<a href="#">213325 at</a>	<a href="#">PVRL3</a>	poliovirus receptor-related 3
0.0001951	0.0005038	7.094	<a href="#">237411 at</a>	<a href="#">ADAMTS6</a>	ADAM metalloproteinase with thrombospondin type 1 motif, 6
4.89E-05	0.0002114	7.033	<a href="#">1555579 s at</a>	<a href="#">PTPRM</a>	protein tyrosine phosphatase, receptor type, M
3.60E-06	6.75E-05	7.003	<a href="#">244503 at</a>	<a href="#">NA</a>	NA
1.51E-05	0.0001177	7.003	<a href="#">209118 s at</a>	<a href="#">TUBA1A</a>	tubulin, alpha 1a
1.16E-05	0.0001021	6.813	<a href="#">227828 s at</a>	<a href="#">FAM176A</a>	family with sequence similarity 176, member A
2.18E-05	0.0001396	6.812	<a href="#">200644 at</a>	<a href="#">MARCKSL1</a>	MARCKS-like 1
0.0001251	0.0003722	6.787	<a href="#">219937 at</a>	<a href="#">TRHDE</a>	thyrotropin-releasing hormone degrading enzyme
1.91E-05	0.0001292	6.763	<a href="#">226751 at</a>	<a href="#">CNRIP1</a>	cannabinoid receptor interacting protein 1
9.05E-05	0.0003099	6.697	<a href="#">223614 at</a>	<a href="#">MMP16</a>	matrix metalloproteinase 16 (membrane-inserted)
1.50E-06	5.65E-05	6.672	<a href="#">205006 s at</a>	<a href="#">NMT2</a>	N-myristoyltransferase 2
5.80E-05	0.0002346	6.662	<a href="#">226834 at</a>	<a href="#">NA</a>	NA
2.60E-06	6.21E-05	6.579	<a href="#">203895 at</a>	<a href="#">PLCB4</a>	phospholipase C, beta 4
5.67E-05	0.0002314	6.551	<a href="#">232458 at</a>	<a href="#">COL3A1</a>	collagen, type III, alpha 1
1.59E-05	0.000119	6.486	<a href="#">218380 at</a>	<a href="#">NLRP1</a>	NLR family, pyrin domain containing 1
0.0001014	0.0003289	6.472	<a href="#">205003 at</a>	<a href="#">DOCK4</a>	dedicator of cytokinesis 4
2.80E-06	6.21E-05	6.462	<a href="#">210517 s at</a>	<a href="#">AKAP12</a>	A kinase (PRKA) anchor protein 12
9.14E-05	0.0003106	6.461	<a href="#">205547 s at</a>	<a href="#">TAGLN</a>	transgelin
2.00E-07	3.57E-05	6.439	<a href="#">227846 at</a>	<a href="#">GPR176</a>	G protein-coupled receptor 176
2.35E-05	0.0001452	6.418	<a href="#">219427 at</a>	<a href="#">FAT4</a>	FAT tumor suppressor homolog 4 (Drosophila)
0.0003268	0.0007258	6.361	<a href="#">210762 s at</a>	<a href="#">DLC1</a>	deleted in liver cancer 1
6.86E-05	0.0002607	6.342	<a href="#">210105 s at</a>	<a href="#">FYN</a>	FYN oncogene related to SRC, FGR, YES
0.0002669	0.0006302	6.320	<a href="#">206382 s at</a>	<a href="#">BDNF</a>	brain-derived neurotrophic factor
6.72E-05	0.0002584	6.301	<a href="#">209031 at</a>	<a href="#">CADM1</a>	cell adhesion molecule 1
4.60E-06	7.04E-05	6.275	<a href="#">228551 at</a>	<a href="#">DENND5B</a>	DENN/MADD domain containing 5B
0.0002058	0.0005234	6.270	<a href="#">219179 at</a>	<a href="#">DACT1</a>	dapper, antagonist of beta-catenin, homolog 1 (Xenopus laevis)
3.98E-05	0.0001887	6.230	<a href="#">205005 s at</a>	<a href="#">NMT2</a>	N-myristoyltransferase 2
1.86E-05	0.0001286	6.204	<a href="#">227503 at</a>	<a href="#">NA</a>	NA
4.48E-05	0.0001978	6.161	<a href="#">225275 at</a>	<a href="#">EDIL3</a>	EGF-like repeats and discoidin I-like domains 3
9.60E-05	0.0003192	6.143	<a href="#">227529 s at</a>	<a href="#">AKAP12</a>	A kinase (PRKA) anchor protein 12
4.43E-05	0.000196	6.123	<a href="#">202952 s at</a>	<a href="#">ADAM12</a>	ADAM metalloproteinase domain 12
1.70E-05	0.000122	6.114	<a href="#">219140 s at</a>	<a href="#">RBP4</a>	retinol binding protein 4, plasma
2.35E-05	0.0001452	5.998	<a href="#">211343 s at</a>	<a href="#">COL13A1</a>	collagen, type XIII, alpha 1
0.0008583	0.0014882	5.954	<a href="#">226142 at</a>	<a href="#">GLIPR1</a>	GLI pathogenesis-related 1
0.0002515	0.0006096	5.886	<a href="#">205083 at</a>	<a href="#">AOX1</a>	aldehyde oxidase 1
6.54E-05	0.0002549	5.833	<a href="#">239367 at</a>	<a href="#">BDNF</a>	brain-derived neurotrophic factor
8.80E-06	9.21E-05	5.832	<a href="#">206924 at</a>	<a href="#">IL11</a>	interleukin 11
0.0007948	0.0014039	5.832	<a href="#">237435 at</a>	<a href="#">NA</a>	NA
0.0001974	0.000506	5.803	<a href="#">205924 at</a>	<a href="#">RAB3B</a>	RAB3B, member RAS oncogene family
6.00E-07	4.84E-05	5.794	<a href="#">203139 at</a>	<a href="#">DAPK1</a>	death-associated protein kinase 1
4.27E-05	0.0001945	5.792	<a href="#">225496 s at</a>	<a href="#">SYTL2</a>	synaptotagmin-like 2
9.67E-05	0.0003206	5.780	<a href="#">212158 at</a>	<a href="#">SDC2</a>	syndecan 2
4.40E-06	6.91E-05	5.764	<a href="#">227530 at</a>	<a href="#">AKAP12</a>	A kinase (PRKA) anchor protein 12

1.89E-05	0.0001291	5.757	<a href="#">219308 s at</a>	<a href="#">AK5</a>	adenylate kinase 5
3.29E-05	0.000171	5.736	<a href="#">210764 s at</a>	<a href="#">CYR61</a>	cysteine-rich, angiogenic inducer, 61
2.45E-05	0.0001487	5.679	<a href="#">229566 at</a>	<a href="#">LOC645638</a>	similar to WDNM1-like protein
0.0002105	0.0005336	5.590	<a href="#">227458 at</a>	NA	NA
8.77E-05	0.0003051	5.567	<a href="#">229641 at</a>	NA	NA
1.61E-05	0.0001197	5.548	<a href="#">235570 at</a>	NA	NA
0.0008001	0.0014089	5.526	<a href="#">229437 at</a>	<a href="#">MIRHG2</a>	microRNA host gene 2 (non-protein coding)
8.00E-07	5.36E-05	5.521	<a href="#">202555 s at</a>	<a href="#">MYLK</a>	myosin light chain kinase
4.64E-05	0.0002025	5.463	<a href="#">210220 at</a>	<a href="#">FZD2</a>	frizzled homolog 2 (Drosophila)
7.60E-06	8.38E-05	5.413	<a href="#">1558280 s at</a>	<a href="#">ARHGAP29</a>	Rho GTPase activating protein 29
1.26E-05	0.0001063	5.386	<a href="#">235696 at</a>	NA	NA
4.31E-05	0.0001945	5.358	<a href="#">236824 at</a>	<a href="#">TMEM132B</a>	transmembrane protein 132B
1.69E-05	0.000122	5.308	<a href="#">204537 s at</a>	<a href="#">GABRE</a>	gamma-aminobutyric acid (GABA) A receptor, epsilon
2.61E-05	0.0001529	5.277	<a href="#">230560 at</a>	<a href="#">STXBP6</a>	syntaxin binding protein 6 (amisyn)
3.19E-05	0.0001701	5.233	<a href="#">224822 at</a>	<a href="#">DLC1</a>	deleted in liver cancer 1
1.10E-05	9.99E-05	5.196	<a href="#">201289 at</a>	<a href="#">CYR61</a>	cysteine-rich, angiogenic inducer, 61
2.10E-06	5.78E-05	5.181	<a href="#">202957 at</a>	<a href="#">HCLS1</a>	hematopoietic cell-specific Lyn substrate 1
0.0002039	0.0005197	5.181	<a href="#">215143 at</a>	<a href="#">DPY19L2P2</a>	dpy-19-like 2 pseudogene 2 (C. elegans)
6.62E-05	0.0002563	5.127	<a href="#">201058 s at</a>	<a href="#">MYL9</a>	myosin, light chain 9, regulatory
0.0002705	0.0006364	5.085	<a href="#">1560573 at</a>	<a href="#">LOC387895</a>	hypothetical gene supported by BC040060
1.36E-05	0.000111	5.061	<a href="#">204955 at</a>	<a href="#">SRPX</a>	sushi-repeat-containing protein, X-linked
2.61E-05	0.0001529	5.035	<a href="#">222162 s at</a>	<a href="#">ADAMTS1</a>	ADAM metalloproteinase with thrombospondin type 1 motif, 1
0.0008332	0.0014547	5.005	<a href="#">226051 at</a>	<a href="#">SELM</a>	selenoprotein M
0.0002277	0.0005651	4.969	<a href="#">229580 at</a>	NA	NA
2.00E-07	3.57E-05	4.917	<a href="#">212256 at</a>	<a href="#">GALNT10</a>	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 10 (GalNAc-T10)
9.65E-05	0.0003204	4.916	<a href="#">205931 s at</a>	<a href="#">CREB5</a>	cAMP responsive element binding protein 5
8.43E-05	0.0002992	4.869	<a href="#">226756 at</a>	NA	NA
6.70E-05	0.0002581	4.848	<a href="#">201312 s at</a>	<a href="#">SH3BGRL</a>	SH3 domain binding glutamic acid-rich protein like
1.03E-05	9.66E-05	4.841	<a href="#">203562 at</a>	<a href="#">FEZ1</a>	fasciculation and elongation protein zeta 1 (zygin I)
0.0002412	0.0005928	4.839	<a href="#">209909 s at</a>	<a href="#">TGFB2</a>	transforming growth factor, beta 2
0.0001323	0.0003881	4.828	<a href="#">216005 at</a>	<a href="#">TNC</a>	tenascin C
2.95E-05	0.0001655	4.827	<a href="#">232268 at</a>	NA	NA
7.50E-06	8.35E-05	4.825	<a href="#">203329 at</a>	<a href="#">PTPRM</a>	protein tyrosine phosphatase, receptor type, M
1.45E-05	0.0001142	4.823	<a href="#">212419 at</a>	<a href="#">ZCCHC24</a>	zinc finger, CCHC domain containing 24
6.70E-06	8.10E-05	4.810	<a href="#">203434 s at</a>	<a href="#">MME</a>	membrane metallo-endopeptidase
0.000378	0.0008052	4.808	<a href="#">241970 at</a>	NA	NA
0.0002811	0.0006545	4.807	<a href="#">232231 at</a>	<a href="#">RUNX2</a>	runt-related transcription factor 2
6.60E-06	8.10E-05	4.802	<a href="#">216033 s at</a>	<a href="#">FYN</a>	FYN oncogene related to SRC, FGR, YES
1.16E-05	0.0001021	4.727	<a href="#">239108 at</a>	<a href="#">FAR2</a>	fatty acyl CoA reductase 2
5.00E-06	7.16E-05	4.723	<a href="#">212488 at</a>	<a href="#">COL5A1</a>	collagen, type V, alpha 1
0.0001628	0.0004398	4.720	<a href="#">225288 at</a>	<a href="#">COL27A1</a>	collagen, type XXVII, alpha 1
1.40E-06	5.65E-05	4.716	<a href="#">201540 at</a>	<a href="#">FHL1</a>	four and a half LIM domains 1
0.0006946	0.0012675	4.701	<a href="#">235342 at</a>	<a href="#">SPOCK3</a>	sparc/osteonectin, cwcv and kazal-like domains proteoglycan (testican) 3

8.00E-05	0.0002894	4.701	<a href="#">209946_at</a>	<a href="#">VEGFC</a>	vascular endothelial growth factor C
0.0005452	0.0010527	4.692	<a href="#">212609_s_at</a>	<a href="#">AKT3</a>	v-akt murine thymoma viral oncogene homolog 3 (protein kinase B, gamma)
0.0004284	0.0008774	4.686	<a href="#">202202_s_at</a>	<a href="#">LAMA4</a>	laminin, alpha 4
0.0002472	0.0006047	4.665	<a href="#">205579_at</a>	<a href="#">HRH1</a>	histamine receptor H1
< 1e-07	< 1e-07	4.646	<a href="#">208782_at</a>	<a href="#">FSTL1</a>	folliculin-like 1
0.0007327	0.001319	4.622	<a href="#">218678_at</a>	<a href="#">NES</a>	nestin
0.0002754	0.0006435	4.612	<a href="#">213429_at</a>	<a href="#">NA</a>	NA
0.0001148	0.000358	4.589	<a href="#">210299_s_at</a>	<a href="#">FHL1</a>	four and a half LIM domains 1
0.0001067	0.0003393	4.569	<a href="#">229307_at</a>	<a href="#">ANKRD28</a>	ankyrin repeat domain 28
2.50E-06	6.21E-05	4.501	<a href="#">205207_at</a>	<a href="#">IL6</a>	interleukin 6 (interferon, beta 2)
2.20E-05	0.0001401	4.498	<a href="#">203570_at</a>	<a href="#">LOXL1</a>	lysyl oxidase-like 1
0.0003879	0.0008195	4.497	<a href="#">1552487_a_at</a>	<a href="#">BNC1</a>	basonuclin 1
3.52E-05	0.0001774	4.486	<a href="#">204083_s_at</a>	<a href="#">TPM2</a>	tropomyosin 2 (beta)
0.0005285	0.001031	4.469	<a href="#">220615_s_at</a>	<a href="#">FAR2</a>	fatty acyl CoA reductase 2
0.0008766	0.001513	4.415	<a href="#">223395_at</a>	<a href="#">ABI3BP</a>	ABI family, member 3 (NESH) binding protein
7.16E-05	0.0002685	4.373	<a href="#">225664_at</a>	<a href="#">COL12A1</a>	collagen, type XII, alpha 1
0.0007751	0.0013794	4.319	<a href="#">235367_at</a>	<a href="#">MYPN</a>	myopalladin
3.11E-05	0.0001676	4.312	<a href="#">212843_at</a>	<a href="#">NCAM1</a>	neural cell adhesion molecule 1
9.46E-05	0.0003168	4.291	<a href="#">1560698_a_at</a>	<a href="#">LOC283392</a>	hypothetical LOC283392
1.92E-05	0.0001292	4.282	<a href="#">203910_at</a>	<a href="#">ARHGAP29</a>	Rho GTPase activating protein 29
3.22E-05	0.0001702	4.272	<a href="#">228438_at</a>	<a href="#">LOC100132891</a>	hypothetical protein LOC100132891
0.000413	0.0008551	4.249	<a href="#">204236_at</a>	<a href="#">FLI1</a>	Friend leukemia virus integration 1
1.64E-05	0.0001203	4.249	<a href="#">220994_s_at</a>	<a href="#">STXBP6</a>	syntaxin binding protein 6 (amisyn)
6.45E-05	0.0002523	4.242	<a href="#">209348_s_at</a>	<a href="#">MAF</a>	v-maf musculoaponeurotic fibrosarcoma oncogene homolog (avian)
0.0001553	0.0004308	4.221	<a href="#">203963_at</a>	<a href="#">CA12</a>	carbonic anhydrase XII
3.90E-06	6.75E-05	4.199	<a href="#">212489_at</a>	<a href="#">COL5A1</a>	collagen, type V, alpha 1
0.0002471	0.0006047	4.187	<a href="#">213603_s_at</a>	<a href="#">RAC2</a>	ras-related C3 botulinum toxin substrate 2 (rho family, small GTP binding protein Rac2)
1.74E-05	0.0001233	4.180	<a href="#">230418_s_at</a>	<a href="#">GALNTL1</a>	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase-like 1
0.0005247	0.0010272	4.176	<a href="#">203980_at</a>	<a href="#">FABP4</a>	fatty acid binding protein 4, adipocyte
0.0002122	0.0005366	4.176	<a href="#">213397_x_at</a>	<a href="#">ANG</a>	angiogenin, ribonuclease, RNase A family, 5
5.01E-05	0.0002149	4.160	<a href="#">219737_s_at</a>	<a href="#">PCDH9</a>	protocadherin 9
8.19E-05	0.0002949	4.159	<a href="#">223723_at</a>	<a href="#">MF12</a>	antigen p97 (melanoma associated) identified by monoclonal antibodies 133.2 and 96.5
3.53E-05	0.0001774	4.148	<a href="#">226932_at</a>	<a href="#">SSPN</a>	sarcospan (Kras oncogene-associated gene)
0.0005252	0.0010272	4.128	<a href="#">225241_at</a>	<a href="#">CCDC80</a>	coiled-coil domain containing 80
3.34E-05	0.0001724	4.122	<a href="#">225524_at</a>	<a href="#">ANTXR2</a>	anthrax toxin receptor 2
0.0001064	0.0003388	4.111	<a href="#">207303_at</a>	<a href="#">PDE1C</a>	phosphodiesterase 1C, calmodulin-dependent 70kDa
0.0002519	0.0006096	4.083	<a href="#">230883_at</a>	<a href="#">NA</a>	NA
5.30E-05	0.0002223	4.079	<a href="#">212423_at</a>	<a href="#">ZCCHC24</a>	zinc finger, CCHC domain containing 24
0.0001031	0.0003321	4.066	<a href="#">1562415_a_at</a>	<a href="#">SPOCD1</a>	SPOC domain containing 1
8.37E-05	0.0002992	4.014	<a href="#">224964_s_at</a>	<a href="#">GNG2</a>	guanine nucleotide binding protein (G protein), gamma 2
1.09E-05	9.98E-05	3.978	<a href="#">205019_s_at</a>	<a href="#">VIPR1</a>	vasoactive intestinal peptide receptor 1
2.61E-05	0.0001529	3.973	<a href="#">219702_at</a>	<a href="#">PLAC1</a>	placenta-specific 1
9.00E-07	5.36E-05	3.964	<a href="#">209357_at</a>	<a href="#">CITED2</a>	Cbp/p300-interacting transactivator, with Glu/Asp-rich carboxy-terminal domain, 2

2.00E-06	5.72E-05	3.964	<a href="#">226066 at</a>	<a href="#">MITF</a>	microphthalmia-associated transcription factor
1.82E-05	0.0001274	3.963	<a href="#">226614 s at</a>	<a href="#">FAM167A</a>	family with sequence similarity 167, member A
0.0002766	0.0006454	3.948	<a href="#">230968 at</a>	NA	NA
4.00E-06	6.75E-05	3.930	<a href="#">218717 s at</a>	<a href="#">LEPREL1</a>	leprecan-like 1
0.0009165	0.0015651	3.917	<a href="#">220407 s at</a>	<a href="#">TGFB2</a>	transforming growth factor, beta 2
9.23E-05	0.0003122	3.912	<a href="#">206363 at</a>	<a href="#">MAF</a>	v-maf musculoaponeurotic fibrosarcoma oncogene homolog (avian)
0.000116	0.0003595	3.910	<a href="#">229273 at</a>	<a href="#">SALL1</a>	sal-like 1 (Drosophila)
6.29E-05	0.0002469	3.902	<a href="#">207876 s at</a>	<a href="#">FLNC</a>	filamin C, gamma (actin binding protein 280)
4.43E-05	0.000196	3.879	<a href="#">230425 at</a>	<a href="#">EPHB1</a>	EPH receptor B1
6.58E-05	0.0002561	3.823	<a href="#">236656 s at</a>	<a href="#">LOC100130506</a>	hypothetical protein LOC100130506
2.00E-05	0.0001322	3.816	<a href="#">202686 s at</a>	<a href="#">AXL</a>	AXL receptor tyrosine kinase
1.86E-05	0.0001286	3.800	<a href="#">1556138 a at</a>	NA	NA
9.48E-05	0.0003168	3.793	<a href="#">1555724 s at</a>	<a href="#">TAGLN</a>	transgelin
0.0008734	0.0015086	3.768	<a href="#">223618 at</a>	<a href="#">FMN2</a>	formin 2
3.55E-05	0.0001776	3.754	<a href="#">230183 at</a>	<a href="#">EXT1</a>	exostoses (multiple) 1
0.0008325	0.0014547	3.749	<a href="#">225381 at</a>	<a href="#">LOC399959</a>	hypothetical gene supported by BX647608
0.0002584	0.0006182	3.744	<a href="#">1556499 s at</a>	<a href="#">COL1A1</a>	collagen, type I, alpha 1
0.0005686	0.001084	3.737	<a href="#">226358 at</a>	NA	NA
0.0001539	0.0004291	3.732	<a href="#">206290 s at</a>	<a href="#">RGS7</a>	regulator of G-protein signaling 7
0.000193	0.0004999	3.729	<a href="#">215867 x at</a>	<a href="#">CA12</a>	carbonic anhydrase XII
3.60E-06	6.75E-05	3.725	<a href="#">201041 s at</a>	<a href="#">DUSP1</a>	dual specificity phosphatase 1
3.90E-06	6.75E-05	3.724	<a href="#">211071 s at</a>	<a href="#">MLLT11</a>	myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 11
2.40E-06	6.21E-05	3.716	<a href="#">226905 at</a>	<a href="#">FAM101B</a>	family with sequence similarity 101, member B
1.19E-05	0.0001035	3.714	<a href="#">218820 at</a>	<a href="#">C14orf132</a>	chromosome 14 open reading frame 132
9.70E-06	9.41E-05	3.707	<a href="#">219295 s at</a>	<a href="#">PCOLCE2</a>	procollagen C-endopeptidase enhancer 2
3.00E-07	3.57E-05	3.707	<a href="#">209949 at</a>	<a href="#">NCF2</a>	neutrophil cytosolic factor 2
6.75E-05	0.0002591	3.702	<a href="#">204748 at</a>	<a href="#">PTGS2</a>	prostaglandin-endoperoxide synthase 2 (prostaglandin G/H synthase and cyclooxygenase)
2.28E-05	0.0001416	3.700	<a href="#">226633 at</a>	<a href="#">RAB8B</a>	RAB8B, member RAS oncogene family
2.96E-05	0.0001657	3.698	<a href="#">228121 at</a>	<a href="#">TGFB2</a>	transforming growth factor, beta 2
0.0008303	0.001453	3.697	<a href="#">202765 s at</a>	<a href="#">FBN1</a>	fibrillin 1
7.51E-05	0.000277	3.696	<a href="#">221031 s at</a>	<a href="#">APOLD1</a>	apolipoprotein L domain containing 1
2.49E-05	0.0001493	3.692	<a href="#">225975 at</a>	<a href="#">PCDH18</a>	protocadherin 18
1.61E-05	0.0001197	3.678	<a href="#">225989 at</a>	<a href="#">HERC4</a>	hect domain and RLD 4
7.50E-06	8.35E-05	3.671	<a href="#">224925 at</a>	<a href="#">PREX1</a>	phosphatidylinositol-3,4,5-trisphosphate-dependent Rac exchange factor 1
4.62E-05	0.000202	3.663	<a href="#">229092 at</a>	<a href="#">NR2F2</a>	nuclear receptor subfamily 2, group F, member 2
0.000909	0.0015558	3.661	<a href="#">203153 at</a>	<a href="#">IFIT1</a>	interferon-induced protein with tetratricopeptide repeats 1
1.38E-05	0.000111	3.652	<a href="#">219304 s at</a>	<a href="#">PDGFD</a>	platelet derived growth factor D
0.000156	0.0004313	3.642	<a href="#">1558692 at</a>	<a href="#">C1orf85</a>	chromosome 1 open reading frame 85
0.0006406	0.0011843	3.623	<a href="#">231067 s at</a>	NA	NA
0.0002714	0.0006379	3.623	<a href="#">221016 s at</a>	<a href="#">TCF7L1</a>	transcription factor 7-like 1 (T-cell specific, HMG-box)
0.0001595	0.0004361	3.606	<a href="#">227646 at</a>	<a href="#">EBF1</a>	early B-cell factor 1
4.80E-06	7.05E-05	3.599	<a href="#">204417 at</a>	<a href="#">GALC</a>	galactosylceramidase
0.0003917	0.0008246	3.585	<a href="#">227140 at</a>	NA	NA

6.60E-05	0.0002563	3.581	<a href="#">203665 at</a>	<a href="#">HMOX1</a>	heme oxygenase (decycling) 1
0.0003573	0.000773	3.566	<a href="#">239227 at</a>	NA	NA
0.0006936	0.0012667	3.553	<a href="#">202936 s at</a>	<a href="#">SOX9</a>	SRY (sex determining region Y)-box 9
1.47E-05	0.000115	3.541	<a href="#">223218 s at</a>	<a href="#">NFKBIZ</a>	nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, zeta
0.0001023	0.0003309	3.538	<a href="#">203423 at</a>	<a href="#">RBP1</a>	retinol binding protein 1, cellular
1.80E-06	5.65E-05	3.533	<a href="#">227386 s at</a>	<a href="#">TMEM200B</a>	transmembrane protein 200B
4.00E-07	3.77E-05	3.501	<a href="#">210095 s at</a>	<a href="#">IGFBP3</a>	insulin-like growth factor binding protein 3
0.0003795	0.0008063	3.501	<a href="#">202310 s at</a>	<a href="#">COL1A1</a>	collagen, type I, alpha 1
0.00061	0.0011432	3.501	<a href="#">219855 at</a>	<a href="#">NUDT11</a>	nudix (nucleoside diphosphate linked moiety X)-type motif 11
3.23E-05	0.0001702	3.454	<a href="#">203325 s at</a>	<a href="#">COL5A1</a>	collagen, type V, alpha 1
7.72E-05	0.0002824	3.454	<a href="#">204529 s at</a>	<a href="#">TOX</a>	thymocyte selection-associated high mobility group box
0.0001847	0.0004847	3.441	<a href="#">206375 s at</a>	<a href="#">HSPB3</a>	heat shock 27kDa protein 3
1.43E-05	0.000113	3.425	<a href="#">202037 s at</a>	<a href="#">SFRP1</a>	secreted frizzled-related protein 1
2.20E-06	5.78E-05	3.417	<a href="#">214612 x at</a>	<a href="#">MAGEA6</a>	melanoma antigen family A, 6
1.80E-06	5.65E-05	3.407	<a href="#">225681 at</a>	<a href="#">CTHRC1</a>	collagen triple helix repeat containing 1
0.0001544	0.0004297	3.402	<a href="#">202859 x at</a>	<a href="#">IL8</a>	interleukin 8
1.77E-05	0.0001251	3.396	<a href="#">206117 at</a>	<a href="#">TPM1</a>	tropomyosin 1 (alpha)
1.42E-05	0.000113	3.394	<a href="#">229019 at</a>	<a href="#">ZNF385B</a>	zinc finger protein 385B
4.20E-06	6.83E-05	3.384	<a href="#">202252 at</a>	<a href="#">RAB13</a>	RAB13, member RAS oncogene family
3.30E-06	6.60E-05	3.381	<a href="#">204984 at</a>	<a href="#">GPC4</a>	glypican 4
0.0001057	0.0003374	3.374	<a href="#">44702 at</a>	<a href="#">SYDE1</a>	synapse defective 1, Rho GTPase, homolog 1 (C. elegans)
7.84E-05	0.0002845	3.372	<a href="#">204963 at</a>	<a href="#">SSPN</a>	sarcospan (Kras oncogene-associated gene)
0.0006748	0.0012361	3.372	<a href="#">204422 s at</a>	<a href="#">FGF2</a>	fibroblast growth factor 2 (basic)
0.0004995	0.0009907	3.368	<a href="#">203821 at</a>	<a href="#">HBEGF</a>	heparin-binding EGF-like growth factor
0.000548	0.0010572	3.357	<a href="#">230130 at</a>	NA	NA
9.30E-06	9.40E-05	3.356	<a href="#">201069 at</a>	<a href="#">MMP2</a>	matrix metalloproteinase 2 (gelatinase A, 72kDa gelatinase, 72kDa type IV collagenase)
8.43E-05	0.0002992	3.351	<a href="#">235548 at</a>	<a href="#">APCDD1L</a>	adenomatosis polyposis coli down-regulated 1-like
0.0002841	0.0006593	3.347	<a href="#">233002 at</a>	<a href="#">PPP4R4</a>	protein phosphatase 4, regulatory subunit 4
8.31E-05	0.0002982	3.334	<a href="#">228007 at</a>	<a href="#">C6orf204</a>	chromosome 6 open reading frame 204
0.0005206	0.0010218	3.329	<a href="#">213712 at</a>	<a href="#">ELOVL2</a>	elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like 2
0.0001325	0.0003881	3.318	<a href="#">236115 at</a>	NA	NA
6.96E-05	0.0002627	3.317	<a href="#">221530 s at</a>	<a href="#">BHLHE41</a>	basic helix-loop-helix family, member e41
0.0001048	0.0003356	3.295	<a href="#">208998 at</a>	<a href="#">UCP2</a>	uncoupling protein 2 (mitochondrial, proton carrier)
0.0001565	0.0004313	3.292	<a href="#">206825 at</a>	<a href="#">OXTR</a>	oxytocin receptor
2.90E-06	6.30E-05	3.289	<a href="#">209942 x at</a>	<a href="#">MAGEA3</a>	melanoma antigen family A, 3
0.0009593	0.0016209	3.281	<a href="#">202158 s at</a>	<a href="#">CUGBP2</a>	CUG triplet repeat, RNA binding protein 2
0.0001201	0.0003667	3.279	<a href="#">209598 at</a>	<a href="#">PNMA2</a>	paraneoplastic antigen MA2
7.16E-05	0.0002685	3.277	<a href="#">203729 at</a>	<a href="#">EMP3</a>	epithelial membrane protein 3
0.000628	0.0011667	3.270	<a href="#">224909 s at</a>	<a href="#">PREX1</a>	phosphatidylinositol-3,4,5-trisphosphate-dependent Rac exchange factor 1
2.36E-05	0.0001452	3.269	<a href="#">228171 s at</a>	<a href="#">PLEKHG4</a>	pleckstrin homology domain containing, family G (with RhoGef domain) member 4
2.60E-06	6.21E-05	3.267	<a href="#">226342 at</a>	<a href="#">SPTBN1</a>	spectrin, beta, non-erythrocytic 1
2.70E-06	6.21E-05	3.259	<a href="#">215388 s at</a>	<a href="#">CFHR1</a>	complement factor H-related 1
4.02E-05	0.0001895	3.256	<a href="#">214930 at</a>	<a href="#">SLITRK5</a>	SLIT and NTRK-like family, member 5



1.38E-05	0.000111	3.200	<a href="#">213438 at</a>	<a href="#">NFASC</a>	neurofascin homolog (chicken)
9.08E-05	0.0003101	3.199	<a href="#">212143 s at</a>	<a href="#">IGFBP3</a>	insulin-like growth factor binding protein 3
3.06E-05	0.0001676	3.193	<a href="#">231175 at</a>	<a href="#">BEND6</a>	BEN domain containing 6
8.80E-05	0.0003056	3.193	<a href="#">201508 at</a>	<a href="#">IGFBP4</a>	insulin-like growth factor binding protein 4
2.44E-05	0.0001487	3.185	<a href="#">232080 at</a>	<a href="#">HECW2</a>	HECT, C2 and WW domain containing E3 ubiquitin protein ligase 2
0.0009599	0.0016209	3.184	<a href="#">207980 s at</a>	<a href="#">CITED2</a>	Cbp/p300-interacting transactivator, with Glu/Asp-rich carboxy-terminal domain, 2
0.0005053	0.0009995	3.182	<a href="#">229065 at</a>	<a href="#">SLC35F3</a>	solute carrier family 35, member F3
0.0003002	0.0006877	3.165	<a href="#">213262 at</a>	<a href="#">SACS</a>	spastic ataxia of Charlevoix-Saguenay (sacsin)
7.29E-05	0.0002715	3.160	<a href="#">229256 at</a>	<a href="#">PGM2L1</a>	phosphoglucomutase 2-like 1
2.80E-06	6.21E-05	3.159	<a href="#">229354 at</a>	<a href="#">AHRR</a>	aryl-hydrocarbon receptor repressor
0.0008507	0.0014793	3.156	<a href="#">239519 at</a>	<a href="#">NA</a>	NA
0.0008184	0.0014355	3.144	<a href="#">202935 s at</a>	<a href="#">SOX9</a>	SRY (sex determining region Y)-box 9
0.0001215	0.0003678	3.144	<a href="#">218720 x at</a>	<a href="#">SEZ6L2</a>	seizure related 6 homolog (mouse)-like 2
7.64E-05	0.0002808	3.132	<a href="#">214913 at</a>	<a href="#">ADAMTS3</a>	ADAM metalloproteinase with thrombospondin type 1 motif, 3
1.00E-06	5.51E-05	3.130	<a href="#">212771 at</a>	<a href="#">FAM171A1</a>	family with sequence similarity 171, member A1
0.0009635	0.0016233	3.124	<a href="#">223622 s at</a>	<a href="#">HYI</a>	hydroxypyruvate isomerase homolog (E. coli)
3.90E-06	6.75E-05	3.112	<a href="#">227279 at</a>	<a href="#">TCEAL3</a>	transcription elongation factor A (SII)-like 3
0.0005303	0.0010327	3.110	<a href="#">226875 at</a>	<a href="#">DOCK11</a>	dedicator of cytokinesis 11
0.0001249	0.0003721	3.107	<a href="#">206243 at</a>	<a href="#">TIMP4</a>	TIMP metalloproteinase inhibitor 4
8.57E-05	0.0003017	3.107	<a href="#">228724 at</a>	<a href="#">NA</a>	NA
3.10E-05	0.0001676	3.100	<a href="#">227713 at</a>	<a href="#">KATNAL1</a>	katanin p60 subunit A-like 1
0.0001845	0.0004847	3.070	<a href="#">229948 at</a>	<a href="#">NA</a>	NA
0.0004648	0.0009308	3.067	<a href="#">205381 at</a>	<a href="#">LRRC17</a>	leucine rich repeat containing 17
6.30E-06	7.96E-05	3.060	<a href="#">205583 s at</a>	<a href="#">ALG13</a>	asparagine-linked glycosylation 13 homolog (S. cerevisiae)
0.000492	0.0009784	3.051	<a href="#">213158 at</a>	<a href="#">NA</a>	NA
0.0003206	0.0007174	3.045	<a href="#">232174 at</a>	<a href="#">NA</a>	NA
0.0001926	0.0004994	3.042	<a href="#">237116 at</a>	<a href="#">LOC646903</a>	hypothetical LOC646903
0.0002163	0.0005446	3.041	<a href="#">215206 at</a>	<a href="#">NA</a>	NA
0.0007379	0.0013262	3.035	<a href="#">209156 s at</a>	<a href="#">COL6A2</a>	collagen, type VI, alpha 2
0.0007052	0.0012812	3.029	<a href="#">236129 at</a>	<a href="#">GALNT5</a>	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 5 (GalNAc-T5)
0.0003029	0.0006911	3.024	<a href="#">206170 at</a>	<a href="#">ADR2</a>	adrenergic, beta-2-, receptor, surface
8.06E-05	0.0002911	3.012	<a href="#">228783 at</a>	<a href="#">BVES</a>	blood vessel epicardial substance
0.000328	0.0007261	3.002	<a href="#">212298 at</a>	<a href="#">NRP1</a>	neuropilin 1
4.29E-05	0.0001945	3.000	<a href="#">225782 at</a>	<a href="#">MSRB3</a>	methionine sulfoxide reductase B3
0.0001588	0.0004347	2.996	<a href="#">218723 s at</a>	<a href="#">C13orf15</a>	chromosome 13 open reading frame 15
1.53E-05	0.0001177	2.994	<a href="#">241353 s at</a>	<a href="#">LOC100129105</a>	similar to hCG1821214
0.0002209	0.000553	2.993	<a href="#">219410 at</a>	<a href="#">TMEM45A</a>	transmembrane protein 45A
2.50E-06	6.21E-05	2.992	<a href="#">219170 at</a>	<a href="#">FSD1</a>	fibronectin type III and SPRY domain containing 1
0.0003508	0.0007634	2.985	<a href="#">1553708 at</a>	<a href="#">MGC16075</a>	hypothetical protein MGC16075
1.70E-06	5.65E-05	2.984	<a href="#">204872 at</a>	<a href="#">TLE4</a>	transducin-like enhancer of split 4 (E(sp1) homolog, Drosophila)
0.000227	0.000564	2.978	<a href="#">231726 at</a>	<a href="#">PCDHB14</a>	protocadherin beta 14
0.00025	0.0006074	2.976	<a href="#">213428 s at</a>	<a href="#">COL6A1</a>	collagen, type VI, alpha 1
0.0004105	0.0008523	2.974	<a href="#">223082 at</a>	<a href="#">SH3KBP1</a>	SH3-domain kinase binding protein 1



0.0001389	0.0003996	2.972	<a href="#">205637 s at</a>	<a href="#">SH3GL3</a>	SH3-domain GRB2-like 3
4.41E-05	0.000196	2.966	<a href="#">227272 at</a>	<a href="#">C15orf52</a>	chromosome 15 open reading frame 52
0.0002518	0.0006096	2.954	<a href="#">225185 at</a>	<a href="#">MRAS</a>	muscle RAS oncogene homolog
2.58E-05	0.0001527	2.946	<a href="#">224002 s at</a>	<a href="#">FKBP7</a>	FK506 binding protein 7
0.0003276	0.0007261	2.946	<a href="#">221870 at</a>	<a href="#">EHD2</a>	EH-domain containing 2
0.0006227	0.0011597	2.940	<a href="#">231766 s at</a>	<a href="#">COL12A1</a>	collagen, type XII, alpha 1
0.0001579	0.0004327	2.940	<a href="#">202796 at</a>	<a href="#">SYNPO</a>	synaptopodin
9.73E-05	0.0003215	2.933	<a href="#">223960 s at</a>	<a href="#">C16orf5</a>	chromosome 16 open reading frame 5
0.0001609	0.0004378	2.928	<a href="#">230706 s at</a>	<a href="#">CAMK2N2</a>	calcium/calmodulin-dependent protein kinase II inhibitor 2
0.0001599	0.0004361	2.922	<a href="#">212190 at</a>	<a href="#">SERPINE2</a>	serpin peptidase inhibitor, clade E (nexin, plasminogen activator inhibitor type 1), member 2
6.10E-06	7.93E-05	2.918	<a href="#">226771 at</a>	<a href="#">ATP8B2</a>	ATPase, class I, type 8B, member 2
0.0005734	0.0010904	2.916	<a href="#">228843 at</a>	NA	NA
0.000287	0.0006649	2.913	<a href="#">235171 at</a>	NA	NA
0.0005142	0.0010127	2.913	<a href="#">226311 at</a>	NA	NA
7.26E-05	0.0002709	2.911	<a href="#">210102 at</a>	<a href="#">VWA5A</a>	von Willebrand factor A domain containing 5A
0.0003256	0.0007239	2.909	<a href="#">204774 at</a>	<a href="#">EVI2A</a>	ecotropic viral integration site 2A
0.0001289	0.0003813	2.908	<a href="#">45297 at</a>	<a href="#">EHD2</a>	EH-domain containing 2
1.30E-06	5.65E-05	2.907	<a href="#">202664 at</a>	<a href="#">WIPF1</a>	WAS/WASL interacting protein family, member 1
0.0001712	0.0004565	2.899	<a href="#">219578 s at</a>	<a href="#">CPEB1</a>	cytoplasmic polyadenylation element binding protein 1
0.0001205	0.0003667	2.886	<a href="#">200632 s at</a>	<a href="#">NDRG1</a>	N-myc downstream regulated 1
0.000133	0.000389	2.885	<a href="#">228573 at</a>	NA	NA
0.0009223	0.0015733	2.877	<a href="#">209604 s at</a>	<a href="#">GATA3</a>	GATA binding protein 3
0.0002884	0.0006661	2.877	<a href="#">202422 s at</a>	<a href="#">ACSL4</a>	acyl-CoA synthetase long-chain family member 4
4.40E-06	6.91E-05	2.862	<a href="#">218651 s at</a>	<a href="#">LARP6</a>	La ribonucleoprotein domain family, member 6
7.82E-05	0.0002843	2.860	<a href="#">218376 s at</a>	<a href="#">MICAL1</a>	microtubule associated monooxygenase, calponin and LIM domain containing 1
4.40E-06	6.91E-05	2.845	<a href="#">201995 at</a>	<a href="#">EXT1</a>	exostoses (multiple) 1
1.55E-05	0.000118	2.845	<a href="#">201482 at</a>	<a href="#">QSOX1</a>	quiescin Q6 sulfhydryl oxidase 1
3.10E-05	0.0001676	2.841	<a href="#">226018 at</a>	<a href="#">C7orf41</a>	chromosome 7 open reading frame 41
0.0003583	0.000773	2.838	<a href="#">225045 at</a>	<a href="#">CCDC88A</a>	coiled-coil domain containing 88A
1.29E-05	0.0001084	2.834	<a href="#">224967 at</a>	<a href="#">UGCG</a>	UDP-glucose ceramide glucosyltransferase
0.0003569	0.000773	2.830	<a href="#">212154 at</a>	<a href="#">SDC2</a>	syndecan 2
6.88E-05	0.0002609	2.830	<a href="#">221276 s at</a>	<a href="#">SYNC</a>	syncoilin, intermediate filament protein
3.30E-06	6.60E-05	2.829	<a href="#">225988 at</a>	<a href="#">HERC4</a>	hect domain and RLD 4
5.17E-05	0.0002193	2.828	<a href="#">227290 at</a>	NA	NA
0.000152	0.0004253	2.825	<a href="#">226876 at</a>	<a href="#">FAM101B</a>	family with sequence similarity 101, member B
0.0009268	0.0015779	2.818	<a href="#">224833 at</a>	<a href="#">ETS1</a>	v-ets erythroblastosis virus E26 oncogene homolog 1 (avian)
0.0002521	0.0006096	2.815	<a href="#">239153 at</a>	<a href="#">HOTAIR</a>	hox transcript antisense RNA (non-protein coding)
9.97E-05	0.0003262	2.810	<a href="#">212414 s at</a>	<a href="#">6-Sep</a>	septin 6
5.63E-05	0.000231	2.809	<a href="#">209883 at</a>	<a href="#">GLT25D2</a>	glycosyltransferase 25 domain containing 2
0.0004256	0.0008732	2.794	<a href="#">225102 at</a>	<a href="#">MGLL</a>	monoglyceride lipase
0.0006108	0.0011432	2.793	<a href="#">201311 s at</a>	<a href="#">SH3BGRL</a>	SH3 domain binding glutamic acid-rich protein like
3.61E-05	0.0001786	2.783	<a href="#">224352 s at</a>	<a href="#">CFL2</a>	cofilin 2 (muscle)
2.22E-05	0.000141	2.781	<a href="#">226152 at</a>	<a href="#">TTC7B</a>	tetratricopeptide repeat domain 7B

7.48E-05	0.0002763	2.777	<a href="#">225575_at</a>	<a href="#">LIFR</a>	leukemia inhibitory factor receptor alpha
0.0001371	0.0003974	2.774	<a href="#">226462_at</a>	<a href="#">STXBP6</a>	syntaxin binding protein 6 (amysyn)
2.55E-05	0.0001517	2.771	<a href="#">219279_at</a>	<a href="#">DOCK10</a>	dedicator of cytokinesis 10
0.0006505	0.0011987	2.771	<a href="#">215069_at</a>	<a href="#">NMT2</a>	N-myristoyltransferase 2
1.37E-05	0.000111	2.771	<a href="#">238575_at</a>	<a href="#">OSBPL6</a>	oxysterol binding protein-like 6
0.0002492	0.0006074	2.769	<a href="#">219747_at</a>	<a href="#">C4orf31</a>	chromosome 4 open reading frame 31
0.000659	0.0012134	2.767	<a href="#">241435_at</a>	<a href="#">NA</a>	NA
0.0002566	0.0006159	2.765	<a href="#">1552972_at</a>	<a href="#">NA</a>	NA
0.0003076	0.0006973	2.765	<a href="#">242100_at</a>	<a href="#">CHSY3</a>	chondroitin sulfate synthase 3
0.0006432	0.0011881	2.763	<a href="#">242876_at</a>	<a href="#">AKT3</a>	v-akt murine thymoma viral oncogene homolog 3 (protein kinase B, gamma)
0.0001644	0.0004409	2.760	<a href="#">212736_at</a>	<a href="#">C16orf45</a>	chromosome 16 open reading frame 45
0.0002693	0.0006343	2.757	<a href="#">227839_at</a>	<a href="#">MBD5</a>	methyl-CpG binding domain protein 5
0.0008023	0.0014117	2.755	<a href="#">203921_at</a>	<a href="#">CHST2</a>	carbohydrate (N-acetylglucosamine-6-O) sulfotransferase 2
1.01E-05	9.64E-05	2.753	<a href="#">212651_at</a>	<a href="#">RHOBTB1</a>	Rho-related BTB domain containing 1
4.09E-05	0.0001895	2.747	<a href="#">219785_s_at</a>	<a href="#">FBXO31</a>	F-box protein 31
0.0008678	0.0015012	2.744	<a href="#">228128_x_at</a>	<a href="#">PAPPA</a>	pregnancy-associated plasma protein A, pappalysin 1
0.0001039	0.0003332	2.741	<a href="#">223315_at</a>	<a href="#">NTN4</a>	netrin 4
8.80E-06	9.21E-05	2.741	<a href="#">224558_s_at</a>	<a href="#">MALAT1</a>	metastasis associated lung adenocarcinoma transcript 1 (non-protein coding)
1.15E-05	0.0001021	2.741	<a href="#">213268_at</a>	<a href="#">CAMTA1</a>	calmodulin binding transcription activator 1
0.0002741	0.0006422	2.736	<a href="#">223393_s_at</a>	<a href="#">TSHZ3</a>	teashirt zinc finger homeobox 3
0.0001161	0.0003595	2.735	<a href="#">209633_at</a>	<a href="#">PPP2R3A</a>	protein phosphatase 2 (formerly 2A), regulatory subunit B@#%&@#%&, alpha
1.59E-05	0.000119	2.730	<a href="#">213029_at</a>	<a href="#">NFIB</a>	nuclear factor I/B
0.0004045	0.0008445	2.719	<a href="#">204602_at</a>	<a href="#">DKK1</a>	dickkopf homolog 1 (Xenopus laevis)
0.0002551	0.0006136	2.713	<a href="#">209708_at</a>	<a href="#">MOXD1</a>	monoxygenase, DBH-like 1
1.89E-05	0.0001291	2.708	<a href="#">225401_at</a>	<a href="#">C1orf85</a>	chromosome 1 open reading frame 85
6.64E-05	0.0002566	2.699	<a href="#">204337_at</a>	<a href="#">RGS4</a>	regulator of G-protein signaling 4
6.08E-05	0.0002416	2.699	<a href="#">206706_at</a>	<a href="#">NTF3</a>	neurotrophin 3
4.30E-06	6.90E-05	2.690	<a href="#">203650_at</a>	<a href="#">PROCR</a>	protein C receptor, endothelial (EPCR)
0.0009615	0.0016211	2.680	<a href="#">229435_at</a>	<a href="#">GLIS3</a>	GLIS family zinc finger 3
0.0005281	0.001031	2.679	<a href="#">222453_at</a>	<a href="#">CYBRD1</a>	cytochrome b reductase 1
1.63E-05	0.00012	2.678	<a href="#">233085_s_at</a>	<a href="#">OBFC2A</a>	oligonucleotide/oligosaccharide-binding fold containing 2A
0.0007062	0.0012815	2.676	<a href="#">204823_at</a>	<a href="#">NAV3</a>	neuron navigator 3
0.0003045	0.0006919	2.672	<a href="#">220987_s_at</a>	<a href="#">NUAK2</a>	NUAK family, SNF1-like kinase, 2
4.19E-05	0.0001929	2.670	<a href="#">204359_at</a>	<a href="#">FLRT2</a>	fibronectin leucine rich transmembrane protein 2
5.20E-05	0.0002202	2.665	<a href="#">233337_s_at</a>	<a href="#">SEZ6L2</a>	seizure related 6 homolog (mouse)-like 2
0.0006752	0.0012361	2.664	<a href="#">203435_s_at</a>	<a href="#">MME</a>	membrane metallo-endopeptidase
0.0002924	0.0006739	2.662	<a href="#">56256_at</a>	<a href="#">SIDT2</a>	SID1 transmembrane family, member 2
7.20E-06	8.26E-05	2.660	<a href="#">203097_s_at</a>	<a href="#">RAPGEF2</a>	Rap guanine nucleotide exchange factor (GEF) 2
5.21E-05	0.0002202	2.658	<a href="#">214164_x_at</a>	<a href="#">CA12</a>	carbonic anhydrase XII
0.0003347	0.0007358	2.657	<a href="#">218309_at</a>	<a href="#">CAMK2N1</a>	calcium/calmodulin-dependent protein kinase II inhibitor 1
0.0003588	0.0007734	2.654	<a href="#">205880_at</a>	<a href="#">PRKD1</a>	protein kinase D1
0.000702	0.0012769	2.651	<a href="#">237183_at</a>	<a href="#">GALNT5</a>	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 5 (GalNAc-T5)
0.0009284	0.0015795	2.637	<a href="#">211026_s_at</a>	<a href="#">MGLL</a>	monoglyceride lipase

4.17E-05	0.0001924	2.635	<a href="#">219501_at</a>	<a href="#">ENOX1</a>	ecto-NOX disulfide-thiol exchanger 1
0.000185	0.0004847	2.633	<a href="#">205832_at</a>	<a href="#">CPA4</a>	carboxypeptidase A4
6.80E-05	0.0002597	2.633	<a href="#">212813_at</a>	<a href="#">JAM3</a>	junctional adhesion molecule 3
0.0003696	0.0007918	2.632	<a href="#">223769_x_at</a>	<a href="#">HYI</a>	hydroxypyruvate isomerase homolog (E. coli)
5.90E-06	7.80E-05	2.627	<a href="#">227396_at</a>	<a href="#">PTPRJ</a>	protein tyrosine phosphatase, receptor type, J
0.0009455	0.0016058	2.620	<a href="#">224940_s_at</a>	<a href="#">PAPPA</a>	pregnancy-associated plasma protein A, pappalysin 1
0.0006375	0.0011799	2.619	<a href="#">243041_s_at</a>	<a href="#">RBM53</a>	RNA binding motif, single stranded interacting protein
2.68E-05	0.0001554	2.615	<a href="#">201849_at</a>	<a href="#">BNIP3</a>	BCL2/adenovirus E1B 19kDa interacting protein 3
0.0002581	0.0006182	2.609	<a href="#">205498_at</a>	<a href="#">GHR</a>	growth hormone receptor
1.87E-05	0.0001289	2.600	<a href="#">222872_x_at</a>	<a href="#">OBFC2A</a>	oligonucleotide/oligosaccharide-binding fold containing 2A
0.0002651	0.0006296	2.600	<a href="#">209632_at</a>	<a href="#">PPP2R3A</a>	protein phosphatase 2 (formerly 2A), regulatory subunit B@#%&@#%&, alpha
0.0003459	0.0007542	2.594	<a href="#">202552_s_at</a>	<a href="#">CRIM1</a>	cysteine rich transmembrane BMP regulator 1 (chordin-like)
0.0001848	0.0004847	2.591	<a href="#">221760_at</a>	<a href="#">MAN1A1</a>	mannosidase, alpha, class 1A, member 1
5.10E-06	7.25E-05	2.580	<a href="#">207233_s_at</a>	<a href="#">MITF</a>	microphthalmia-associated transcription factor
5.10E-05	0.0002176	2.578	<a href="#">201105_at</a>	<a href="#">LGALS1</a>	lectin, galactoside-binding, soluble, 1
0.0003563	0.000773	2.572	<a href="#">228890_at</a>	<a href="#">ATOH8</a>	atonal homolog 8 (Drosophila)
0.0001635	0.0004398	2.572	<a href="#">239202_at</a>	<a href="#">NA</a>	NA
0.0002873	0.0006649	2.570	<a href="#">209894_at</a>	<a href="#">LEPR</a>	leptin receptor
0.0007971	0.0014058	2.569	<a href="#">227966_s_at</a>	<a href="#">CCDC74B</a>	coiled-coil domain containing 74B
2.24E-05	0.0001414	2.568	<a href="#">228234_at</a>	<a href="#">TICAM2</a>	toll-like receptor adaptor molecule 2
1.54E-05	0.000118	2.565	<a href="#">216869_at</a>	<a href="#">PDE1C</a>	phosphodiesterase 1C, calmodulin-dependent 70kDa
0.0001599	0.0004361	2.563	<a href="#">209651_at</a>	<a href="#">TGFB11</a>	transforming growth factor beta 1 induced transcript 1
3.55E-05	0.0001776	2.551	<a href="#">225871_at</a>	<a href="#">STEAP2</a>	six transmembrane epithelial antigen of the prostate 2
5.62E-05	0.000231	2.550	<a href="#">203282_at</a>	<a href="#">GBE1</a>	glucan (1,4-alpha-), branching enzyme 1
0.0003026	0.0006911	2.534	<a href="#">1558693_s_at</a>	<a href="#">C1orf85</a>	chromosome 1 open reading frame 85
0.0002066	0.0005243	2.532	<a href="#">237310_at</a>	<a href="#">NA</a>	NA
1.30E-05	0.0001089	2.528	<a href="#">209210_s_at</a>	<a href="#">FERMT2</a>	fermitin family homolog 2 (Drosophila)
0.0001751	0.0004647	2.524	<a href="#">205479_s_at</a>	<a href="#">PLAU</a>	plasminogen activator, urokinase
0.0002211	0.000553	2.519	<a href="#">218330_s_at</a>	<a href="#">NAV2</a>	neuron navigator 2
0.0001167	0.00036	2.511	<a href="#">229784_at</a>	<a href="#">MGC16121</a>	hypothetical protein MGC16121
2.78E-05	0.0001591	2.505	<a href="#">212646_at</a>	<a href="#">RFTN1</a>	raftlin, lipid raft linker 1
0.0001884	0.0004925	2.503	<a href="#">204099_at</a>	<a href="#">SMARCD3</a>	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily d, member 3
0.000101	0.0003288	2.501	<a href="#">223805_at</a>	<a href="#">OSBPL6</a>	oxysterol binding protein-like 6
1.68E-05	0.0001217	2.493	<a href="#">200808_s_at</a>	<a href="#">ZYX</a>	zyxin
2.12E-05	0.0001377	2.491	<a href="#">213035_at</a>	<a href="#">ANKRD28</a>	ankyrin repeat domain 28
0.0001972	0.000506	2.484	<a href="#">225442_at</a>	<a href="#">DDR2</a>	discoidin domain receptor tyrosine kinase 2
9.30E-06	9.40E-05	2.483	<a href="#">212012_at</a>	<a href="#">PXDN</a>	peroxidasin homolog (Drosophila)
8.49E-05	0.0003004	2.476	<a href="#">203882_at</a>	<a href="#">IRF9</a>	interferon regulatory factor 9
6.00E-06	7.89E-05	2.475	<a href="#">209365_s_at</a>	<a href="#">ECM1</a>	extracellular matrix protein 1
3.80E-06	6.75E-05	2.472	<a href="#">226609_at</a>	<a href="#">DCBLD1</a>	discoidin, CUB and LCCL domain containing 1
4.07E-05	0.0001895	2.467	<a href="#">227533_at</a>	<a href="#">NA</a>	NA
7.30E-06	8.32E-05	2.465	<a href="#">209289_at</a>	<a href="#">NFIB</a>	nuclear factor I/B
1.46E-05	0.0001146	2.465	<a href="#">209351_at</a>	<a href="#">KRT14</a>	keratin 14

4.70E-06	7.04E-05	2.464	<a href="#">225798 at</a>	<a href="#">JAZF1</a>	JAZF zinc finger 1
7.40E-06	8.32E-05	2.460	<a href="#">203438 at</a>	<a href="#">STC2</a>	stanniocalcin 2
1.34E-05	0.000111	2.457	<a href="#">228708 at</a>	<a href="#">RAB27B</a>	RAB27B, member RAS oncogene family
0.0003879	0.0008195	2.456	<a href="#">243366 s at</a>	<a href="#">NA</a>	NA
1.09E-05	9.98E-05	2.455	<a href="#">215706 x at</a>	<a href="#">ZYX</a>	zyxin
0.0003241	0.000722	2.452	<a href="#">214633 at</a>	<a href="#">SOX3</a>	SRY (sex determining region Y)-box 3
0.0001226	0.0003687	2.444	<a href="#">235308 at</a>	<a href="#">ZBTB20</a>	zinc finger and BTB domain containing 20
2.26E-05	0.0001414	2.435	<a href="#">203408 s at</a>	<a href="#">SATB1</a>	SATB homeobox 1
6.14E-05	0.0002423	2.434	<a href="#">226773 at</a>	<a href="#">NA</a>	NA
3.23E-05	0.0001702	2.434	<a href="#">211538 s at</a>	<a href="#">HSPA2</a>	heat shock 70kDa protein 2
0.0007933	0.0014024	2.432	<a href="#">227166 at</a>	<a href="#">DNAJC18</a>	DnaJ (Hsp40) homolog, subfamily C, member 18
0.0006161	0.0011503	2.430	<a href="#">232752 at</a>	<a href="#">NA</a>	NA
0.0001538	0.0004291	2.429	<a href="#">238934 at</a>	<a href="#">NA</a>	NA
1.08E-05	9.98E-05	2.425	<a href="#">223170 at</a>	<a href="#">TMEM98</a>	transmembrane protein 98
0.0003975	0.0008337	2.423	<a href="#">230240 at</a>	<a href="#">NA</a>	NA
2.99E-05	0.0001665	2.422	<a href="#">209290 s at</a>	<a href="#">NFIB</a>	nuclear factor I/B
0.000265	0.0006296	2.421	<a href="#">228555 at</a>	<a href="#">CAMK2D</a>	calcium/calmodulin-dependent protein kinase II delta
6.40E-06	8.04E-05	2.420	<a href="#">205330 at</a>	<a href="#">MN1</a>	meningioma (disrupted in balanced translocation) 1
4.80E-06	7.05E-05	2.419	<a href="#">242775 at</a>	<a href="#">NA</a>	NA
5.96E-05	0.0002381	2.419	<a href="#">209522 s at</a>	<a href="#">CRAT</a>	carnitine acetyltransferase
5.90E-05	0.0002369	2.416	<a href="#">225710 at</a>	<a href="#">GNB4</a>	guanine nucleotide binding protein (G protein), beta polypeptide 4
0.0001623	0.0004398	2.408	<a href="#">225897 at</a>	<a href="#">MARCKS</a>	myristoylated alanine-rich protein kinase C substrate
0.0001084	0.0003428	2.407	<a href="#">219882 at</a>	<a href="#">TTL7</a>	tubulin tyrosine ligase-like family, member 7
0.0001433	0.000407	2.407	<a href="#">213800 at</a>	<a href="#">CFH</a>	complement factor H
0.0003782	0.0008052	2.407	<a href="#">217892 s at</a>	<a href="#">LIMA1</a>	LIM domain and actin binding 1
2.20E-06	5.78E-05	2.407	<a href="#">209732 at</a>	<a href="#">CLEC2B</a>	C-type lectin domain family 2, member B
0.0004195	0.000867	2.405	<a href="#">1569150 x at</a>	<a href="#">PDLIM7</a>	PDZ and LIM domain 7 (enigma)
0.0008805	0.0015159	2.403	<a href="#">209030 s at</a>	<a href="#">CADM1</a>	cell adhesion molecule 1
0.0001956	0.0005043	2.397	<a href="#">227771 at</a>	<a href="#">LIFR</a>	leukemia inhibitory factor receptor alpha
5.44E-05	0.0002265	2.396	<a href="#">238917 s at</a>	<a href="#">DENND5B</a>	DENN/MADD domain containing 5B
8.53E-05	0.0003009	2.392	<a href="#">203184 at</a>	<a href="#">FBN2</a>	fibrillin 2
0.0005346	0.0010384	2.391	<a href="#">219346 at</a>	<a href="#">LRFN3</a>	leucine rich repeat and fibronectin type III domain containing 3
2.06E-05	0.0001354	2.387	<a href="#">210143 at</a>	<a href="#">ANXA10</a>	annexin A10
6.79E-05	0.0002597	2.387	<a href="#">1553997 a at</a>	<a href="#">ASPHD1</a>	aspartate beta-hydroxylase domain containing 1
0.0001547	0.0004297	2.384	<a href="#">213103 at</a>	<a href="#">STARD13</a>	StAR-related lipid transfer (START) domain containing 13
0.0004617	0.0009271	2.382	<a href="#">224819 at</a>	<a href="#">TCEAL8</a>	transcription elongation factor A (SII)-like 8
0.0006015	0.0011333	2.381	<a href="#">204881 s at</a>	<a href="#">UGCG</a>	UDP-glucose ceramide glucosyltransferase
0.0003161	0.0007097	2.378	<a href="#">210510 s at</a>	<a href="#">NRP1</a>	neuropilin 1
2.90E-06	6.30E-05	2.378	<a href="#">227115 at</a>	<a href="#">NA</a>	NA
1.00E-06	5.51E-05	2.375	<a href="#">221036 s at</a>	<a href="#">APH1B</a>	anterior pharynx defective 1 homolog B (C. elegans)
0.000173	0.0004607	2.375	<a href="#">208510 s at</a>	<a href="#">PPARG</a>	peroxisome proliferator-activated receptor gamma
3.15E-05	0.0001684	2.370	<a href="#">212345 s at</a>	<a href="#">CREB3L2</a>	cAMP responsive element binding protein 3-like 2
4.20E-06	6.83E-05	2.368	<a href="#">213093 at</a>	<a href="#">PRKCA</a>	protein kinase C, alpha

9.00E-07	5.36E-05	2.366	<a href="#">226989 at</a>	<a href="#">RGMB</a>	RGM domain family, member B
0.0007224	0.0013046	2.364	<a href="#">215790 at</a>	<a href="#">AJAP1</a>	adherens junctions associated protein 1
0.0004316	0.0008831	2.357	<a href="#">226058 at</a>	<a href="#">B3GNT9</a>	UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 9
7.70E-05	0.0002822	2.353	<a href="#">213227 at</a>	<a href="#">PGRMC2</a>	progesterone receptor membrane component 2
0.0001823	0.0004815	2.351	<a href="#">213418 at</a>	<a href="#">HSPA6</a>	heat shock 70kDa protein 6 (HSP70B@#%&)
0.0001962	0.0005049	2.345	<a href="#">231406 at</a>	<a href="#">ORAI2</a>	ORAI calcium release-activated calcium modulator 2
0.0003557	0.0007726	2.343	<a href="#">209119 x at</a>	<a href="#">NR2F2</a>	nuclear receptor subfamily 2, group F, member 2
0.0001444	0.0004084	2.332	<a href="#">213906 at</a>	<a href="#">MYBL1</a>	v-myb myeloblastosis viral oncogene homolog (avian)-like 1
0.0003477	0.0007574	2.332	<a href="#">213122 at</a>	<a href="#">TSPYL5</a>	TSPY-like 5
0.0002411	0.0005928	2.331	<a href="#">212013 at</a>	<a href="#">PXDN</a>	peroxidasin homolog (Drosophila)
2.85E-05	0.0001619	2.330	<a href="#">236029 at</a>	<a href="#">FAT3</a>	FAT tumor suppressor homolog 3 (Drosophila)
8.64E-05	0.0003024	2.328	<a href="#">235911 at</a>	<a href="#">LOC440995</a>	hypothetical gene supported by BC034933; BC068085
0.0001228	0.0003687	2.326	<a href="#">227370 at</a>	<a href="#">FAM171B</a>	family with sequence similarity 171, member B
9.11E-05	0.0003106	2.321	<a href="#">228046 at</a>	<a href="#">ZNF827</a>	zinc finger protein 827
6.03E-05	0.0002405	2.319	<a href="#">217989 at</a>	<a href="#">HSD17B11</a>	hydroxysteroid (17-beta) dehydrogenase 11
0.0006337	0.0011754	2.313	<a href="#">206460 at</a>	<a href="#">AJAP1</a>	adherens junctions associated protein 1
0.0004079	0.0008484	2.312	<a href="#">202073 at</a>	<a href="#">OPTN</a>	optineurin
0.0003128	0.0007037	2.309	<a href="#">215613 at</a>	<a href="#">ADAM12</a>	ADAM metalloproteinase domain 12
1.92E-05	0.0001292	2.307	<a href="#">205120 s at</a>	<a href="#">SGCB</a>	sarcoglycan, beta (43kDa dystrophin-associated glycoprotein)
2.19E-05	0.0001399	2.307	<a href="#">230463 at</a>	<a href="#">NA</a>	NA
0.0002593	0.0006197	2.306	<a href="#">224911 s at</a>	<a href="#">DCBLD2</a>	discoidin, CUB and LCCL domain containing 2
0.0006183	0.0011534	2.304	<a href="#">222288 at</a>	<a href="#">NA</a>	NA
0.0002559	0.0006149	2.304	<a href="#">231725 at</a>	<a href="#">PCDHB2</a>	protocadherin beta 2
2.82E-05	0.0001606	2.302	<a href="#">205383 s at</a>	<a href="#">ZBTB20</a>	zinc finger and BTB domain containing 20
0.0003117	0.0007028	2.296	<a href="#">204966 at</a>	<a href="#">BAI2</a>	brain-specific angiogenesis inhibitor 2
9.02E-05	0.0003095	2.295	<a href="#">220326 s at</a>	<a href="#">FLJ10357</a>	hypothetical protein FLJ10357
5.57E-05	0.0002302	2.291	<a href="#">212774 at</a>	<a href="#">ZNF238</a>	zinc finger protein 238
0.0001963	0.0005049	2.291	<a href="#">204303 s at</a>	<a href="#">KIAA0427</a>	KIAA0427
0.0007306	0.0013162	2.282	<a href="#">204421 s at</a>	<a href="#">FGF2</a>	fibroblast growth factor 2 (basic)
0.0004531	0.0009131	2.281	<a href="#">203136 at</a>	<a href="#">RABAC1</a>	Rab acceptor 1 (prenylated)
5.46E-05	0.0002265	2.280	<a href="#">203390 s at</a>	<a href="#">KIF3C</a>	kinesin family member 3C
8.30E-06	8.89E-05	2.279	<a href="#">212829 at</a>	<a href="#">PIP4K2A</a>	phosphatidylinositol-5-phosphate 4-kinase, type II, alpha
7.40E-06	8.32E-05	2.279	<a href="#">201389 at</a>	<a href="#">ITGA5</a>	integrin, alpha 5 (fibronectin receptor, alpha polypeptide)
0.000101	0.0003288	2.278	<a href="#">229521 at</a>	<a href="#">FLJ36031</a>	hypothetical protein FLJ36031
5.00E-06	7.16E-05	2.277	<a href="#">221059 s at</a>	<a href="#">CHST6</a>	carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 6
0.0001447	0.0004084	2.274	<a href="#">1561064 a at</a>	<a href="#">NA</a>	NA
0.0001355	0.0003948	2.270	<a href="#">220559 at</a>	<a href="#">EN1</a>	engrailed homeobox 1
8.61E-05	0.0003022	2.268	<a href="#">230099 at</a>	<a href="#">NA</a>	NA
1.91E-05	0.0001292	2.266	<a href="#">227489 at</a>	<a href="#">SMURF2</a>	SMAD specific E3 ubiquitin protein ligase 2
2.54E-05	0.0001515	2.264	<a href="#">212120 at</a>	<a href="#">RHOQ</a>	ras homolog gene family, member Q
0.0005605	0.0010733	2.263	<a href="#">221024 s at</a>	<a href="#">SLC2A10</a>	solute carrier family 2 (facilitated glucose transporter), member 10
0.0005379	0.0010423	2.259	<a href="#">213644 at</a>	<a href="#">CCDC46</a>	coiled-coil domain containing 46
5.90E-06	7.80E-05	2.257	<a href="#">224583 at</a>	<a href="#">COTL1</a>	coactosin-like 1 (Dictyostelium)

0.00018	0.000476	2.253	<a href="#">218627_at</a>	<a href="#">DRAM</a>	damage-regulated autophagy modulator
5.40E-06	7.54E-05	2.252	<a href="#">227326_at</a>	<a href="#">MXRA7</a>	matrix-remodelling associated 7
0.0007809	0.001387	2.252	<a href="#">221953_s_at</a>	<a href="#">MT1P3</a>	metallothionein 1 pseudogene 3
9.30E-06	9.40E-05	2.248	<a href="#">210987_x_at</a>	<a href="#">TPM1</a>	tropomyosin 1 (alpha)
0.000436	0.0008897	2.247	<a href="#">225163_at</a>	<a href="#">FRMD4A</a>	FERM domain containing 4A
2.20E-06	5.78E-05	2.247	<a href="#">228739_at</a>	<a href="#">LOC100131506</a>	hypothetical LOC100131506
0.0001492	0.0004185	2.245	<a href="#">202551_s_at</a>	<a href="#">CRIM1</a>	cysteine rich transmembrane BMP regulator 1 (chordin-like)
8.08E-05	0.0002914	2.239	<a href="#">201162_at</a>	<a href="#">IGFBP7</a>	insulin-like growth factor binding protein 7
0.0002012	0.000514	2.236	<a href="#">203813_s_at</a>	<a href="#">SLIT3</a>	slit homolog 3 (Drosophila)
0.0005447	0.0010526	2.235	<a href="#">209340_at</a>	<a href="#">UAP1</a>	UDP-N-acetylglucosamine pyrophosphorylase 1
6.85E-05	0.0002607	2.226	<a href="#">204589_at</a>	<a href="#">NUAK1</a>	NUAK family, SNF1-like kinase, 1
0.0004431	0.0009018	2.226	<a href="#">1569149_at</a>	<a href="#">PDLIM7</a>	PDZ and LIM domain 7 (enigma)
2.71E-05	0.0001559	2.224	<a href="#">53991_at</a>	<a href="#">DENND2A</a>	DENN/MADD domain containing 2A
0.0005526	0.0010633	2.224	<a href="#">1555471_a_at</a>	<a href="#">FMN2</a>	formin 2
0.000325	0.0007233	2.217	<a href="#">227261_at</a>	<a href="#">KLF12</a>	Kruppel-like factor 12
0.0001128	0.0003547	2.217	<a href="#">207719_x_at</a>	<a href="#">CEP170</a>	centrosomal protein 170kDa
0.0003965	0.0008324	2.216	<a href="#">205407_at</a>	<a href="#">RECK</a>	reversion-inducing-cysteine-rich protein with kazal motifs
0.0001446	0.0004084	2.210	<a href="#">239638_at</a>	<a href="#">NA</a>	NA
0.0003865	0.0008182	2.208	<a href="#">243864_at</a>	<a href="#">CCDC80</a>	coiled-coil domain containing 80
0.0003688	0.0007911	2.207	<a href="#">241058_at</a>	<a href="#">NA</a>	NA
0.0004088	0.0008495	2.206	<a href="#">226694_at</a>	<a href="#">AKAP2</a>	A kinase (PRKA) anchor protein 2
0.0001422	0.0004054	2.204	<a href="#">36553_at</a>	<a href="#">ASMTL</a>	acetylserotonin O-methyltransferase-like
0.0004528	0.0009131	2.202	<a href="#">237301_at</a>	<a href="#">NA</a>	NA
1.33E-05	0.0001106	2.202	<a href="#">226197_at</a>	<a href="#">NA</a>	NA
3.65E-05	0.000179	2.201	<a href="#">232067_at</a>	<a href="#">C6orf168</a>	chromosome 6 open reading frame 168
0.0001903	0.0004963	2.200	<a href="#">203874_s_at</a>	<a href="#">SMARCA1</a>	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 1
0.0004505	0.0009103	2.199	<a href="#">224229_s_at</a>	<a href="#">AKT3</a>	v-akt murine thymoma viral oncogene homolog 3 (protein kinase B, gamma)
0.0004005	0.0008387	2.199	<a href="#">243606_at</a>	<a href="#">NA</a>	NA
1.80E-05	0.0001264	2.198	<a href="#">226507_at</a>	<a href="#">PAK1</a>	p21 protein (Cdc42/Rac)-activated kinase 1
0.0002658	0.0006302	2.196	<a href="#">212561_at</a>	<a href="#">DENND5A</a>	DENN/MADD domain containing 5A
2.62E-05	0.0001531	2.194	<a href="#">235343_at</a>	<a href="#">VASH2</a>	vasohibin 2
9.14E-05	0.0003106	2.186	<a href="#">227224_at</a>	<a href="#">RALGPS2</a>	Ral GEF with PH domain and SH3 binding motif 2
0.0005152	0.0010129	2.185	<a href="#">205596_s_at</a>	<a href="#">SMURF2</a>	SMAD specific E3 ubiquitin protein ligase 2
0.0001429	0.0004064	2.185	<a href="#">202440_s_at</a>	<a href="#">ST5</a>	suppression of tumorigenicity 5
0.000297	0.0006817	2.184	<a href="#">213241_at</a>	<a href="#">PLXNC1</a>	plexin C1
1.74E-05	0.0001233	2.181	<a href="#">210135_s_at</a>	<a href="#">SHOX2</a>	short stature homeobox 2
0.0002474	0.0006047	2.179	<a href="#">204604_at</a>	<a href="#">PFTK1</a>	PFTAIRE protein kinase 1
4.37E-05	0.000196	2.178	<a href="#">218656_s_at</a>	<a href="#">LHFP</a>	lipoma HMGIC fusion partner
1.63E-05	0.00012	2.176	<a href="#">226546_at</a>	<a href="#">NA</a>	NA
2.27E-05	0.0001414	2.171	<a href="#">219895_at</a>	<a href="#">FAM70A</a>	family with sequence similarity 70, member A
0.0001226	0.0003687	2.169	<a href="#">205741_s_at</a>	<a href="#">DTNA</a>	dystrobrevin, alpha
0.0002722	0.0006391	2.168	<a href="#">203370_s_at</a>	<a href="#">PDLIM7</a>	PDZ and LIM domain 7 (enigma)
0.0004518	0.0009121	2.167	<a href="#">235061_at</a>	<a href="#">PPM1K</a>	protein phosphatase 1K (PP2C domain containing)

0.0005238	0.0010272	2.166	<a href="#">222900_at</a>	<a href="#">NA</a>	NA
0.0003396	0.0007426	2.166	<a href="#">238532_at</a>	<a href="#">DPF3</a>	D4, zinc and double PHD fingers, family 3
3.60E-06	6.75E-05	2.163	<a href="#">235205_at</a>	<a href="#">LOC100128259</a>	hypothetical protein LOC100128259
0.0004627	0.0009283	2.162	<a href="#">226853_at</a>	<a href="#">BMP2K</a>	BMP2 inducible kinase
1.19E-05	0.0001035	2.160	<a href="#">210139_s_at</a>	<a href="#">PMP22</a>	peripheral myelin protein 22
0.0001827	0.0004817	2.160	<a href="#">227088_at</a>	<a href="#">PDE5A</a>	phosphodiesterase 5A, cGMP-specific
0.0007303	0.0013162	2.160	<a href="#">225817_at</a>	<a href="#">CGNL1</a>	cingulin-like 1
0.0003927	0.0008259	2.156	<a href="#">201701_s_at</a>	<a href="#">PGRMC2</a>	progesterone receptor membrane component 2
0.0008437	0.0014696	2.152	<a href="#">221261_x_at</a>	<a href="#">MAGED4B</a>	melanoma antigen family D, 4B
9.00E-06	9.33E-05	2.150	<a href="#">217889_s_at</a>	<a href="#">CYBRD1</a>	cytochrome b reductase 1
8.74E-05	0.0003045	2.145	<a href="#">213484_at</a>	<a href="#">NA</a>	NA
0.0005973	0.0011264	2.143	<a href="#">227226_at</a>	<a href="#">MRAP2</a>	melanocortin 2 receptor accessory protein 2
6.25E-05	0.0002462	2.141	<a href="#">241484_x_at</a>	<a href="#">NA</a>	NA
0.0002363	0.0005833	2.139	<a href="#">226955_at</a>	<a href="#">AFAP1L1</a>	actin filament associated protein 1-like 1
1.53E-05	0.0001177	2.139	<a href="#">208370_s_at</a>	<a href="#">RCAN1</a>	regulator of calcineurin 1
6.20E-06	7.96E-05	2.139	<a href="#">224663_s_at</a>	<a href="#">CFL2</a>	cofilin 2 (muscle)
3.60E-05	0.0001785	2.139	<a href="#">224791_at</a>	<a href="#">ASAP1</a>	ArfGAP with SH3 domain, ankyrin repeat and PH domain 1
0.0005427	0.0010497	2.138	<a href="#">204368_at</a>	<a href="#">SLCO2A1</a>	solute carrier organic anion transporter family, member 2A1
0.0003078	0.0006973	2.135	<a href="#">239162_at</a>	<a href="#">NA</a>	NA
0.000584	0.0011077	2.134	<a href="#">210319_x_at</a>	<a href="#">MSX2</a>	msh homeobox 2
9.74E-05	0.0003215	2.129	<a href="#">201719_s_at</a>	<a href="#">EPB41L2</a>	erythrocyte membrane protein band 4.1-like 2
0.0005096	0.0010063	2.128	<a href="#">230722_at</a>	<a href="#">BNC2</a>	basonuclin 2
0.0005868	0.0011112	2.127	<a href="#">1555976_s_at</a>	<a href="#">MYL12A</a>	myosin, light chain 12A, regulatory, non-sarcomeric
0.0001952	0.0005038	2.123	<a href="#">213213_at</a>	<a href="#">DIDO1</a>	death inducer-obliterator 1
0.0007629	0.0013636	2.121	<a href="#">237452_at</a>	<a href="#">NA</a>	NA
0.0009117	0.0015593	2.119	<a href="#">239431_at</a>	<a href="#">TICAM2</a>	toll-like receptor adaptor molecule 2
0.0001011	0.0003288	2.118	<a href="#">227929_at</a>	<a href="#">NA</a>	NA
0.0003086	0.0006984	2.118	<a href="#">231411_at</a>	<a href="#">LHFP</a>	lipoma HMGIC fusion partner
0.0001641	0.0004407	2.117	<a href="#">225860_at</a>	<a href="#">MT1P3</a>	metallothionein 1 pseudogene 3
5.13E-05	0.000218	2.117	<a href="#">202074_s_at</a>	<a href="#">OPTN</a>	optineurin
7.00E-04	0.0012743	2.114	<a href="#">221435_x_at</a>	<a href="#">HYI</a>	hydroxypyruvate isomerase homolog (E. coli)
0.0001399	0.0004014	2.111	<a href="#">224927_at</a>	<a href="#">KIAA1949</a>	KIAA1949
4.29E-05	0.0001945	2.109	<a href="#">240312_at</a>	<a href="#">LOC389895</a>	hypothetical LOC389895
5.39E-05	0.0002248	2.100	<a href="#">230741_at</a>	<a href="#">NA</a>	NA
0.0001761	0.0004668	2.097	<a href="#">223041_at</a>	<a href="#">CD99L2</a>	CD99 molecule-like 2
9.85E-05	0.0003242	2.093	<a href="#">220136_s_at</a>	<a href="#">CRYBA2</a>	crystallin, beta A2
0.0005295	0.0010321	2.092	<a href="#">204891_s_at</a>	<a href="#">LCK</a>	lymphocyte-specific protein tyrosine kinase
0.0002176	0.0005473	2.087	<a href="#">218486_at</a>	<a href="#">KLF11</a>	Kruppel-like factor 11
0.0004168	0.0008622	2.082	<a href="#">202920_at</a>	<a href="#">ANK2</a>	ankyrin 2, neuronal
0.0002331	0.0005773	2.081	<a href="#">225784_s_at</a>	<a href="#">ZC4H2</a>	zinc finger, C4H2 domain containing
3.63E-05	0.0001789	2.080	<a href="#">202665_s_at</a>	<a href="#">WIPF1</a>	WAS/WASL interacting protein family, member 1
0.0002129	0.0005372	2.080	<a href="#">228667_at</a>	<a href="#">AGPAT4</a>	1-acylglycerol-3-phosphate O-acyltransferase 4 (lysophosphatidic acid acyltransferase, delta)
3.98E-05	0.0001887	2.075	<a href="#">202014_at</a>	<a href="#">PPP1R15A</a>	protein phosphatase 1, regulatory (inhibitor) subunit 15A



0.0006094	0.0011432	2.073	<a href="#">205347 s at</a>	<a href="#">TMSB15A</a>	thymosin beta 15a
0.0005606	0.0010733	2.072	<a href="#">239474 at</a>	NA	NA
4.30E-05	0.0001945	2.067	<a href="#">201825 s at</a>	<a href="#">SCCPDH</a>	saccharopine dehydrogenase (putative)
0.0004235	0.0008713	2.064	<a href="#">210151 s at</a>	<a href="#">DYRK3</a>	dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 3
0.0004617	0.0009271	2.062	<a href="#">210619 s at</a>	<a href="#">HYAL1</a>	hyaluronoglucosaminidase 1
0.0001319	0.0003878	2.062	<a href="#">232341 x at</a>	<a href="#">HABP4</a>	hyaluronan binding protein 4
0.000304	0.0006915	2.055	<a href="#">1556346 at</a>	NA	NA
0.000223	0.0005559	2.054	<a href="#">202973 x at</a>	<a href="#">FAM13A</a>	family with sequence similarity 13, member A
0.0004213	0.0008691	2.041	<a href="#">209602 s at</a>	<a href="#">GATA3</a>	GATA binding protein 3
8.86E-05	0.0003063	2.039	<a href="#">219061 s at</a>	<a href="#">LAGE3</a>	L antigen family, member 3
0.0007895	0.0013979	2.031	<a href="#">1554127 s at</a>	<a href="#">MSRB3</a>	methionine sulfoxide reductase B3
0.0003716	0.0007941	2.027	<a href="#">230264 s at</a>	<a href="#">AP1S2</a>	adaptor-related protein complex 1, sigma 2 subunit
0.000597	0.0011264	2.021	<a href="#">1552287 s at</a>	<a href="#">AFG3L1</a>	AFG3 ATPase family gene 3-like 1 (S. cerevisiae)
0.00033	0.0007294	2.021	<a href="#">37028 at</a>	<a href="#">PPP1R15A</a>	protein phosphatase 1, regulatory (inhibitor) subunit 15A
0.0005671	0.001083	2.018	<a href="#">236167 at</a>	NA	NA
7.22E-05	0.0002698	2.018	<a href="#">225355 at</a>	<a href="#">NEURL1B</a>	neutralized homolog 1B (Drosophila)
0.0003909	0.0008237	2.007	<a href="#">1553960 at</a>	<a href="#">SNX21</a>	sorting nexin family member 21
0.0003749	0.0008004	2.007	<a href="#">207264 at</a>	<a href="#">KDELR3</a>	KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 3
0.0002045	0.0005207	0.499	<a href="#">204879 at</a>	<a href="#">PDPN</a>	podoplanin
0.0002675	0.0006307	0.497	<a href="#">218204 s at</a>	<a href="#">FYCO1</a>	FYVE and coiled-coil domain containing 1
8.62E-05	0.0003022	0.496	<a href="#">225726 s at</a>	<a href="#">PLEKHH1</a>	pleckstrin homology domain containing, family H (with MyTH4 domain) member 1
0.0007987	0.0014075	0.496	<a href="#">238461 at</a>	<a href="#">EIF4E3</a>	eukaryotic translation initiation factor 4E family member 3
0.000554	0.0010642	0.496	<a href="#">203962 s at</a>	<a href="#">NEBL</a>	nebulette
0.0007581	0.0013561	0.496	<a href="#">235542 at</a>	<a href="#">TET3</a>	tet oncogene family member 3
0.000448	0.0009076	0.494	<a href="#">226313 at</a>	<a href="#">C10orf35</a>	chromosome 10 open reading frame 35
0.0003118	0.0007028	0.494	<a href="#">219625 s at</a>	<a href="#">COL4A3BP</a>	collagen, type IV, alpha 3 (Goodpasture antigen) binding protein
0.0004962	0.0009856	0.494	<a href="#">1554246 at</a>	<a href="#">C1orf210</a>	chromosome 1 open reading frame 210
0.0009701	0.0016296	0.491	<a href="#">239309 at</a>	<a href="#">DLX6</a>	distal-less homeobox 6
0.0006377	0.0011799	0.491	<a href="#">239412 at</a>	<a href="#">IRF5</a>	interferon regulatory factor 5
0.0003121	0.0007028	0.489	<a href="#">1554894 a at</a>	<a href="#">PCBD2</a>	pterin-4 alpha-carbinolamine dehydratase/dimerization cofactor of hepatocyte nuclear factor 1 alpha (TCF1) 2
0.0001303	0.0003841	0.489	<a href="#">238451 at</a>	<a href="#">MPP7</a>	membrane protein, palmitoylated 7 (MAGUK p55 subfamily member 7)
0.0009084	0.0015558	0.488	<a href="#">225299 at</a>	<a href="#">MYO5B</a>	myosin VB
0.0007881	0.0013965	0.488	<a href="#">235165 at</a>	<a href="#">PARD6B</a>	par-6 partitioning defective 6 homolog beta (C. elegans)
0.0001159	0.0003595	0.486	<a href="#">223466 x at</a>	<a href="#">COL4A3BP</a>	collagen, type IV, alpha 3 (Goodpasture antigen) binding protein
8.83E-05	0.0003062	0.482	<a href="#">221636 s at</a>	<a href="#">MOSC2</a>	MOCO sulphurase C-terminal domain containing 2
0.0007847	0.0013915	0.481	<a href="#">202342 s at</a>	<a href="#">TRIM2</a>	tripartite motif-containing 2
0.0002339	0.0005786	0.480	<a href="#">201200 at</a>	<a href="#">CREG1</a>	cellular repressor of E1A-stimulated genes 1
0.0004454	0.0009048	0.480	<a href="#">1556773 at</a>	NA	NA
0.000853	0.0014801	0.480	<a href="#">227404 s at</a>	<a href="#">EGR1</a>	early growth response 1
0.0004223	0.0008704	0.479	<a href="#">205770 at</a>	<a href="#">GSR</a>	glutathione reductase
0.0001089	0.0003439	0.479	<a href="#">239273 s at</a>	<a href="#">MMP28</a>	matrix metalloproteinase 28
7.80E-05	0.0002843	0.479	<a href="#">213260 at</a>	<a href="#">FOXC1</a>	forkhead box C1

2.80E-05	0.0001599	0.479	<a href="#">1553055_a_at</a>	<a href="#">SLFN5</a>	schlafen family member 5
0.0001694	0.0004522	0.478	<a href="#">203021_at</a>	<a href="#">SLPI</a>	secretory leukocyte peptidase inhibitor
5.69E-05	0.0002314	0.478	<a href="#">219527_at</a>	<a href="#">MOSC2</a>	MOCO sulphurase C-terminal domain containing 2
0.0006617	0.0012173	0.478	<a href="#">223232_s_at</a>	<a href="#">CGN</a>	cingulin
0.0001031	0.0003321	0.478	<a href="#">212686_at</a>	<a href="#">PPM1H</a>	protein phosphatase 1H (PP2C domain containing)
0.0004239	0.0008713	0.477	<a href="#">222587_s_at</a>	<a href="#">GALNT7</a>	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 7 (GalNAc-T7)
0.0002639	0.0006281	0.476	<a href="#">204922_at</a>	<a href="#">C11orf80</a>	chromosome 11 open reading frame 80
6.08E-05	0.0002416	0.476	<a href="#">229377_at</a>	<a href="#">GRTP1</a>	growth hormone regulated TBC protein 1
4.31E-05	0.0001945	0.476	<a href="#">209885_at</a>	<a href="#">RHOD</a>	ras homolog gene family, member D
0.0001084	0.0003428	0.475	<a href="#">218510_x_at</a>	<a href="#">FAM134B</a>	family with sequence similarity 134, member B
5.23E-05	0.0002206	0.474	<a href="#">41660_at</a>	<a href="#">CELSR1</a>	cadherin, EGF LAG seven-pass G-type receptor 1 (flamingo homolog, Drosophila)
0.0005554	0.001066	0.474	<a href="#">209980_s_at</a>	<a href="#">SHMT1</a>	serine hydroxymethyltransferase 1 (soluble)
0.0008976	0.0015422	0.474	<a href="#">242726_at</a>	NA	NA
0.0004639	0.0009299	0.473	<a href="#">227875_at</a>	<a href="#">KLHL13</a>	kelch-like 13 (Drosophila)
0.0006999	0.0012743	0.473	<a href="#">235026_at</a>	<a href="#">C12orf66</a>	chromosome 12 open reading frame 66
0.0009919	0.0016612	0.473	<a href="#">239653_at</a>	NA	NA
9.76E-05	0.0003217	0.472	<a href="#">244665_at</a>	NA	NA
0.0002634	0.0006276	0.472	<a href="#">209385_s_at</a>	<a href="#">PROSC</a>	proline synthetase co-transcribed homolog (bacterial)
0.0002964	0.0006817	0.472	<a href="#">215724_at</a>	<a href="#">PLD1</a>	phospholipase D1, phosphatidylcholine-specific
0.0001273	0.0003777	0.471	<a href="#">202409_at</a>	<a href="#">IGF2</a>	insulin-like growth factor 2 (somatomedin A)
0.0001189	0.0003653	0.471	<a href="#">240354_at</a>	<a href="#">C12orf54</a>	chromosome 12 open reading frame 54
0.0001017	0.0003294	0.470	<a href="#">205678_at</a>	<a href="#">AP3B2</a>	adaptor-related protein complex 3, beta 2 subunit
6.89E-05	0.0002609	0.470	<a href="#">222269_at</a>	<a href="#">APOOL</a>	apolipoprotein O-like
6.68E-05	0.0002577	0.469	<a href="#">218657_at</a>	<a href="#">RAPGEFL1</a>	Rap guanine nucleotide exchange factor (GEF)-like 1
0.0008512	0.0014793	0.468	<a href="#">214040_s_at</a>	<a href="#">GSN</a>	gelsolin (amyloidosis, Finnish type)
5.69E-05	0.0002314	0.467	<a href="#">204636_at</a>	<a href="#">COL17A1</a>	collagen, type XVII, alpha 1
0.0001135	0.0003554	0.467	<a href="#">1569201_a_at</a>	NA	NA
0.0003789	0.0008059	0.466	<a href="#">211367_s_at</a>	<a href="#">CASP1</a>	caspase 1, apoptosis-related cysteine peptidase (interleukin 1, beta, convertase)
0.0008122	0.0014258	0.465	<a href="#">212339_at</a>	<a href="#">EPB41L1</a>	erythrocyte membrane protein band 4.1-like 1
0.0001211	0.000367	0.465	<a href="#">223183_at</a>	<a href="#">AGPAT3</a>	1-acylglycerol-3-phosphate O-acyltransferase 3
0.0002362	0.0005833	0.464	<a href="#">208165_s_at</a>	<a href="#">PRSS16</a>	protease, serine, 16 (thymus)
0.0003282	0.0007261	0.464	<a href="#">200766_at</a>	<a href="#">CTSD</a>	cathepsin D
0.0009957	0.0016664	0.463	<a href="#">216969_s_at</a>	<a href="#">KIF22</a>	kinesin family member 22
0.0003152	0.0007084	0.462	<a href="#">213912_at</a>	<a href="#">TBC1D30</a>	TBC1 domain family, member 30
2.59E-05	0.0001529	0.462	<a href="#">1556194_a_at</a>	NA	NA
0.0001739	0.0004626	0.461	<a href="#">226622_at</a>	<a href="#">MUC20</a>	mucin 20, cell surface associated
0.0005036	0.0009971	0.460	<a href="#">235684_s_at</a>	<a href="#">SESN3</a>	sestrin 3
0.0007415	0.0013316	0.460	<a href="#">206300_s_at</a>	<a href="#">PTH LH</a>	parathyroid hormone-like hormone
0.0009233	0.0015733	0.460	<a href="#">228338_at</a>	<a href="#">LOC120376</a>	Uncharacterized protein LOC120376
0.0001989	0.0005087	0.460	<a href="#">213618_at</a>	<a href="#">ARAP2</a>	ArfGAP with RhoGAP domain, ankyrin repeat and PH domain 2
3.90E-05	0.0001868	0.460	<a href="#">226789_at</a>	<a href="#">LOC647121</a>	embigin homolog (mouse) pseudogene
0.0001578	0.0004327	0.460	<a href="#">226213_at</a>	<a href="#">ERBB3</a>	v-erb-b2 erythroblastic leukemia viral oncogene homolog 3 (avian)
0.0007687	0.0013707	0.459	<a href="#">230788_at</a>	<a href="#">GCNT2</a>	glucosaminyl (N-acetyl) transferase 2, I-branching enzyme (I blood group)

0.0005804	0.0011018	0.459	<a href="#">242055 at</a>	<a href="#">NA</a>	NA
0.000752	0.0013484	0.458	<a href="#">1555812 a at</a>	<a href="#">ARHGDI B</a>	Rho GDP dissociation inhibitor (GDI) beta
2.20E-06	5.78E-05	0.458	<a href="#">204475 at</a>	<a href="#">MMP1</a>	matrix metalloproteinase 1 (interstitial collagenase)
0.0002906	0.0006705	0.458	<a href="#">228220 at</a>	<a href="#">FCHO2</a>	FCH domain only 2
0.0005115	0.0010092	0.457	<a href="#">220266 s at</a>	<a href="#">KLF4</a>	Kruppel-like factor 4 (gut)
1.83E-05	0.0001277	0.457	<a href="#">227346 at</a>	<a href="#">IKZF1</a>	IKAROS family zinc finger 1 (Ikaros)
0.000946	0.0016058	0.457	<a href="#">218792 s at</a>	<a href="#">BSPRY</a>	B-box and SPRY domain containing
3.01E-05	0.0001672	0.457	<a href="#">206023 at</a>	<a href="#">NMU</a>	neuromedin U
0.0002493	0.0006074	0.456	<a href="#">217924 at</a>	<a href="#">C6orf106</a>	chromosome 6 open reading frame 106
9.02E-05	0.0003095	0.455	<a href="#">225407 at</a>	<a href="#">MBP</a>	myelin basic protein
5.01E-05	0.0002149	0.455	<a href="#">223194 s at</a>	<a href="#">SLC22A23</a>	solute carrier family 22, member 23
0.0005409	0.0010471	0.454	<a href="#">215386 at</a>	<a href="#">NA</a>	NA
0.0004269	0.0008751	0.454	<a href="#">225412 at</a>	<a href="#">TMEM87B</a>	transmembrane protein 87B
5.06E-05	0.0002163	0.453	<a href="#">229296 at</a>	<a href="#">LOC100128501</a>	hypothetical protein LOC100128501
3.53E-05	0.0001774	0.453	<a href="#">224164 at</a>	<a href="#">TPM3</a>	tropomyosin 3
0.0003804	0.0008068	0.453	<a href="#">213849 s at</a>	<a href="#">PPP2R2B</a>	protein phosphatase 2 (formerly 2A), regulatory subunit B, beta isoform
0.0001366	0.0003965	0.453	<a href="#">201566 x at</a>	<a href="#">ID2</a>	inhibitor of DNA binding 2, dominant negative helix-loop-helix protein
3.30E-06	6.60E-05	0.452	<a href="#">227475 at</a>	<a href="#">FOXQ1</a>	forkhead box Q1
0.0003035	0.0006915	0.452	<a href="#">217974 at</a>	<a href="#">TM7SF3</a>	transmembrane 7 superfamily member 3
1.80E-06	5.65E-05	0.451	<a href="#">202744 at</a>	<a href="#">SLC20A2</a>	solute carrier family 20 (phosphate transporter), member 2
0.000267	0.0006302	0.451	<a href="#">203726 s at</a>	<a href="#">LAMA3</a>	laminin, alpha 3
0.0001672	0.0004469	0.451	<a href="#">216782 at</a>	<a href="#">NA</a>	NA
0.000489	0.0009733	0.451	<a href="#">219764 at</a>	<a href="#">FZD10</a>	frizzled homolog 10 (Drosophila)
0.0005686	0.001084	0.451	<a href="#">1558404 at</a>	<a href="#">LOC644242</a>	hypothetical protein LOC644242
0.0001399	0.0004014	0.450	<a href="#">207414 s at</a>	<a href="#">PCSK6</a>	proprotein convertase subtilisin/kexin type 6
1.42E-05	0.000113	0.449	<a href="#">224840 at</a>	<a href="#">FKBP5</a>	FK506 binding protein 5
0.0002755	0.0006435	0.449	<a href="#">219985 at</a>	<a href="#">HS3ST3A1</a>	heparan sulfate (glucosamine) 3-O-sulfotransferase 3A1
8.58E-05	0.0003017	0.448	<a href="#">207076 s at</a>	<a href="#">ASS1</a>	argininosuccinate synthetase 1
0.0002228	0.0005559	0.447	<a href="#">213340 s at</a>	<a href="#">KIAA0495</a>	KIAA0495
0.0001988	0.0005087	0.447	<a href="#">217744 s at</a>	<a href="#">PERP</a>	PERP, TP53 apoptosis effector
0.0001387	0.0003995	0.447	<a href="#">235112 at</a>	<a href="#">NA</a>	NA
9.15E-05	0.0003106	0.446	<a href="#">213900 at</a>	<a href="#">C9orf61</a>	chromosome 9 open reading frame 61
6.31E-05	0.0002473	0.446	<a href="#">201691 s at</a>	<a href="#">TPD52</a>	tumor protein D52
3.52E-05	0.0001774	0.446	<a href="#">223092 at</a>	<a href="#">ANKH</a>	ankylosis, progressive homolog (mouse)
0.0004128	0.0008551	0.444	<a href="#">230193 at</a>	<a href="#">WDR66</a>	WD repeat domain 66
2.67E-05	0.0001554	0.444	<a href="#">203837 at</a>	<a href="#">MAP3K5</a>	mitogen-activated protein kinase kinase kinase 5
0.0001446	0.0004084	0.444	<a href="#">232017 at</a>	<a href="#">TJP2</a>	tight junction protein 2 (zona occludens 2)
3.91E-05	0.0001869	0.444	<a href="#">230406 at</a>	<a href="#">NA</a>	NA
4.30E-05	0.0001945	0.444	<a href="#">202436 s at</a>	<a href="#">CYP1B1</a>	cytochrome P450, family 1, subfamily B, polypeptide 1
0.0002822	0.0006564	0.443	<a href="#">242417 at</a>	<a href="#">PLEKHA7</a>	pleckstrin homology domain containing, family A member 7
0.0001912	0.0004975	0.443	<a href="#">226113 at</a>	<a href="#">ZNF436</a>	zinc finger protein 436
2.16E-05	0.0001391	0.443	<a href="#">203964 at</a>	<a href="#">NMI</a>	N-myc (and STAT) interactor
9.10E-06	9.35E-05	0.443	<a href="#">202085 at</a>	<a href="#">TJP2</a>	tight junction protein 2 (zona occludens 2)

8.90E-06	9.27E-05	0.443	<a href="#">222477_s_at</a>	<a href="#">TM7SF3</a>	transmembrane 7 superfamily member 3
1.20E-05	0.0001036	0.442	<a href="#">212964_at</a>	<a href="#">HIC2</a>	hypermethylated in cancer 2
0.0003846	0.000815	0.441	<a href="#">223182_s_at</a>	<a href="#">AGPAT3</a>	1-acylglycerol-3-phosphate O-acyltransferase 3
7.59E-05	0.0002795	0.441	<a href="#">218776_s_at</a>	<a href="#">TMEM62</a>	transmembrane protein 62
6.61E-05	0.0002563	0.439	<a href="#">225619_at</a>	<a href="#">SLAIN1</a>	SLAIN motif family, member 1
4.98E-05	0.0002145	0.439	<a href="#">203143_s_at</a>	<a href="#">KIAA0040</a>	KIAA0040
0.0009234	0.0015733	0.439	<a href="#">1553989_a_at</a>	<a href="#">ATP6V1C2</a>	ATPase, H+ transporting, lysosomal 42kDa, V1 subunit C2
0.0001751	0.0004647	0.439	<a href="#">225440_at</a>	<a href="#">AGPAT3</a>	1-acylglycerol-3-phosphate O-acyltransferase 3
0.000129	0.0003813	0.439	<a href="#">212240_s_at</a>	<a href="#">PIK3R1</a>	phosphoinositide-3-kinase, regulatory subunit 1 (alpha)
0.0001058	0.0003374	0.438	<a href="#">232406_at</a>	<a href="#">NA</a>	NA
0.0009866	0.0016536	0.438	<a href="#">230493_at</a>	<a href="#">SHISA2</a>	shisa homolog 2 (Xenopus laevis)
3.09E-05	0.0001676	0.437	<a href="#">201900_s_at</a>	<a href="#">AKR1A1</a>	aldo-keto reductase family 1, member A1 (aldehyde reductase)
0.0003235	0.0007213	0.436	<a href="#">207382_at</a>	<a href="#">TP63</a>	tumor protein p63
0.0002203	0.000553	0.436	<a href="#">1560800_at</a>	<a href="#">NA</a>	NA
0.000113	0.0003549	0.435	<a href="#">203855_at</a>	<a href="#">WDR47</a>	WD repeat domain 47
0.0004017	0.0008402	0.435	<a href="#">220038_at</a>	<a href="#">SGK3</a>	serum/glucocorticoid regulated kinase family, member 3
4.32E-05	0.0001946	0.435	<a href="#">233487_s_at</a>	<a href="#">LRRC8A</a>	leucine rich repeat containing 8 family, member A
7.40E-05	0.0002743	0.434	<a href="#">221610_s_at</a>	<a href="#">STAP2</a>	signal transducing adaptor family member 2
0.0005319	0.0010341	0.434	<a href="#">238673_at</a>	<a href="#">SAMD12</a>	sterile alpha motif domain containing 12
1.06E-05	9.90E-05	0.434	<a href="#">218677_at</a>	<a href="#">S100A14</a>	S100 calcium binding protein A14
0.0002535	0.000611	0.434	<a href="#">230769_at</a>	<a href="#">DENND2C</a>	DENN/MADD domain containing 2C
9.80E-06	9.43E-05	0.434	<a href="#">214058_at</a>	<a href="#">MYCL1</a>	v-myc myelocytomatosis viral oncogene homolog 1, lung carcinoma derived (avian)
0.0001545	0.0004297	0.433	<a href="#">218035_s_at</a>	<a href="#">RBM47</a>	RNA binding motif protein 47
2.70E-05	0.0001559	0.432	<a href="#">226803_at</a>	<a href="#">CHMP4C</a>	chromatin modifying protein 4C
0.0006218	0.001159	0.432	<a href="#">216488_s_at</a>	<a href="#">ATP11A</a>	ATPase, class VI, type 11A
0.0007189	0.0013003	0.432	<a href="#">212128_s_at</a>	<a href="#">DAG1</a>	dystroglycan 1 (dystrophin-associated glycoprotein 1)
3.11E-05	0.0001676	0.432	<a href="#">220225_at</a>	<a href="#">IRX4</a>	iroquois homeobox 4
0.000628	0.0011667	0.432	<a href="#">210237_at</a>	<a href="#">ARTN</a>	artemin
0.0002937	0.0006762	0.431	<a href="#">1559607_s_at</a>	<a href="#">GBP6</a>	guanylate binding protein family, member 6
0.0002977	0.0006827	0.431	<a href="#">205455_at</a>	<a href="#">MST1R</a>	macrophage stimulating 1 receptor (c-met-related tyrosine kinase)
9.34E-05	0.0003143	0.430	<a href="#">205109_s_at</a>	<a href="#">ARHGEF4</a>	Rho guanine nucleotide exchange factor (GEF) 4
0.000909	0.0015558	0.430	<a href="#">223500_at</a>	<a href="#">CPLX1</a>	complexin 1
0.0001152	0.0003588	0.429	<a href="#">1562226_at</a>	<a href="#">VWDE</a>	von Willebrand factor D and EGF domains
4.54E-05	0.0002001	0.429	<a href="#">208908_s_at</a>	<a href="#">CAST</a>	calpastatin
9.23E-05	0.0003122	0.428	<a href="#">225133_at</a>	<a href="#">KLF3</a>	Kruppel-like factor 3 (basic)
4.06E-05	0.0001895	0.428	<a href="#">235141_at</a>	<a href="#">MARVELD2</a>	MARVEL domain containing 2
0.0004356	0.0008897	0.428	<a href="#">223094_s_at</a>	<a href="#">ANKH</a>	ankylosis, progressive homolog (mouse)
4.07E-05	0.0001895	0.428	<a href="#">242354_at</a>	<a href="#">NA</a>	NA
0.0002298	0.0005697	0.427	<a href="#">204667_at</a>	<a href="#">FOXA1</a>	forkhead box A1
3.12E-05	0.0001676	0.427	<a href="#">220354_at</a>	<a href="#">hCG_1774568</a>	similar to hCG1774568
0.0001157	0.0003595	0.427	<a href="#">212719_at</a>	<a href="#">PHLPP</a>	PH domain and leucine rich repeat protein phosphatase
4.27E-05	0.0001945	0.427	<a href="#">209748_at</a>	<a href="#">SPAST</a>	spastin
4.08E-05	0.0001895	0.426	<a href="#">212812_at</a>	<a href="#">NA</a>	NA

3.80E-05	0.0001836	0.425	<a href="#">238455 at</a>	<a href="#">NA</a>	NA
0.000438	0.000893	0.425	<a href="#">220285 at</a>	<a href="#">FAM108B1</a>	family with sequence similarity 108, member B1
0.0001654	0.0004431	0.425	<a href="#">204573 at</a>	<a href="#">CROT</a>	carnitine O-octanoyltransferase
3.25E-05	0.0001705	0.425	<a href="#">1556518 at</a>	<a href="#">NA</a>	NA
4.05E-05	0.0001895	0.424	<a href="#">243256 at</a>	<a href="#">MKNK1</a>	MAP kinase interacting serine/threonine kinase 1
0.0001405	0.0004016	0.424	<a href="#">226499 at</a>	<a href="#">NRARP</a>	NOTCH-regulated ankyrin repeat protein
0.0001828	0.0004817	0.424	<a href="#">206582 s at</a>	<a href="#">GPR56</a>	G protein-coupled receptor 56
0.0001266	0.0003761	0.423	<a href="#">228575 at</a>	<a href="#">IL20RB</a>	interleukin 20 receptor beta
0.0003572	0.000773	0.423	<a href="#">204967 at</a>	<a href="#">SHROOM2</a>	shroom family member 2
0.0001173	0.0003613	0.422	<a href="#">1552477 a at</a>	<a href="#">IRF6</a>	interferon regulatory factor 6
0.0001417	0.0004045	0.422	<a href="#">222603 at</a>	<a href="#">ERMP1</a>	endoplasmic reticulum metalloproteinase 1
1.08E-05	9.98E-05	0.422	<a href="#">203509 at</a>	<a href="#">SORL1</a>	sortilin-related receptor, L(DLR class) A repeats-containing
2.08E-05	0.0001359	0.422	<a href="#">218802 at</a>	<a href="#">CCDC109B</a>	coiled-coil domain containing 109B
0.0003893	0.0008211	0.422	<a href="#">1559425 at</a>	<a href="#">NA</a>	NA
0.0009665	0.0016259	0.421	<a href="#">203210 s at</a>	<a href="#">RFC5</a>	replication factor C (activator 1) 5, 36.5kDa
0.0008999	0.0015449	0.420	<a href="#">212974 at</a>	<a href="#">DENND3</a>	DENN/MADD domain containing 3
0.0002501	0.0006074	0.420	<a href="#">243000 at</a>	<a href="#">CDK6</a>	cyclin-dependent kinase 6
0.0007084	0.0012838	0.420	<a href="#">201009 s at</a>	<a href="#">TXNIP</a>	thioredoxin interacting protein
9.50E-05	0.0003168	0.420	<a href="#">207030 s at</a>	<a href="#">CSRP2</a>	cysteine and glycine-rich protein 2
5.04E-05	0.0002158	0.420	<a href="#">202761 s at</a>	<a href="#">SYNE2</a>	spectrin repeat containing, nuclear envelope 2
0.000626	0.0011649	0.419	<a href="#">234331 s at</a>	<a href="#">FAM84A</a>	family with sequence similarity 84, member A
0.0001636	0.0004398	0.418	<a href="#">206099 at</a>	<a href="#">PRKCH</a>	protein kinase C, eta
2.25E-05	0.0001414	0.417	<a href="#">228894 at</a>	<a href="#">LOC253842</a>	hypothetical protein LOC253842
0.0003457	0.0007542	0.417	<a href="#">202435 s at</a>	<a href="#">CYP1B1</a>	cytochrome P450, family 1, subfamily B, polypeptide 1
6.70E-06	8.10E-05	0.416	<a href="#">219211 at</a>	<a href="#">USP18</a>	ubiquitin specific peptidase 18
3.80E-05	0.0001836	0.415	<a href="#">205560 at</a>	<a href="#">PCSK5</a>	proprotein convertase subtilisin/kexin type 5
0.0003021	0.0006906	0.415	<a href="#">205968 at</a>	<a href="#">KCNS3</a>	potassium voltage-gated channel, delayed-rectifier, subfamily S, member 3
0.0002754	0.0006435	0.414	<a href="#">236381 s at</a>	<a href="#">WDR8</a>	WD repeat domain 8
5.49E-05	0.0002273	0.414	<a href="#">215023 s at</a>	<a href="#">PEX1</a>	peroxisomal biogenesis factor 1
1.33E-05	0.0001106	0.414	<a href="#">235068 at</a>	<a href="#">ZDHHC21</a>	zinc finger, DHHC-type containing 21
0.0001344	0.0003926	0.414	<a href="#">232081 at</a>	<a href="#">NA</a>	NA
0.0007541	0.00135	0.413	<a href="#">231950 at</a>	<a href="#">ZNF658</a>	zinc finger protein 658
4.04E-05	0.0001895	0.412	<a href="#">236009 at</a>	<a href="#">NA</a>	NA
0.0003315	0.0007312	0.412	<a href="#">238320 at</a>	<a href="#">NCRNA00084</a>	non-protein coding RNA 84
0.0003338	0.0007351	0.412	<a href="#">225354 s at</a>	<a href="#">SH3BGRL2</a>	SH3 domain binding glutamic acid-rich protein like 2
0.0005034	0.0009971	0.411	<a href="#">207675 x at</a>	<a href="#">ARTN</a>	artemin
0.0002735	0.0006415	0.410	<a href="#">235092 at</a>	<a href="#">NA</a>	NA
0.0003199	0.0007174	0.410	<a href="#">209631 s at</a>	<a href="#">GPR37</a>	G protein-coupled receptor 37 (endothelin receptor type B-like)
0.0003758	0.0008016	0.410	<a href="#">210372 s at</a>	<a href="#">TPD52L1</a>	tumor protein D52-like 1
1.90E-06	5.65E-05	0.410	<a href="#">201764 at</a>	<a href="#">TMEM106C</a>	transmembrane protein 106C
0.0001619	0.0004394	0.410	<a href="#">207949 s at</a>	<a href="#">ICA1</a>	islet cell autoantigen 1, 69kDa
0.0009611	0.0016211	0.409	<a href="#">233924 s at</a>	<a href="#">EXOC6</a>	exocyst complex component 6
0.0006961	0.0012693	0.407	<a href="#">216918 s at</a>	<a href="#">DST</a>	dystonin

0.0003112	0.0007028	0.406	<a href="#">206683_at</a>	<a href="#">ZNF165</a>	zinc finger protein 165
0.0001003	0.0003277	0.406	<a href="#">216258_s_at</a>	<a href="#">SERPINB13</a>	serpin peptidase inhibitor, clade B (ovalbumin), member 13
7.14E-05	0.0002685	0.406	<a href="#">236279_at</a>	NA	NA
0.000163	0.0004398	0.405	<a href="#">224097_s_at</a>	<a href="#">F11R</a>	F11 receptor
0.0003645	0.0007834	0.405	<a href="#">223632_s_at</a>	<a href="#">BCAN</a>	brevican
3.22E-05	0.0001702	0.405	<a href="#">218499_at</a>	<a href="#">RP6-213H19.1</a>	serine/threonine protein kinase MST4
0.0001792	0.0004744	0.404	<a href="#">228653_at</a>	<a href="#">SAMD5</a>	sterile alpha motif domain containing 5
1.72E-05	0.0001227	0.404	<a href="#">235583_at</a>	<a href="#">ILDR1</a>	immunoglobulin-like domain containing receptor 1
1.21E-05	0.0001036	0.404	<a href="#">219476_at</a>	<a href="#">C1orf116</a>	chromosome 1 open reading frame 116
0.0003712	0.000794	0.403	<a href="#">206515_at</a>	<a href="#">CYP4F3</a>	cytochrome P450, family 4, subfamily F, polypeptide 3
9.41E-05	0.000316	0.403	<a href="#">200965_s_at</a>	<a href="#">ABLIM1</a>	actin binding LIM protein 1
0.0006032	0.0011335	0.402	<a href="#">243862_at</a>	NA	NA
6.12E-05	0.0002423	0.401	<a href="#">212599_at</a>	<a href="#">AUTS2</a>	autism susceptibility candidate 2
0.0004657	0.0009318	0.401	<a href="#">1565863_at</a>	NA	NA
0.0001625	0.0004398	0.401	<a href="#">225009_at</a>	<a href="#">CMTM4</a>	CKLF-like MARVEL transmembrane domain containing 4
0.0006721	0.0012325	0.401	<a href="#">200946_x_at</a>	<a href="#">GLUD1</a>	glutamate dehydrogenase 1
3.95E-05	0.000188	0.401	<a href="#">225846_at</a>	<a href="#">RBM35A</a>	RNA binding motif protein 35A
4.24E-05	0.0001945	0.400	<a href="#">212560_at</a>	<a href="#">SORL1</a>	sortilin-related receptor, L(DLR class) A repeats-containing
0.0006107	0.0011432	0.400	<a href="#">219115_s_at</a>	<a href="#">IL20RA</a>	interleukin 20 receptor, alpha
2.49E-05	0.0001493	0.400	<a href="#">226597_at</a>	<a href="#">REEP6</a>	receptor accessory protein 6
0.0001376	0.0003977	0.400	<a href="#">205286_at</a>	<a href="#">TFAP2C</a>	transcription factor AP-2 gamma (activating enhancer binding protein 2 gamma)
0.0001427	0.0004064	0.399	<a href="#">224460_s_at</a>	<a href="#">L2HGDH</a>	L-2-hydroxyglutarate dehydrogenase
0.0004496	0.0009093	0.399	<a href="#">231257_at</a>	<a href="#">TCERG1L</a>	transcription elongation regulator 1-like
0.0007967	0.0014058	0.398	<a href="#">228010_at</a>	<a href="#">PPP2R2C</a>	protein phosphatase 2 (formerly 2A), regulatory subunit B, gamma isoform
0.0002135	0.0005382	0.398	<a href="#">238599_at</a>	<a href="#">IRAK1BP1</a>	interleukin-1 receptor-associated kinase 1 binding protein 1
0.0009678	0.0016269	0.398	<a href="#">225123_at</a>	NA	NA
0.0007784	0.0013836	0.398	<a href="#">210480_s_at</a>	<a href="#">MYO6</a>	myosin VI
0.0002208	0.000553	0.398	<a href="#">227642_at</a>	<a href="#">TFCP2L1</a>	transcription factor CP2-like 1
0.0001667	0.000446	0.398	<a href="#">219265_at</a>	<a href="#">MOBK12B</a>	MOB1, Mps One Binder kinase activator-like 2B (yeast)
0.0005803	0.0011018	0.397	<a href="#">226226_at</a>	<a href="#">TMEM45B</a>	transmembrane protein 45B
2.89E-05	0.0001634	0.396	<a href="#">208792_s_at</a>	<a href="#">CLU</a>	clusterin
9.02E-05	0.0003095	0.396	<a href="#">1555765_a_at</a>	<a href="#">GNG4</a>	guanine nucleotide binding protein (G protein), gamma 4
2.50E-06	6.21E-05	0.396	<a href="#">231916_at</a>	<a href="#">EXOSC6</a>	exosome component 6
3.00E-07	3.57E-05	0.396	<a href="#">218706_s_at</a>	<a href="#">GRAMD3</a>	GRAM domain containing 3
0.0001925	0.0004994	0.395	<a href="#">242899_at</a>	NA	NA
5.29E-05	0.0002223	0.395	<a href="#">206600_s_at</a>	<a href="#">SLC16A5</a>	solute carrier family 16, member 5 (monocarboxylic acid transporter 6)
0.0001461	0.0004114	0.395	<a href="#">214045_at</a>	<a href="#">LIAS</a>	lipoic acid synthetase
4.58E-05	0.0002011	0.394	<a href="#">220262_s_at</a>	<a href="#">DLK2</a>	delta-like 2 homolog (Drosophila)
0.0001217	0.0003679	0.393	<a href="#">215723_s_at</a>	<a href="#">PLD1</a>	phospholipase D1, phosphatidylcholine-specific
0.0003314	0.0007312	0.393	<a href="#">201565_s_at</a>	<a href="#">ID2</a>	inhibitor of DNA binding 2, dominant negative helix-loop-helix protein
0.0004965	0.0009856	0.392	<a href="#">213075_at</a>	<a href="#">OLFML2A</a>	olfactomedin-like 2A
0.0003882	0.0008195	0.392	<a href="#">209212_s_at</a>	<a href="#">KLF5</a>	Kruppel-like factor 5 (intestinal)
7.43E-05	0.0002749	0.392	<a href="#">226907_at</a>	<a href="#">PPP1R14C</a>	protein phosphatase 1, regulatory (inhibitor) subunit 14C

5.77E-05	0.0002338	0.391	<a href="#">208791_at</a>	<a href="#">CLU</a>	clusterin
4.72E-05	0.0002048	0.391	<a href="#">243278_at</a>	<a href="#">FOXP2</a>	forkhead box P2
0.0003416	0.0007462	0.391	<a href="#">208937_s_at</a>	<a href="#">ID1</a>	inhibitor of DNA binding 1, dominant negative helix-loop-helix protein
6.00E-07	4.84E-05	0.391	<a href="#">204971_at</a>	<a href="#">CSTA</a>	cystatin A (stefin A)
0.000602	0.0011333	0.391	<a href="#">211776_s_at</a>	<a href="#">EPB41L3</a>	erythrocyte membrane protein band 4.1-like 3
3.03E-05	0.0001675	0.390	<a href="#">235075_at</a>	<a href="#">DSG3</a>	desmoglein 3 (pemphigus vulgaris antigen)
3.56E-05	0.0001777	0.390	<a href="#">220520_s_at</a>	<a href="#">NUP62CL</a>	nucleoporin 62kDa C-terminal like
0.0001012	0.0003288	0.390	<a href="#">235201_at</a>	<a href="#">FOXP2</a>	forkhead box P2
0.0002264	0.0005631	0.390	<a href="#">223574_x_at</a>	<a href="#">PPP2R2C</a>	protein phosphatase 2 (formerly 2A), regulatory subunit B, gamma isoform
0.0006129	0.0011462	0.390	<a href="#">227379_at</a>	<a href="#">MBOAT1</a>	membrane bound O-acyltransferase domain containing 1
1.52E-05	0.0001177	0.389	<a href="#">238689_at</a>	<a href="#">GPR110</a>	G protein-coupled receptor 110
0.0002825	0.0006565	0.389	<a href="#">226489_at</a>	<a href="#">TMCC3</a>	transmembrane and coiled-coil domain family 3
0.0004236	0.0008713	0.389	<a href="#">239945_at</a>	<a href="#">NA</a>	NA
0.0001634	0.0004398	0.387	<a href="#">214434_at</a>	<a href="#">HSPA12A</a>	heat shock 70kDa protein 12A
4.61E-05	0.000202	0.387	<a href="#">238593_at</a>	<a href="#">C11orf80</a>	chromosome 11 open reading frame 80
0.0004057	0.0008454	0.386	<a href="#">213996_at</a>	<a href="#">YPEL1</a>	yippee-like 1 (Drosophila)
4.05E-05	0.0001895	0.384	<a href="#">219481_at</a>	<a href="#">TTC13</a>	tetratricopeptide repeat domain 13
0.0007754	0.0013794	0.384	<a href="#">235050_at</a>	<a href="#">SLC2A12</a>	solute carrier family 2 (facilitated glucose transporter), member 12
1.38E-05	0.000111	0.384	<a href="#">239132_at</a>	<a href="#">NOS1</a>	nitric oxide synthase 1 (neuronal)
0.0001319	0.0003878	0.383	<a href="#">203126_at</a>	<a href="#">IMPA2</a>	inositol(myo)-1(or 4)-monophosphatase 2
9.25E-05	0.0003122	0.383	<a href="#">233634_at</a>	<a href="#">MARVELD3</a>	MARVEL domain containing 3
0.0004006	0.0008387	0.383	<a href="#">227228_s_at</a>	<a href="#">CCDC88C</a>	coiled-coil domain containing 88C
4.60E-06	7.04E-05	0.383	<a href="#">202421_at</a>	<a href="#">IGSF3</a>	immunoglobulin superfamily, member 3
0.0008373	0.0014596	0.382	<a href="#">205724_at</a>	<a href="#">PKP1</a>	plakophilin 1 (ectodermal dysplasia/skin fragility syndrome)
0.000335	0.0007358	0.382	<a href="#">224221_s_at</a>	<a href="#">VAV3</a>	vav 3 guanine nucleotide exchange factor
0.0003698	0.0007918	0.381	<a href="#">227032_at</a>	<a href="#">PLXNA2</a>	plexin A2
1.72E-05	0.0001227	0.380	<a href="#">202454_s_at</a>	<a href="#">ERBB3</a>	v-erb-b2 erythroblastic leukemia viral oncogene homolog 3 (avian)
0.000572	0.0010886	0.380	<a href="#">231727_s_at</a>	<a href="#">MIF4GD</a>	MIF4G domain containing
7.00E-06	8.24E-05	0.380	<a href="#">204224_s_at</a>	<a href="#">GCH1</a>	GTP cyclohydrolase 1
0.0003389	0.0007418	0.380	<a href="#">226114_at</a>	<a href="#">ZNF436</a>	zinc finger protein 436
8.43E-05	0.0002992	0.380	<a href="#">36499_at</a>	<a href="#">CELSR2</a>	cadherin, EGF LAG seven-pass G-type receptor 2 (flamingo homolog, Drosophila)
3.90E-06	6.75E-05	0.380	<a href="#">229764_at</a>	<a href="#">TPRG1</a>	tumor protein p63 regulated 1
0.0003582	0.000773	0.380	<a href="#">204199_at</a>	<a href="#">RALGPS1</a>	Ral GEF with PH domain and SH3 binding motif 1
2.57E-05	0.0001525	0.379	<a href="#">233198_at</a>	<a href="#">GOLGA2L1</a>	golgi autoantigen, golgin subfamily a, 2-like 1
0.0003099	0.0007007	0.379	<a href="#">222496_s_at</a>	<a href="#">RBM47</a>	RNA binding motif protein 47
3.00E-06	6.40E-05	0.379	<a href="#">225618_at</a>	<a href="#">ARHGAP27</a>	Rho GTPase activating protein 27
0.0001447	0.0004084	0.379	<a href="#">210547_x_at</a>	<a href="#">ICA1</a>	islet cell autoantigen 1, 69kDa
0.0007086	0.0012838	0.379	<a href="#">225611_at</a>	<a href="#">MAST4</a>	microtubule associated serine/threonine kinase family member 4
4.70E-06	7.04E-05	0.378	<a href="#">223245_at</a>	<a href="#">STRBP</a>	spermatid perinuclear RNA binding protein
0.000107	0.0003393	0.377	<a href="#">1555716_a_at</a>	<a href="#">CXADR</a>	coxsackie virus and adenovirus receptor
1.37E-05	0.000111	0.377	<a href="#">202831_at</a>	<a href="#">GPX2</a>	glutathione peroxidase 2 (gastrointestinal)
7.70E-06	8.41E-05	0.377	<a href="#">209596_at</a>	<a href="#">MXRA5</a>	matrix-remodelling associated 5
0.0001906	0.0004965	0.376	<a href="#">207549_x_at</a>	<a href="#">CD46</a>	CD46 molecule, complement regulatory protein



5.71E-05	0.0002318	0.376	<a href="#">1438_at</a>	<a href="#">EPHB3</a>	EPH receptor B3
2.86E-05	0.0001621	0.376	<a href="#">218692_at</a>	<a href="#">GOLSYN</a>	Golgi-localized protein
0.0002439	0.0005981	0.375	<a href="#">210145_at</a>	<a href="#">PLA2G4A</a>	phospholipase A2, group IVA (cytosolic, calcium-dependent)
0.0009764	0.0016389	0.375	<a href="#">212681_at</a>	<a href="#">EPB41L3</a>	erythrocyte membrane protein band 4.1-like 3
8.27E-05	0.0002973	0.375	<a href="#">235146_at</a>	<a href="#">TMCC3</a>	transmembrane and coiled-coil domain family 3
6.70E-06	8.10E-05	0.375	<a href="#">224650_at</a>	<a href="#">MAL2</a>	mal, T-cell differentiation protein 2
0.0003583	0.000773	0.374	<a href="#">219344_at</a>	<a href="#">SLC29A3</a>	solute carrier family 29 (nucleoside transporters), member 3
0.0007183	0.0013003	0.374	<a href="#">227863_at</a>	<a href="#">CTSD</a>	cathepsin D
9.06E-05	0.0003099	0.374	<a href="#">1569114_at</a>	<a href="#">NA</a>	NA
0.0002843	0.0006593	0.373	<a href="#">219676_at</a>	<a href="#">ZSCAN16</a>	zinc finger and SCAN domain containing 16
8.67E-05	0.000303	0.373	<a href="#">1554544_a_at</a>	<a href="#">MBP</a>	myelin basic protein
0.000411	0.0008525	0.372	<a href="#">244353_s_at</a>	<a href="#">SLC2A12</a>	solute carrier family 2 (facilitated glucose transporter), member 12
4.25E-05	0.0001945	0.372	<a href="#">218764_at</a>	<a href="#">PRKCH</a>	protein kinase C, eta
0.0001249	0.0003721	0.372	<a href="#">211806_s_at</a>	<a href="#">KCNJ15</a>	potassium inwardly-rectifying channel, subfamily J, member 15
0.0001519	0.0004253	0.371	<a href="#">225436_at</a>	<a href="#">FAM108C1</a>	family with sequence similarity 108, member C1
0.0001564	0.0004313	0.371	<a href="#">211126_s_at</a>	<a href="#">CSRP2</a>	cysteine and glycine-rich protein 2
0.0001135	0.0003554	0.371	<a href="#">209950_s_at</a>	<a href="#">VILL</a>	villin-like
0.0002525	0.0006099	0.370	<a href="#">234980_at</a>	<a href="#">TMEM56</a>	transmembrane protein 56
5.00E-06	7.16E-05	0.370	<a href="#">220161_s_at</a>	<a href="#">EPB41L4B</a>	erythrocyte membrane protein band 4.1 like 4B
0.0001903	0.0004963	0.370	<a href="#">231969_at</a>	<a href="#">STOX2</a>	storkhead box 2
0.0001276	0.0003781	0.370	<a href="#">209800_at</a>	<a href="#">KRT16</a>	keratin 16
0.0001209	0.0003669	0.370	<a href="#">218865_at</a>	<a href="#">MOSC1</a>	MOCO sulphurase C-terminal domain containing 1
0.0009642	0.0016233	0.369	<a href="#">230006_s_at</a>	<a href="#">SVIP</a>	small VCP/p97-interacting protein
0.0004486	0.000908	0.367	<a href="#">240911_at</a>	<a href="#">NOS1</a>	nitric oxide synthase 1 (neuronal)
6.20E-06	7.96E-05	0.367	<a href="#">209581_at</a>	<a href="#">PLA2G16</a>	phospholipase A2, group XVI
9.94E-05	0.0003257	0.366	<a href="#">221664_s_at</a>	<a href="#">F11R</a>	F11 receptor
0.0006032	0.0011335	0.366	<a href="#">1555829_at</a>	<a href="#">FAM62B</a>	family with sequence similarity 62 (C2 domain containing) member B
0.0002211	0.000553	0.365	<a href="#">222699_s_at</a>	<a href="#">PLEKHF2</a>	pleckstrin homology domain containing, family F (with FYVE domain) member 2
2.50E-06	6.21E-05	0.363	<a href="#">39248_at</a>	<a href="#">AQP3</a>	aquaporin 3 (Gill blood group)
9.91E-05	0.0003257	0.363	<a href="#">226912_at</a>	<a href="#">ZDHHC23</a>	zinc finger, DHHC-type containing 23
1.09E-05	9.98E-05	0.363	<a href="#">226817_at</a>	<a href="#">DSC2</a>	desmocollin 2
0.0003228	0.0007205	0.362	<a href="#">206191_at</a>	<a href="#">ENTPD3</a>	ectonucleoside triphosphate diphosphohydrolase 3
0.0002661	0.0006302	0.361	<a href="#">234970_at</a>	<a href="#">TC2N</a>	tandem C2 domains, nuclear
2.91E-05	0.0001641	0.361	<a href="#">229396_at</a>	<a href="#">OVOL1</a>	ovo-like 1 (Drosophila)
0.0007302	0.0013162	0.360	<a href="#">230005_at</a>	<a href="#">SVIP</a>	small VCP/p97-interacting protein
0.0005361	0.0010404	0.360	<a href="#">234488_s_at</a>	<a href="#">GMCL1L</a>	germ cell-less homolog 1 (Drosophila)-like
0.0006857	0.0012533	0.360	<a href="#">205552_s_at</a>	<a href="#">OAS1</a>	2-5'-oligoadenylate synthetase 1, 40/46kDa
0.0001881	0.0004922	0.360	<a href="#">219132_at</a>	<a href="#">PELI2</a>	pellino homolog 2 (Drosophila)
1.24E-05	0.0001054	0.360	<a href="#">231365_at</a>	<a href="#">NA</a>	NA
5.60E-06	7.63E-05	0.360	<a href="#">218885_s_at</a>	<a href="#">GALNT12</a>	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 12 (GalNAc-T12)
0.0002111	0.0005345	0.359	<a href="#">218960_at</a>	<a href="#">TMPRSS4</a>	transmembrane protease, serine 4
0.0001619	0.0004394	0.359	<a href="#">235955_at</a>	<a href="#">MARVELD2</a>	MARVEL domain containing 2
7.74E-05	0.0002827	0.359	<a href="#">224856_at</a>	<a href="#">FKBP5</a>	FK506 binding protein 5

3.22E-05	0.0001702	0.357	<a href="#">210119_at</a>	<a href="#">KCNJ15</a>	potassium inwardly-rectifying channel, subfamily J, member 15
0.0006841	0.0012514	0.357	<a href="#">219551_at</a>	<a href="#">EAF2</a>	ELL associated factor 2
3.43E-05	0.0001755	0.355	<a href="#">228865_at</a>	<a href="#">C1orf116</a>	chromosome 1 open reading frame 116
0.0008605	0.0014909	0.354	<a href="#">202525_at</a>	<a href="#">PRSS8</a>	protease, serine, 8
3.03E-05	0.0001675	0.354	<a href="#">227892_at</a>	<a href="#">PRKAA2</a>	protein kinase, AMP-activated, alpha 2 catalytic subunit
0.0007529	0.0013489	0.353	<a href="#">209784_s_at</a>	<a href="#">JAG2</a>	jagged 2
0.0001177	0.0003621	0.352	<a href="#">206400_at</a>	<a href="#">LGALS7</a>	lectin, galactoside-binding, soluble, 7
5.50E-06	7.63E-05	0.351	<a href="#">209270_at</a>	<a href="#">LAMB3</a>	laminin, beta 3
6.09E-05	0.0002416	0.350	<a href="#">1552685_a_at</a>	<a href="#">GRHL1</a>	grainyhead-like 1 (Drosophila)
0.000337	0.0007383	0.350	<a href="#">210652_s_at</a>	<a href="#">TTC39A</a>	tetratricopeptide repeat domain 39A
0.0007819	0.0013877	0.349	<a href="#">221245_s_at</a>	<a href="#">FZD5</a>	frizzled homolog 5 (Drosophila)
6.29E-05	0.0002469	0.349	<a href="#">213424_at</a>	<a href="#">KIAA0895</a>	KIAA0895
2.09E-05	0.0001362	0.349	<a href="#">228977_at</a>	<a href="#">LOC729680</a>	hypothetical protein LOC729680
4.60E-06	7.04E-05	0.348	<a href="#">1555007_s_at</a>	<a href="#">WDR66</a>	WD repeat domain 66
0.0005537	0.0010642	0.348	<a href="#">229225_at</a>	<a href="#">NRP2</a>	neuropilin 2
1.10E-05	9.99E-05	0.347	<a href="#">203747_at</a>	<a href="#">AQP3</a>	aquaporin 3 (Gill blood group)
3.43E-05	0.0001755	0.347	<a href="#">211259_s_at</a>	<a href="#">BMP7</a>	bone morphogenetic protein 7
0.0001605	0.0004372	0.347	<a href="#">205543_at</a>	<a href="#">HSPA4L</a>	heat shock 70kDa protein 4-like
0.0004597	0.0009247	0.345	<a href="#">214203_s_at</a>	<a href="#">PRODH</a>	proline dehydrogenase (oxidase) 1
0.0001965	0.0005049	0.345	<a href="#">242871_at</a>	<a href="#">PAQR5</a>	progesterin and adipoQ receptor family member V
5.90E-06	7.80E-05	0.344	<a href="#">45288_at</a>	<a href="#">ABHD6</a>	abhydrolase domain containing 6
4.80E-06	7.05E-05	0.344	<a href="#">201820_at</a>	<a href="#">KRT5</a>	keratin 5
5.90E-05	0.0002369	0.342	<a href="#">1552283_s_at</a>	<a href="#">ZDHHC11</a>	zinc finger, DHHC-type containing 11
0.0009312	0.001583	0.342	<a href="#">1554835_a_at</a>	<a href="#">B3GNT5</a>	UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 5
6.13E-05	0.0002423	0.341	<a href="#">213456_at</a>	<a href="#">SOSTDC1</a>	sclerostin domain containing 1
0.0002062	0.0005238	0.341	<a href="#">221552_at</a>	<a href="#">ABHD6</a>	abhydrolase domain containing 6
5.83E-05	0.0002354	0.340	<a href="#">223704_s_at</a>	<a href="#">DMRT2</a>	doublesex and mab-3 related transcription factor 2
0.000136	0.0003955	0.339	<a href="#">204039_at</a>	<a href="#">CEBPA</a>	CCAAT/enhancer binding protein (C/EBP), alpha
5.96E-05	0.0002381	0.338	<a href="#">210136_at</a>	<a href="#">MBP</a>	myelin basic protein
1.15E-05	0.0001021	0.337	<a href="#">1559606_at</a>	<a href="#">GBP6</a>	guanylate binding protein family, member 6
0.000225	0.0005603	0.337	<a href="#">32137_at</a>	<a href="#">JAG2</a>	jagged 2
3.26E-05	0.0001706	0.337	<a href="#">203836_s_at</a>	<a href="#">MAP3K5</a>	mitogen-activated protein kinase kinase kinase 5
5.31E-05	0.0002223	0.336	<a href="#">233252_s_at</a>	<a href="#">STRBP</a>	spermatid perinuclear RNA binding protein
8.30E-06	8.89E-05	0.336	<a href="#">226068_at</a>	<a href="#">SYK</a>	spleen tyrosine kinase
0.0006144	0.0011481	0.336	<a href="#">213974_at</a>	<a href="#">ADAMTSL3</a>	ADAMTS-like 3
0.0007214	0.0013038	0.335	<a href="#">228241_at</a>	<a href="#">AGR3</a>	anterior gradient homolog 3 (Xenopus laevis)
1.50E-06	5.65E-05	0.335	<a href="#">202890_at</a>	<a href="#">MAP7</a>	microtubule-associated protein 7
9.56E-05	0.0003183	0.333	<a href="#">237063_at</a>	<a href="#">NA</a>	NA
0.0001402	0.0004016	0.333	<a href="#">212978_at</a>	<a href="#">LRRC8B</a>	leucine rich repeat containing 8 family, member B
3.75E-05	0.0001827	0.331	<a href="#">211401_s_at</a>	<a href="#">FGFR2</a>	fibroblast growth factor receptor 2
5.60E-06	7.63E-05	0.331	<a href="#">219756_s_at</a>	<a href="#">POF1B</a>	premature ovarian failure, 1B
0.0001227	0.0003687	0.330	<a href="#">213100_at</a>	<a href="#">UNC5B</a>	unc-5 homolog B (C. elegans)
7.82E-05	0.0002843	0.330	<a href="#">226863_at</a>	<a href="#">FAM110C</a>	family with sequence similarity 110, member C

3.30E-06	6.60E-05	0.330	<a href="#">218559_s_at</a>	<a href="#">MAFB</a>	v-maf musculoaponeurotic fibrosarcoma oncogene homolog B (avian)
0.000425	0.0008728	0.329	<a href="#">213285_at</a>	<a href="#">TMEM30B</a>	transmembrane protein 30B
0.0001568	0.0004313	0.327	<a href="#">225822_at</a>	<a href="#">TMEM125</a>	transmembrane protein 125
0.000445	0.0009048	0.327	<a href="#">229546_at</a>	<a href="#">LOC653602</a>	hypothetical LOC653602
7.36E-05	0.0002733	0.326	<a href="#">1566764_at</a>	<a href="#">MACC1</a>	metastasis associated in colon cancer 1
3.80E-05	0.0001836	0.326	<a href="#">213030_s_at</a>	<a href="#">PLXNA2</a>	plexin A2
0.0001361	0.0003955	0.326	<a href="#">212966_at</a>	<a href="#">HIC2</a>	hypermethylated in cancer 2
2.54E-05	0.0001515	0.325	<a href="#">235651_at</a>	NA	NA
0.0001056	0.0003374	0.324	<a href="#">226302_at</a>	<a href="#">ATP8B1</a>	ATPase, class I, type 8B, member 1
1.43E-05	0.000113	0.324	<a href="#">235683_at</a>	<a href="#">SESN3</a>	sestrin 3
0.0003962	0.0008324	0.324	<a href="#">208084_at</a>	<a href="#">ITGB6</a>	integrin, beta 6
5.27E-05	0.0002219	0.322	<a href="#">219856_at</a>	<a href="#">C1orf116</a>	chromosome 1 open reading frame 116
0.0008863	0.0015239	0.321	<a href="#">227803_at</a>	<a href="#">ENPP5</a>	ectonucleotide pyrophosphatase/phosphodiesterase 5 (putative function)
0.0001561	0.0004313	0.321	<a href="#">218813_s_at</a>	<a href="#">SH3GLB2</a>	SH3-domain GRB2-like endophilin B2
0.000336	0.0007369	0.320	<a href="#">205180_s_at</a>	<a href="#">ADAM8</a>	ADAM metallopeptidase domain 8
0.0008279	0.0014499	0.319	<a href="#">210505_at</a>	<a href="#">ADH7</a>	alcohol dehydrogenase 7 (class IV), mu or sigma polypeptide
3.10E-05	0.0001676	0.318	<a href="#">204351_at</a>	<a href="#">S100P</a>	S100 calcium binding protein P
8.38E-05	0.0002992	0.318	<a href="#">204503_at</a>	<a href="#">EVPL</a>	envoplakin
7.20E-06	8.26E-05	0.317	<a href="#">231381_at</a>	<a href="#">HESRG</a>	hypothetical LOC790952
3.53E-05	0.0001774	0.317	<a href="#">223246_s_at</a>	<a href="#">STRBP</a>	spermatid perinuclear RNA binding protein
2.00E-07	3.57E-05	0.316	<a href="#">204734_at</a>	<a href="#">KRT15</a>	keratin 15
2.37E-05	0.0001452	0.316	<a href="#">227241_at</a>	<a href="#">MUC15</a>	mucin 15, cell surface associated
9.69E-05	0.0003208	0.316	<a href="#">220318_at</a>	<a href="#">EPN3</a>	epsin 3
9.40E-06	9.40E-05	0.314	<a href="#">1554921_a_at</a>	<a href="#">SCEL</a>	sciellin
0.0005485	0.0010573	0.314	<a href="#">200841_s_at</a>	<a href="#">EPRS</a>	glutamyl-prolyl-tRNA synthetase
1.40E-05	0.0001122	0.314	<a href="#">221646_s_at</a>	<a href="#">ZDHC11</a>	zinc finger, DHHC-type containing 11
0.0002603	0.0006208	0.314	<a href="#">228485_s_at</a>	<a href="#">SLC44A1</a>	solute carrier family 44, member 1
7.31E-05	0.0002718	0.314	<a href="#">227461_at</a>	<a href="#">STON2</a>	stonin 2
1.84E-05	0.000128	0.313	<a href="#">226278_at</a>	<a href="#">SVIP</a>	small VCP/p97-interacting protein
0.0001164	0.0003595	0.313	<a href="#">235150_at</a>	NA	NA
0.0005135	0.0010122	0.313	<a href="#">228221_at</a>	<a href="#">SLC44A3</a>	solute carrier family 44, member 3
0.0001405	0.0004016	0.313	<a href="#">219648_at</a>	<a href="#">MREG</a>	melanoregulin
0.0002124	0.0005366	0.312	<a href="#">208153_s_at</a>	<a href="#">FAT2</a>	FAT tumor suppressor homolog 2 (Drosophila)
0.0002665	0.0006302	0.311	<a href="#">205769_at</a>	<a href="#">SLC27A2</a>	solute carrier family 27 (fatty acid transporter), member 2
9.50E-06	9.41E-05	0.311	<a href="#">205595_at</a>	<a href="#">DSG3</a>	desmoglein 3 (pemphigus vulgaris antigen)
6.49E-05	0.0002534	0.311	<a href="#">218693_at</a>	<a href="#">TSPAN15</a>	tetraspanin 15
0.0002969	0.0006817	0.310	<a href="#">233251_at</a>	<a href="#">STRBP</a>	spermatid perinuclear RNA binding protein
0.0002497	0.0006074	0.310	<a href="#">219976_at</a>	<a href="#">HOOK1</a>	hook homolog 1 (Drosophila)
4.16E-05	0.0001923	0.309	<a href="#">222150_s_at</a>	<a href="#">PION</a>	pigeon homolog (Drosophila)
4.20E-06	6.83E-05	0.308	<a href="#">223075_s_at</a>	<a href="#">AIF1L</a>	allograft inflammatory factor 1-like
3.60E-06	6.75E-05	0.308	<a href="#">225864_at</a>	<a href="#">FAM84B</a>	family with sequence similarity 84, member B
0.0003606	0.0007758	0.308	<a href="#">235247_at</a>	NA	NA
2.70E-06	6.21E-05	0.307	<a href="#">212070_at</a>	<a href="#">GPR56</a>	G protein-coupled receptor 56

3.00E-07	3.57E-05	0.307	<a href="#">209283_at</a>	<a href="#">CRYAB</a>	crystallin, alpha B
0.0003593	0.0007737	0.307	<a href="#">218807_at</a>	<a href="#">VAV3</a>	vav 3 guanine nucleotide exchange factor
0.0001247	0.0003721	0.307	<a href="#">205032_at</a>	<a href="#">ITGA2</a>	integrin, alpha 2 (CD49B, alpha 2 subunit of VLA-2 receptor)
6.80E-06	8.13E-05	0.306	<a href="#">205559_s_at</a>	<a href="#">PCSK5</a>	proprotein convertase subtilisin/kexin type 5
4.10E-06	6.82E-05	0.305	<a href="#">206276_at</a>	<a href="#">LY6D</a>	lymphocyte antigen 6 complex, locus D
0.0007369	0.0013255	0.305	<a href="#">232122_s_at</a>	<a href="#">VEPH1</a>	ventricular zone expressed PH domain homolog 1 (zebrafish)
1.80E-06	5.65E-05	0.304	<a href="#">205603_s_at</a>	<a href="#">DIAPH2</a>	diaphanous homolog 2 (Drosophila)
0.0005899	0.0011161	0.304	<a href="#">214545_s_at</a>	<a href="#">PROSC</a>	proline synthetase co-transcribed homolog (bacterial)
0.0001102	0.0003475	0.303	<a href="#">227717_at</a>	<a href="#">FLJ41603</a>	FLJ41603 protein
0.000135	0.0003939	0.303	<a href="#">212315_s_at</a>	<a href="#">NUP210</a>	nucleoporin 210kDa
0.0005604	0.0010733	0.302	<a href="#">205184_at</a>	<a href="#">GNG4</a>	guanine nucleotide binding protein (G protein), gamma 4
0.0003798	0.0008063	0.301	<a href="#">207655_s_at</a>	<a href="#">BLNK</a>	B-cell linker
0.000836	0.0014585	0.301	<a href="#">211194_s_at</a>	<a href="#">TP63</a>	tumor protein p63
4.09E-05	0.0001895	0.301	<a href="#">215471_s_at</a>	<a href="#">MAP7</a>	microtubule-associated protein 7
0.0009566	0.0016189	0.300	<a href="#">215785_s_at</a>	<a href="#">CYFIP2</a>	cytoplasmic FMR1 interacting protein 2
2.14E-05	0.0001386	0.300	<a href="#">228360_at</a>	<a href="#">LYPD6B</a>	LY6/PLAUR domain containing 6B
7.20E-06	8.26E-05	0.299	<a href="#">205780_at</a>	<a href="#">BIK</a>	BCL2-interacting killer (apoptosis-inducing)
0.0004706	0.00094	0.298	<a href="#">206204_at</a>	<a href="#">GRB14</a>	growth factor receptor-bound protein 14
5.69E-05	0.0002314	0.298	<a href="#">223895_s_at</a>	<a href="#">EPN3</a>	epsin 3
0.0008529	0.0014801	0.298	<a href="#">1553132_a_at</a>	<a href="#">TC2N</a>	tandem C2 domains, nuclear
1.80E-06	5.65E-05	0.298	<a href="#">213947_s_at</a>	<a href="#">NUP210</a>	nucleoporin 210kDa
0.000107	0.0003393	0.297	<a href="#">227314_at</a>	<a href="#">ITGA2</a>	integrin, alpha 2 (CD49B, alpha 2 subunit of VLA-2 receptor)
3.10E-05	0.0001676	0.297	<a href="#">220196_at</a>	<a href="#">MUC16</a>	mucin 16, cell surface associated
1.20E-06	5.65E-05	0.296	<a href="#">217995_at</a>	<a href="#">SQRD1</a>	sulfide quinone reductase-like (yeast)
0.0002437	0.0005981	0.296	<a href="#">203780_at</a>	<a href="#">MPZL2</a>	myelin protein zero-like 2
4.10E-06	6.82E-05	0.296	<a href="#">238029_s_at</a>	<a href="#">SLC16A14</a>	solute carrier family 16, member 14 (monocarboxylic acid transporter 14)
0.0003516	0.0007644	0.296	<a href="#">219503_s_at</a>	<a href="#">TMEM40</a>	transmembrane protein 40
4.00E-05	0.0001892	0.296	<a href="#">226405_s_at</a>	<a href="#">ARRDC1</a>	arrestin domain containing 1
3.51E-05	0.0001774	0.293	<a href="#">202350_s_at</a>	<a href="#">MATN2</a>	matrilin 2
9.70E-06	9.41E-05	0.293	<a href="#">225613_at</a>	<a href="#">MAST4</a>	microtubule associated serine/threonine kinase family member 4
0.0001195	0.0003661	0.293	<a href="#">239862_at</a>	<a href="#">NA</a>	NA
0.0001232	0.0003692	0.293	<a href="#">1554006_a_at</a>	<a href="#">LLGL2</a>	lethal giant larvae homolog 2 (Drosophila)
0.0003217	0.0007187	0.291	<a href="#">209925_at</a>	<a href="#">OCLN</a>	occludin
0.0003352	0.0007358	0.291	<a href="#">223235_s_at</a>	<a href="#">SMOC2</a>	SPARC related modular calcium binding 2
3.27E-05	0.0001707	0.290	<a href="#">228570_at</a>	<a href="#">BTBD11</a>	BTB (POZ) domain containing 11
9.42E-05	0.000316	0.290	<a href="#">240921_at</a>	<a href="#">NA</a>	NA
0.0003282	0.0007261	0.289	<a href="#">227725_at</a>	<a href="#">ST6GALNAC1</a>	ST6 (alpha-N-acetyl-neuraminy-2,3-beta-galactosyl-1,3)-N-acetylgalactosaminide alpha-2,6-sialyltransferase 1
0.0005851	0.0011089	0.289	<a href="#">216521_s_at</a>	<a href="#">BRCC3</a>	BRCA1/BRCA2-containing complex, subunit 3
0.0008453	0.0014713	0.287	<a href="#">1555330_at</a>	<a href="#">GCLC</a>	glutamate-cysteine ligase, catalytic subunit
0.0004054	0.0008454	0.287	<a href="#">202688_at</a>	<a href="#">TNFSF10</a>	tumor necrosis factor (ligand) superfamily, member 10
3.25E-05	0.0001705	0.286	<a href="#">207540_s_at</a>	<a href="#">SYK</a>	spleen tyrosine kinase
0.0002549	0.0006136	0.284	<a href="#">202488_s_at</a>	<a href="#">FXD3</a>	FXD domain containing ion transport regulator 3

1.91E-05	0.0001292	0.284	<a href="#">203638 s at</a>	<a href="#">FGFR2</a>	fibroblast growth factor receptor 2
2.27E-05	0.0001414	0.284	<a href="#">213142 x at</a>	<a href="#">PION</a>	pigeon homolog (Drosophila)
0.0001297	0.0003828	0.283	<a href="#">219938 s at</a>	<a href="#">PSTPIP2</a>	proline-serine-threonine phosphatase interacting protein 2
0.0001578	0.0004327	0.283	<a href="#">242626 at</a>	<a href="#">SAMDS5</a>	sterile alpha motif domain containing 5
3.10E-06	6.55E-05	0.281	<a href="#">232202 at</a>	<a href="#">NA</a>	NA
0.0002668	0.0006302	0.281	<a href="#">243546 at</a>	<a href="#">NA</a>	NA
0.0006716	0.0012325	0.281	<a href="#">242053 at</a>	<a href="#">TSGA10</a>	testis specific, 10
0.0009559	0.0016189	0.280	<a href="#">234730 s at</a>	<a href="#">RIPK4</a>	receptor-interacting serine-threonine kinase 4
0.0003057	0.000694	0.280	<a href="#">213652 at</a>	<a href="#">PCSK5</a>	proprotein convertase subtilisin/kexin type 5
0.0008101	0.0014232	0.280	<a href="#">209098 s at</a>	<a href="#">JAG1</a>	jagged 1 (Alagille syndrome)
7.40E-06	8.32E-05	0.280	<a href="#">217979 at</a>	<a href="#">TSPAN13</a>	tetraspanin 13
4.00E-06	6.75E-05	0.278	<a href="#">229901 at</a>	<a href="#">ZNF488</a>	zinc finger protein 488
4.41E-05	0.000196	0.278	<a href="#">214598 at</a>	<a href="#">CLDN8</a>	claudin 8
0.0005612	0.0010735	0.277	<a href="#">229842 at</a>	<a href="#">ELF3</a>	E74-like factor 3 (ets domain transcription factor, epithelial-specific )
0.0003038	0.0006915	0.277	<a href="#">228557 at</a>	<a href="#">L3MBTL4</a>	l(3)mbt-like 4 (Drosophila)
0.0003204	0.0007174	0.277	<a href="#">1553986 at</a>	<a href="#">RASEF</a>	RAS and EF-hand domain containing
0.0001037	0.000333	0.275	<a href="#">202889 x at</a>	<a href="#">MAP7</a>	microtubule-associated protein 7
1.62E-05	0.00012	0.275	<a href="#">235144 at</a>	<a href="#">NA</a>	NA
2.73E-05	0.0001567	0.275	<a href="#">212992 at</a>	<a href="#">AHNAK2</a>	AHNAK nucleoprotein 2
0.0005312	0.0010336	0.274	<a href="#">244418 at</a>	<a href="#">NA</a>	NA
0.0001233	0.0003692	0.273	<a href="#">231195 at</a>	<a href="#">KLRG2</a>	killer cell lectin-like receptor subfamily G, member 2
0.0002529	0.0006103	0.273	<a href="#">235744 at</a>	<a href="#">PPTC7</a>	PTC7 protein phosphatase homolog (S. cerevisiae)
4.79E-05	0.0002075	0.271	<a href="#">225016 at</a>	<a href="#">APCDD1</a>	adenomatosis polyposis coli down-regulated 1
0.0005079	0.0010038	0.270	<a href="#">225140 at</a>	<a href="#">KLF3</a>	Kruppel-like factor 3 (basic)
0.0009976	0.0016683	0.269	<a href="#">219714 s at</a>	<a href="#">CACNA2D3</a>	calcium channel, voltage-dependent, alpha 2/delta subunit 3
0.000515	0.0010129	0.268	<a href="#">201008 s at</a>	<a href="#">TXNIP</a>	thioredoxin interacting protein
1.70E-06	5.65E-05	0.268	<a href="#">201131 s at</a>	<a href="#">CDH1</a>	cadherin 1, type 1, E-cadherin (epithelial)
9.10E-06	9.35E-05	0.268	<a href="#">214580 x at</a>	<a href="#">KRT6A</a>	keratin 6A
3.30E-05	0.0001711	0.267	<a href="#">219395 at</a>	<a href="#">RBM35B</a>	RNA binding motif protein 35B
8.91E-05	0.0003076	0.267	<a href="#">222773 s at</a>	<a href="#">GALNT12</a>	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 12 (GalNAc-T12)
9.01E-05	0.0003095	0.267	<a href="#">244023 at</a>	<a href="#">SYK</a>	spleen tyrosine kinase
0.0001926	0.0004994	0.265	<a href="#">218806 s at</a>	<a href="#">VAV3</a>	vav 3 guanine nucleotide exchange factor
0.0002578	0.0006181	0.264	<a href="#">230964 at</a>	<a href="#">FREM2</a>	FRAS1 related extracellular matrix protein 2
0.0001479	0.0004154	0.264	<a href="#">220035 at</a>	<a href="#">NUP210</a>	nucleoporin 210kDa
0.0008713	0.0015061	0.264	<a href="#">1569433 at</a>	<a href="#">SAMDS5</a>	sterile alpha motif domain containing 5
3.47E-05	0.0001767	0.264	<a href="#">39249 at</a>	<a href="#">AQP3</a>	aquaporin 3 (Gill blood group)
3.28E-05	0.0001709	0.263	<a href="#">203397 s at</a>	<a href="#">GALNT3</a>	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 3 (GalNAc-T3)
1.26E-05	0.0001063	0.263	<a href="#">204019 s at</a>	<a href="#">SH3YL1</a>	SH3 domain containing, Ysc84-like 1 (S. cerevisiae)
2.80E-06	6.21E-05	0.262	<a href="#">202712 s at</a>	<a href="#">CKMT1B</a>	creatine kinase, mitochondrial 1B
6.82E-05	0.00026	0.261	<a href="#">223709 s at</a>	<a href="#">WNT10A</a>	wingless-type MMTV integration site family, member 10A
5.92E-05	0.0002373	0.261	<a href="#">219121 s at</a>	<a href="#">RBM35A</a>	RNA binding motif protein 35A
9.00E-07	5.36E-05	0.261	<a href="#">201015 s at</a>	<a href="#">JUP</a>	junction plakoglobin
9.50E-05	0.0003168	0.261	<a href="#">215425 at</a>	<a href="#">BTG3</a>	BTG family, member 3

0.0006374	0.0011799	0.260	<a href="#">227935 s at</a>	<a href="#">PCGF5</a>	polycomb group ring finger 5
4.58E-05	0.0002011	0.260	<a href="#">1553319 at</a>	<a href="#">OXGR1</a>	oxoglutarate (alpha-ketoglutarate) receptor 1
9.50E-06	9.41E-05	0.260	<a href="#">205625 s at</a>	<a href="#">CALB1</a>	calbindin 1, 28kDa
0.000279	0.0006503	0.259	<a href="#">228494 at</a>	<a href="#">PPP1R9A</a>	protein phosphatase 1, regulatory (inhibitor) subunit 9A
0.0005498	0.0010589	0.258	<a href="#">209269 s at</a>	<a href="#">SYK</a>	spleen tyrosine kinase
5.64E-05	0.000231	0.256	<a href="#">239148 at</a>	<a href="#">MARVELD3</a>	MARVEL domain containing 3
4.42E-05	0.000196	0.256	<a href="#">227506 at</a>	<a href="#">SLC16A9</a>	solute carrier family 16, member 9 (monocarboxylic acid transporter 9)
0.0001238	0.0003703	0.255	<a href="#">218804 at</a>	<a href="#">ANO1</a>	anoctamin 1, calcium activated chloride channel
1.99E-05	0.0001322	0.255	<a href="#">222892 s at</a>	<a href="#">TMEM40</a>	transmembrane protein 40
6.30E-06	7.96E-05	0.254	<a href="#">231849 at</a>	<a href="#">KRT80</a>	keratin 80
5.61E-05	0.000231	0.254	<a href="#">212657 s at</a>	<a href="#">IL1RN</a>	interleukin 1 receptor antagonist
0.0001454	0.0004099	0.254	<a href="#">229242 at</a>	<a href="#">NA</a>	NA
2.10E-06	5.78E-05	0.251	<a href="#">205626 s at</a>	<a href="#">CALB1</a>	calbindin 1, 28kDa
0.0001531	0.0004279	0.251	<a href="#">201428 at</a>	<a href="#">CLDN4</a>	claudin 4
2.00E-05	0.0001322	0.250	<a href="#">202489 s at</a>	<a href="#">FXVD3</a>	FXVD domain containing ion transport regulator 3
3.43E-05	0.0001755	0.250	<a href="#">1553613 s at</a>	<a href="#">FOXC1</a>	forkhead box C1
6.98E-05	0.000263	0.247	<a href="#">224646 x at</a>	<a href="#">H19</a>	H19, imprinted maternally expressed transcript (non-protein coding)
1.90E-06	5.65E-05	0.246	<a href="#">201690 s at</a>	<a href="#">TPD52</a>	tumor protein D52
1.57E-05	0.0001187	0.246	<a href="#">221679 s at</a>	<a href="#">ABHD6</a>	abhydrolase domain containing 6
0.0003339	0.0007351	0.246	<a href="#">223832 s at</a>	<a href="#">CAPNS2</a>	calpain, small subunit 2
1.35E-05	0.000111	0.244	<a href="#">242786 at</a>	<a href="#">NA</a>	NA
0.0001025	0.0003311	0.243	<a href="#">1552496 a at</a>	<a href="#">COBL</a>	cordon-bleu homolog (mouse)
0.0005921	0.001119	0.241	<a href="#">1555269 a at</a>	<a href="#">ANO1</a>	anoctamin 1, calcium activated chloride channel
0.000775	0.0013794	0.241	<a href="#">209114 at</a>	<a href="#">TSPAN1</a>	tetraspanin 1
0.0001143	0.000357	0.241	<a href="#">239595 at</a>	<a href="#">GPX2</a>	glutathione peroxidase 2 (gastrointestinal)
0.0009593	0.0016209	0.240	<a href="#">204989 s at</a>	<a href="#">ITGB4</a>	integrin, beta 4
3.20E-06	6.60E-05	0.240	<a href="#">201079 at</a>	<a href="#">SYNGR2</a>	synaptogyrin 2
2.26E-05	0.0001414	0.239	<a href="#">225325 at</a>	<a href="#">MFSD6</a>	major facilitator superfamily domain containing 6
2.70E-06	6.21E-05	0.238	<a href="#">212316 at</a>	<a href="#">NUP210</a>	nucleoporin 210kDa
4.42E-05	0.000196	0.238	<a href="#">231311 at</a>	<a href="#">NA</a>	NA
8.43E-05	0.0002992	0.237	<a href="#">215729 s at</a>	<a href="#">VGLL1</a>	vestigial like 1 (Drosophila)
8.70E-06	9.19E-05	0.237	<a href="#">203407 at</a>	<a href="#">PPL</a>	periplakin
9.40E-06	9.40E-05	0.237	<a href="#">212096 s at</a>	<a href="#">MTUS1</a>	mitochondrial tumor suppressor 1
2.20E-06	5.78E-05	0.237	<a href="#">211002 s at</a>	<a href="#">TRIM29</a>	tripartite motif-containing 29
1.03E-05	9.66E-05	0.236	<a href="#">203642 s at</a>	<a href="#">COBLL1</a>	COBL-like 1
1.65E-05	0.0001203	0.235	<a href="#">208228 s at</a>	<a href="#">FGFR2</a>	fibroblast growth factor receptor 2
4.30E-06	6.90E-05	0.235	<a href="#">202800 at</a>	<a href="#">SLC1A3</a>	solute carrier family 1 (glial high affinity glutamate transporter), member 3
0.0006708	0.0012321	0.235	<a href="#">213722 at</a>	<a href="#">SOX2</a>	SRY (sex determining region Y)-box 2
0.0004323	0.0008838	0.235	<a href="#">201510 at</a>	<a href="#">ELF3</a>	E74-like factor 3 (ets domain transcription factor, epithelial-specific)
9.33E-05	0.0003143	0.233	<a href="#">229223 at</a>	<a href="#">NA</a>	NA
1.65E-05	0.0001203	0.230	<a href="#">228256 s at</a>	<a href="#">EPB41L4A</a>	erythrocyte membrane protein band 4.1 like 4A
8.71E-05	0.0003039	0.230	<a href="#">227238 at</a>	<a href="#">MUC15</a>	mucin 15, cell surface associated
8.60E-06	9.17E-05	0.228	<a href="#">225301 s at</a>	<a href="#">MYO5B</a>	myosin VB

0.0008242	0.0014446	0.227	<a href="#">225911 at</a>	<a href="#">NPNT</a>	nephronectin
0.000538	0.0010423	0.226	<a href="#">205185 at</a>	<a href="#">SPINK5</a>	serine peptidase inhibitor, Kazal type 5
5.11E-05	0.0002176	0.226	<a href="#">1555516 at</a>	<a href="#">FOXP2</a>	forkhead box P2
1.14E-05	0.0001021	0.226	<a href="#">202504 at</a>	<a href="#">TRIM29</a>	tripartite motif-containing 29
2.08E-05	0.0001359	0.226	<a href="#">232165 at</a>	<a href="#">EPPK1</a>	epiplakin 1
8.00E-07	5.36E-05	0.225	<a href="#">228038 at</a>	<a href="#">SOX2</a>	SRY (sex determining region Y)-box 2
0.0004543	0.0009147	0.223	<a href="#">203639 s at</a>	<a href="#">FGFR2</a>	fibroblast growth factor receptor 2
7.00E-07	5.36E-05	0.222	<a href="#">229568 at</a>	<a href="#">MOBKL2B</a>	MOB1, Mps One Binder kinase activator-like 2B (yeast)
3.81E-05	0.0001837	0.221	<a href="#">242836 at</a>	<a href="#">NA</a>	NA
2.00E-05	0.0001322	0.220	<a href="#">210827 s at</a>	<a href="#">ELF3</a>	E74-like factor 3 (ets domain transcription factor, epithelial-specific )
0.0005924	0.001119	0.220	<a href="#">223427 s at</a>	<a href="#">EPB41L4B</a>	erythrocyte membrane protein band 4.1 like 4B
0.0004079	0.0008484	0.219	<a href="#">213707 s at</a>	<a href="#">DLX5</a>	distal-less homeobox 5
0.0001164	0.0003595	0.218	<a href="#">227209 at</a>	<a href="#">CNTN1</a>	contactin 1
0.0001035	0.0003329	0.218	<a href="#">226844 at</a>	<a href="#">MOBKL2B</a>	MOB1, Mps One Binder kinase activator-like 2B (yeast)
0.0001139	0.0003562	0.216	<a href="#">202005 at</a>	<a href="#">ST14</a>	suppression of tumorigenicity 14 (colon carcinoma)
0.0002372	0.0005849	0.216	<a href="#">209493 at</a>	<a href="#">PDZD2</a>	PDZ domain containing 2
1.58E-05	0.000119	0.215	<a href="#">209590 at</a>	<a href="#">BMP7</a>	bone morphogenetic protein 7
3.36E-05	0.0001731	0.212	<a href="#">226374 at</a>	<a href="#">NA</a>	NA
1.94E-05	0.0001302	0.210	<a href="#">207291 at</a>	<a href="#">PRRG4</a>	proline rich Gla (G-carboxyglutamic acid) 4 (transmembrane)
3.83E-05	0.0001842	0.210	<a href="#">202687 s at</a>	<a href="#">TNFSF10</a>	tumor necrosis factor (ligand) superfamily, member 10
5.20E-06	7.35E-05	0.209	<a href="#">219010 at</a>	<a href="#">C1orf106</a>	chromosome 1 open reading frame 106
7.92E-05	0.000287	0.208	<a href="#">211203 s at</a>	<a href="#">CNTN1</a>	contactin 1
0.0001629	0.0004398	0.208	<a href="#">244264 at</a>	<a href="#">KLRG2</a>	killer cell lectin-like receptor subfamily G, member 2
5.00E-06	7.16E-05	0.208	<a href="#">225667 s at</a>	<a href="#">FAM84A</a>	family with sequence similarity 84, member A
9.40E-06	9.40E-05	0.207	<a href="#">209125 at</a>	<a href="#">KRT6A</a>	keratin 6A
6.60E-06	8.10E-05	0.206	<a href="#">209126 x at</a>	<a href="#">KRT6B</a>	keratin 6B
0.0008801	0.0015159	0.205	<a href="#">1552714 at</a>	<a href="#">CREG2</a>	cellular repressor of E1A-stimulated genes 2
1.67E-05	0.0001214	0.205	<a href="#">206698 at</a>	<a href="#">XK</a>	X-linked Kx blood group (McLeod syndrome)
0.0001195	0.0003661	0.204	<a href="#">214549 x at</a>	<a href="#">SPRR1A</a>	small proline-rich protein 1A
1.56E-05	0.0001184	0.204	<a href="#">203797 at</a>	<a href="#">VSNL1</a>	visinin-like 1
1.30E-06	5.65E-05	0.203	<a href="#">221854 at</a>	<a href="#">PKP1</a>	plakophilin 1 (ectodermal dysplasia/skin fragility syndrome)
3.80E-06	6.75E-05	0.203	<a href="#">223869 at</a>	<a href="#">SOST</a>	sclerosteosis
0.0001204	0.0003667	0.202	<a href="#">1566766 a at</a>	<a href="#">MACC1</a>	metastasis associated in colon cancer 1
4.66E-05	0.000203	0.200	<a href="#">224997 x at</a>	<a href="#">H19</a>	H19, imprinted maternally expressed transcript (non-protein coding)
0.0004386	0.0008934	0.199	<a href="#">232682 at</a>	<a href="#">MREG</a>	melanoregulin
9.25E-05	0.0003122	0.198	<a href="#">237159 x at</a>	<a href="#">AP1S3</a>	adaptor-related protein complex 1, sigma 3 subunit
0.0003208	0.0007174	0.197	<a href="#">227202 at</a>	<a href="#">CNTN1</a>	contactin 1
3.11E-05	0.0001676	0.194	<a href="#">231489 x at</a>	<a href="#">NA</a>	NA
2.37E-05	0.0001452	0.193	<a href="#">1555383 a at</a>	<a href="#">POF1B</a>	premature ovarian failure, 1B
7.19E-05	0.0002691	0.192	<a href="#">203528 at</a>	<a href="#">SEMA4D</a>	sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4D
6.90E-06	8.21E-05	0.192	<a href="#">202388 at</a>	<a href="#">RGS2</a>	regulator of G-protein signaling 2, 24kDa
2.00E-07	3.57E-05	0.192	<a href="#">227492 at</a>	<a href="#">NA</a>	NA



0.0006496	0.001198	0.192	<a href="#">206262 at</a>	<a href="#">ADH1C</a>	alcohol dehydrogenase 1C (class I), gamma polypeptide
1.90E-06	5.65E-05	0.188	<a href="#">232164 s at</a>	<a href="#">EPPK1</a>	epiplakin 1
0.0007055	0.0012812	0.187	<a href="#">1559361 at</a>	<a href="#">MACC1</a>	metastasis associated in colon cancer 1
1.03E-05	9.66E-05	0.187	<a href="#">237885 at</a>	<a href="#">NA</a>	NA
1.80E-06	5.65E-05	0.186	<a href="#">201839 s at</a>	<a href="#">EPCAM</a>	epithelial cell adhesion molecule
0.0002225	0.0005559	0.183	<a href="#">204751 x at</a>	<a href="#">DSC2</a>	desmocollin 2
1.35E-05	0.000111	0.177	<a href="#">229513 at</a>	<a href="#">STRBP</a>	spermatid perinuclear RNA binding protein
9.93E-05	0.0003257	0.174	<a href="#">205220 at</a>	<a href="#">NIACR2</a>	niacin receptor 2
3.73E-05	0.0001825	0.174	<a href="#">223423 at</a>	<a href="#">GPR160</a>	G protein-coupled receptor 160
2.46E-05	0.0001487	0.172	<a href="#">213721 at</a>	<a href="#">SOX2</a>	SRY (sex determining region Y)-box 2
5.60E-06	7.63E-05	0.172	<a href="#">203917 at</a>	<a href="#">CXADR</a>	coxsackie virus and adenovirus receptor
3.64E-05	0.0001789	0.171	<a href="#">203641 s at</a>	<a href="#">COBLL1</a>	COBL-like 1
6.93E-05	0.000262	0.171	<a href="#">201689 s at</a>	<a href="#">TPD52</a>	tumor protein D52
2.92E-05	0.0001642	0.170	<a href="#">219209 at</a>	<a href="#">IFIH1</a>	interferon induced with helicase C domain 1
2.18E-05	0.0001396	0.168	<a href="#">223454 at</a>	<a href="#">CXCL16</a>	chemokine (C-X-C motif) ligand 16
5.39E-05	0.0002248	0.168	<a href="#">203713 s at</a>	<a href="#">LLGL2</a>	lethal giant larvae homolog 2 (Drosophila)
3.80E-06	6.75E-05	0.168	<a href="#">226961 at</a>	<a href="#">PRR15</a>	proline rich 15
0.0003008	0.0006884	0.167	<a href="#">204798 at</a>	<a href="#">MYB</a>	v-myb myeloblastosis viral oncogene homolog (avian)
0.0001324	0.0003881	0.166	<a href="#">219412 at</a>	<a href="#">RAB38</a>	RAB38, member RAS oncogene family
0.0001476	0.0004151	0.166	<a href="#">213796 at</a>	<a href="#">SPRR1A</a>	small proline-rich protein 1A
2.71E-05	0.0001559	0.165	<a href="#">212095 s at</a>	<a href="#">MTUS1</a>	mitochondrial tumor suppressor 1
3.80E-06	6.75E-05	0.165	<a href="#">202790 at</a>	<a href="#">CLDN7</a>	claudin 7
3.92E-05	0.000187	0.164	<a href="#">232056 at</a>	<a href="#">SCEL</a>	sciellin
1.01E-05	9.64E-05	0.163	<a href="#">238692 at</a>	<a href="#">BTBD11</a>	BTB (POZ) domain containing 11
6.70E-06	8.10E-05	0.162	<a href="#">221795 at</a>	<a href="#">NTRK2</a>	neurotrophic tyrosine kinase, receptor, type 2
3.31E-05	0.0001713	0.162	<a href="#">226899 at</a>	<a href="#">UNC5B</a>	unc-5 homolog B (C. elegans)
2.00E-07	3.57E-05	0.161	<a href="#">218186 at</a>	<a href="#">RAB25</a>	RAB25, member RAS oncogene family
1.00E-06	5.51E-05	0.161	<a href="#">232151 at</a>	<a href="#">MACC1</a>	metastasis associated in colon cancer 1
7.00E-06	8.24E-05	0.159	<a href="#">221796 at</a>	<a href="#">NTRK2</a>	neurotrophic tyrosine kinase, receptor, type 2
0.0004462	0.0009048	0.158	<a href="#">204320 at</a>	<a href="#">COL11A1</a>	collagen, type XI, alpha 1
1.22E-05	0.0001041	0.158	<a href="#">206884 s at</a>	<a href="#">SCEL</a>	sciellin
0.0004816	0.0009602	0.156	<a href="#">212909 at</a>	<a href="#">LYPD1</a>	LY6/PLAUR domain containing 1
7.70E-06	8.41E-05	0.155	<a href="#">209591 s at</a>	<a href="#">BMP7</a>	bone morphogenetic protein 7
1.12E-05	0.0001009	0.153	<a href="#">204542 at</a>	<a href="#">ST6GALNAC2</a>	ST6 (alpha-N-acetyl-neuraminy-2,3-beta-galactosyl-1,3)-N-acetylgalactosaminide alpha-2,6-sialyltransferase 2
6.80E-05	0.0002597	0.149	<a href="#">204567 s at</a>	<a href="#">ABCG1</a>	ATP-binding cassette, sub-family G (WHITE), member 1
4.41E-05	0.000196	0.149	<a href="#">225792 at</a>	<a href="#">HOOK1</a>	hook homolog 1 (Drosophila)
7.20E-06	8.26E-05	0.146	<a href="#">219429 at</a>	<a href="#">FA2H</a>	fatty acid 2-hydroxylase
5.90E-05	0.0002369	0.146	<a href="#">227048 at</a>	<a href="#">LAMA1</a>	laminin, alpha 1
3.64E-05	0.0001789	0.145	<a href="#">231929 at</a>	<a href="#">IKZF2</a>	IKAROS family zinc finger 2 (Helios)
2.80E-06	6.21E-05	0.145	<a href="#">216905 s at</a>	<a href="#">ST14</a>	suppression of tumorigenicity 14 (colon carcinoma)
0.0001556	0.0004311	0.143	<a href="#">235852 at</a>	<a href="#">STON2</a>	stonin 2
4.00E-06	6.75E-05	0.143	<a href="#">205190 at</a>	<a href="#">PLS1</a>	plastin 1 (I isoform)

6.00E-07	4.84E-05	0.140	<a href="#">227985 at</a>	<a href="#">NA</a>	NA
3.14E-05	0.0001682	0.140	<a href="#">227899 at</a>	<a href="#">VIT</a>	vitrin
0.0008798	0.0015159	0.139	<a href="#">224407 s at</a>	<a href="#">RP6-213H19.1</a>	serine/threonine protein kinase MST4
2.48E-05	0.0001493	0.136	<a href="#">223278 at</a>	<a href="#">GJB2</a>	gap junction protein, beta 2, 26kDa
8.86E-05	0.0003063	0.136	<a href="#">218990 s at</a>	<a href="#">SPRR3</a>	small proline-rich protein 3
2.15E-05	0.0001389	0.136	<a href="#">204364 s at</a>	<a href="#">REEP1</a>	receptor accessory protein 1
0.0002499	0.0006074	0.136	<a href="#">212093 s at</a>	<a href="#">MTUS1</a>	mitochondrial tumor suppressor 1
7.65E-05	0.0002808	0.135	<a href="#">239155 at</a>	<a href="#">CXADRP1</a>	coxsackie virus and adenovirus receptor pseudogene 1
0.000122	0.0003683	0.132	<a href="#">203798 s at</a>	<a href="#">VSNL1</a>	visinin-like 1
3.80E-06	6.75E-05	0.128	<a href="#">209792 s at</a>	<a href="#">KLK10</a>	kallikrein-related peptidase 10
1.03E-05	9.66E-05	0.126	<a href="#">228051 at</a>	<a href="#">LOC202451</a>	hypothetical protein LOC202451
0.000527	0.0010299	0.124	<a href="#">201688 s at</a>	<a href="#">TPD52</a>	tumor protein D52
5.90E-06	7.80E-05	0.117	<a href="#">201130 s at</a>	<a href="#">CDH1</a>	cadherin 1, type 1, E-cadherin (epithelial)
4.00E-06	6.75E-05	0.113	<a href="#">238513 at</a>	<a href="#">PRRG4</a>	proline rich Gla (G-carboxyglutamic acid) 4 (transmembrane)
8.38E-05	0.0002992	0.110	<a href="#">1555731 a at</a>	<a href="#">AP1S3</a>	adaptor-related protein complex 1, sigma 3 subunit
1.60E-06	5.65E-05	0.109	<a href="#">213050 at</a>	<a href="#">COBL</a>	cordon-bleu homolog (mouse)
3.00E-07	3.57E-05	0.101	<a href="#">204469 at</a>	<a href="#">PTPRZ1</a>	protein tyrosine phosphatase, receptor-type, Z polypeptide 1
2.25E-05	0.0001414	0.099	<a href="#">215465 at</a>	<a href="#">ABCA12</a>	ATP-binding cassette, sub-family A (ABC1), member 12
2.41E-05	0.0001473	0.099	<a href="#">212224 at</a>	<a href="#">ALDH1A1</a>	aldehyde dehydrogenase 1 family, member A1
3.84E-05	0.0001843	0.099	<a href="#">239196 at</a>	<a href="#">ANKRD22</a>	ankyrin repeat domain 22
3.45E-05	0.0001761	0.098	<a href="#">205306 x at</a>	<a href="#">KMO</a>	kynurenine 3-monooxygenase (kynurenine 3-hydroxylase)
3.74E-05	0.0001826	0.097	<a href="#">37892 at</a>	<a href="#">COL11A1</a>	collagen, type XI, alpha 1
1.10E-06	5.65E-05	0.094	<a href="#">211138 s at</a>	<a href="#">KMO</a>	kynurenine 3-monooxygenase (kynurenine 3-hydroxylase)
3.00E-07	3.57E-05	0.091	<a href="#">218966 at</a>	<a href="#">MYO5C</a>	myosin VC
3.50E-06	6.75E-05	0.088	<a href="#">202718 at</a>	<a href="#">IGFBP2</a>	insulin-like growth factor binding protein 2, 36kDa
8.00E-07	5.36E-05	0.088	<a href="#">207935 s at</a>	<a href="#">KRT13</a>	keratin 13
1.90E-06	5.65E-05	0.087	<a href="#">222830 at</a>	<a href="#">GRHL1</a>	grainyhead-like 1 (Drosophila)
1.55E-05	0.000118	0.082	<a href="#">205064 at</a>	<a href="#">SPRR1B</a>	small proline-rich protein 1B (cornifin)
1.50E-06	5.65E-05	0.082	<a href="#">203571 s at</a>	<a href="#">C10orf116</a>	chromosome 10 open reading frame 116
3.80E-05	0.0001836	0.078	<a href="#">203453 at</a>	<a href="#">SCNN1A</a>	sodium channel, nonvoltage-gated 1 alpha
6.30E-06	7.96E-05	0.075	<a href="#">211361 s at</a>	<a href="#">SERPINB13</a>	serpin peptidase inhibitor, clade B (ovalbumin), member 13
1.11E-05	0.0001004	0.073	<a href="#">205569 at</a>	<a href="#">LAMP3</a>	lysosomal-associated membrane protein 3
9.60E-06	9.41E-05	0.071	<a href="#">238439 at</a>	<a href="#">ANKRD22</a>	ankyrin repeat domain 22
1.90E-06	5.65E-05	0.068	<a href="#">215692 s at</a>	<a href="#">MPPED2</a>	metallophosphoesterase domain containing 2
4.70E-06	7.04E-05	0.063	<a href="#">217272 s at</a>	<a href="#">SERPINB13</a>	serpin peptidase inhibitor, clade B (ovalbumin), member 13
1.00E-07	3.57E-05	0.042	<a href="#">209173 at</a>	<a href="#">AGR2</a>	anterior gradient homolog 2 (Xenopus laevis)
7.80E-06	8.44E-05	0.036	<a href="#">205413 at</a>	<a href="#">MPPED2</a>	metallophosphoesterase domain containing 2
< 1e-07	< 1e-07	0.032	<a href="#">231771 at</a>	<a href="#">GJB6</a>	gap junction protein, beta 6, 30kDa
2.70E-06	6.21E-05	0.028	<a href="#">213240 s at</a>	<a href="#">KRT4</a>	keratin 4