

**Supplementary Table 3.**List of genes commonly up-regulated in the microdissected cerebral white matter region after exposure to DBDE at 100 and 1000 ppm ( $\geq 2$ -fold)

Accession no.	Gene title	Symbol	DBDE		
			10 ppm	100 ppm	1000 ppm
Up-regulated (669 genes)					
NM_001191992	outer dense fiber of sperm tails 3B	Odf3b	1.55	5.87	29.88
BI288579	EST	–	1.18	84.91	27.81
NM_001106602	deoxyguanosine kinase	Dguok	0.82	28.19	24.35
XR_085573	similar to mKIAA1783 protein	RGD1560214	0.63	9.81	15.39
BM390363	EST	–	0.82	2.20	12.46
NM_031686	sodium channel, voltage-gated, type VII, alpha	Scn7a	0.89	7.34	11.60
NM_001191598	coiled-coil domain containing 113	Ccdc113	1.33	2.08	11.31
BE113493	EST	–	0.64	12.08	11.20
NM_021666	triadin	Trdn	1.38	5.40	10.88
XM_002725158	Hypothetical LOC290577	LOC290577	1.43	3.86	10.76
BM386877	EST	–	1.65	5.38	10.34
NM_138884	aldo-keto reductase family 1, member D1 (delta 4-3-ketosteroid-5-beta-reductase)	Akr1d1	1.64	10.18	10.25
NM_022928	G protein-coupled receptor kinase 4	Grk4	0.61	4.57	9.86
NM_031635	fucosyltransferase 2 (secretor status included)	Fut2	1.14	5.31	9.83
XM_001061953	Similar to paired immunoglobulin-like type 2 receptor alpha	LOC685020	1.72	10.32	9.74
AW433528	EST	–	1.02	5.75	9.32
AI010736	EST	–	1.13	6.35	9.10
NM_024365	5-hydroxytryptamine (serotonin) receptor 6	Htr6	1.15	14.58	8.69
NM_001033073	defensin alpha-related sequence 1	Defa-rs1	1.05	6.78	8.61
AI233574	EST	–	1.74	4.29	8.20
NM_001135807	H2A histone family, member Y2	H2afy2	1.45	3.10	8.06
NM_139184	proline-rich glycoprotein (sgp158)	Sgp158	1.77	6.77	7.93
NM_001141935	ATPase, class V, type 10A	Atp10a	1.50	7.61	7.67
XM_001061900	histone cluster 1, H4m	Hist1h4m	1.87	5.14	7.57
NM_001106689	Zinc finger protein 593	Znf593	1.01	5.07	7.45
NM_001009920	glutathione S-transferase Yc2 subunit	Gsta5	1.03	2.42	7.39
NM_001191758	cache domain containing 1	Cachd1	1.19	6.53	7.22
AA893390	EST	–	0.54	2.68	7.19
XM_001058380	Hypothetical protein LOC680687	LOC680687	1.44	5.41	7.16
BE119818	EST	–	1.35	7.87	7.12
XR_085609	hypothetical LOC100363796	LOC100363796	0.92	2.13	7.07
AI598332	EST	–	1.02	2.19	6.96
NM_147213	alpha-2u globulin PGCL5	LOC259245	1.98	3.02	6.93
NM_001033964	6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 2	Pfkfb2	1.92	6.05	6.89
BF410174	EST	–	1.09	2.76	6.86
XR_085950	similar to Gene model 784	RGD1561693	1.51	5.62	6.85
BI282568	keratinocyte differentiation associated protein	Krtdap	1.18	5.46	6.83
BF420118	EST	–	1.07	4.91	6.74
AI008553	EST	–	1.88	3.64	6.71
NM_001025417	actin-like 7b	Actl7b	1.66	6.41	6.71
AA850953	EST	–	0.78	10.18	6.61
XM_001059437	Methyl-CpG binding domain protein 4	Mbd4	1.97	5.69	6.58
NM_022213	phosphoinositide-3-kinase, regulatory subunit 3 (gamma)	Pik3r3	0.86	2.82	6.50
BF545930	EST	–	0.89	3.20	6.27
NM_001025661	crumbs homolog 3	Crb3	1.70	4.61	6.19
XM_001081628	SRY-box containing gene 9	Sox9	1.39	3.18	6.17

NM_181378	cathepsin M	Ctsm	1.12	7.07	6.14
AA800583	EST	–	1.21	4.46	6.05
NM_031688	synuclein, gamma (breast cancer-specific protein 1)	Sncg	0.97	4.50	6.05
NM_012873	protamine 2	Prm2	1.71	2.15	5.92
BF409344	EST	–	1.15	2.94	5.84
NM_053851	calcium channel, voltage-dependent, beta 2 subunit	Cacnb2	1.42	3.55	5.70
NM_133536	RAB3C, member RAS oncogene family	Rab3c	1.00	2.31	5.68
AI763952	EST	–	1.08	6.14	5.66
NM_001109289	similar to RIKEN cDNA 1700012B09	RGD1561795	1.05	2.20	5.59
BM389765	EST	–	1.19	2.72	5.56
AW536017	EST	–	1.49	3.52	5.55
XM_001066684	similar to ovostatin-2	RGD1565709	1.19	6.73	5.55
NM_001013994	REST corepressor 2	Rcor2	1.92	2.91	5.53
XR_085857	similar to Kiaa0575	RGD1564091	1.60	2.16	5.49
NM_001025713	similar to melanoma antigen family A, 10	MGC114529	0.92	2.04	5.39
NM_012566	growth factor independent 1 transcription repressor	Gfi1	1.50	2.54	5.38
AI639396	EST	–	1.07	4.82	5.37
XM_578186	Similar to C20orf118	RGD1561343	1.95	4.37	5.35
NM_001108069	serine/threonine kinase 11	Stk11	1.23	10.93	5.32
NM_022628	nephrosis 1 homolog	Nphs1	0.86	2.22	5.27
BF399795	EST	–	1.38	4.48	5.26
BF416438	EST	–	1.20	2.29	5.24
BE113044	EST	–	1.09	2.57	5.22
NM_023968	neuropeptide Y receptor Y2	Npy2r	1.65	3.58	5.22
NM_001106544	dolichyl-phosphate mannosyltransferase polypeptide 1, catalytic subunit	Dpm1	1.09	3.04	5.22
NM_183334	lactate dehydrogenase A-like 6B	Ldhal6b	1.25	2.28	5.21
NM_053753	C-type lectin domain family 4, member f	Clec4f	1.04	2.24	5.21
AW528716	EST	–	0.88	3.32	5.19
NM_022599	synaptojanin 2 binding protein	Synj2bp	0.88	7.00	5.14
BE114170	EST	–	1.18	4.22	5.09
BE115208	EST	–	1.15	3.73	5.04
NM_017265	hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 6	Hsd3b6	0.68	3.84	5.03
BE102060	EST	–	1.56	4.90	5.02
XM_002727806	homeobox B2	LOC100361765	1.50	2.36	5.01
BF391102	EST	–	1.02	3.92	5.00
AI556639	EST	–	1.76	2.47	4.98
BG374590	EST	–	0.86	2.10	4.97
NM_001108398	Hyaluronan and proteoglycan link protein 4	Hapln4	1.80	2.66	4.96
BG375582	EST	–	1.12	2.30	4.96
BE100632	EST	–	1.72	4.77	4.95
XM_001059736	similar to chromosome 9 open reading frame 5	RGD1308958	0.66	3.28	4.90
NM_012540	cytochrome P450, family 1, subfamily a, polypeptide 1	Cyp1a1	0.92	3.18	4.89
BF414994	EST	–	0.94	5.21	4.88
NM_001109363	RAB17, member RAS oncogene family	Rab17	0.93	4.36	4.84
NM_031813	myosin binding protein H	Mybph	0.86	2.49	4.82
NM_001109607	histidine triad nucleotide binding protein 1	Hint1	0.97	4.29	4.80
XM_001067323	Hypothetical protein LOC688535	LOC688535	1.58	2.26	4.77
AI044342	EST	–	1.01	3.95	4.77
NM_001109124	Coiled-coil domain containing 112	Ccdc112	0.70	3.67	4.77
NM_001109024	DnaJ (Hsp40) homolog, subfamily C, member 30	Dnajc30	1.49	2.12	4.77
BF403436	EST	–	1.18	7.28	4.75
NM_001033860	carcinoembryonic antigen-related cell adhesion	Ceacam1	0.72	4.52	4.74

	molecule 1				
NM_001083338	translocase of outer mitochondrial membrane 40 homolog B	Tomm40b	0.83	5.21	4.74
BF393107	EST	–	1.27	2.67	4.72
NM_017180	pleckstrin homology-like domain, family A, member 1	Phlda1	1.32	2.91	4.72
BF409856	EST	–	1.54	4.94	4.70
NM_031538	CD8a molecule	Cd8a	0.98	2.40	4.70
XM_001080252	similar to Complement C5 precursor	RGD1308742	1.15	11.97	4.66
BM385451	EST	–	1.19	3.31	4.63
AI059958	EST	–	1.27	3.34	4.62
NM_001108066	AT rich interactive domain 3A (Bright like)	Arid3a	1.05	3.93	4.62
NM_001025274	splicing factor, arginine/serine-rich 18	Sfrs18	0.43	2.90	4.61
NM_001108550	periostin, osteoblast specific factor	Postn	0.51	5.03	4.60
NM_019345	solute carrier family 12 (sodium/chloride transporters), member 3	Slc12a3	0.53	4.51	4.59
AI113313	EST	–	1.43	7.47	4.55
BE099423	EST	–	1.40	3.54	4.52
NM_181364	thyrotropin releasing hormone receptor 2	Trhr2	1.10	3.08	4.47
AI412543	EST	–	0.95	4.54	4.45
BF284535	EST	–	1.41	4.57	4.44
BE099881	EST	–	0.88	2.34	4.44
NM_022696	Heart and neural crest derivatives expressed 2	Hand2	0.88	4.27	4.42
NM_001105920	alkB, alkylation repair homolog 4	Alkbh4	1.16	2.44	4.42
AI555247	EST	–	1.02	4.64	4.42
BE113549	EST	–	1.28	4.81	4.38
X76129	EST	–	1.66	3.50	4.38
BF408794	EST	–	1.36	5.24	4.37
BE097095	EST	–	0.97	2.71	4.37
NM_001100659	CDC23 (cell division cycle 23)	Cdc23	1.97	4.07	4.36
BF415420	EST	–	0.85	2.37	4.35
NM_173321	hemiferrin, transferrin-like protein	LOC286987	1.03	3.34	4.35
NM_001037357	leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 3-like	Lilrb3l	1.50	5.33	4.32
NM_001109892	fibroblast growth factor receptor 2	Fgfr2	1.26	2.70	4.30
NM_021659	synaptotagmin VII	Syt7	0.69	4.49	4.30
NM_001166020	E030032D13Rik gene	E030032D13Rik	1.67	4.31	4.29
NM_001025115	signal transducing adaptor family member 1	Stap1	1.18	2.43	4.27
NM_001164396	similar to 20-alpha-hydroxysteroid dehydrogenase	RGD1564865	0.69	2.83	4.27
BF405078	EST	–	0.77	2.52	4.27
NM_001079712	lysine (K)-specific demethylase 4D	Kdm4d	1.54	2.38	4.27
AA997287	EST	–	1.62	3.41	4.26
XM_001075943	ER degradation enhancer, mannosidase alpha-like 1	Edem1	1.49	4.15	4.21
BF404359	EST	–	1.98	3.42	4.21
NM_021583	prostaglandin E synthase	Ptges	0.67	3.48	4.21
NM_001101008	Similar to Fgfr1 oncogene partner	LOC683722	1.54	2.58	4.19
AI598405	EST	–	0.41	4.45	4.19
BF391867	EST	–	1.41	2.73	4.18
BE106653	EST	–	1.25	3.88	4.18
NM_031584	solute carrier family 22 (organic cation transporter), member 2	Slc22a2	1.68	3.52	4.15
BE097012	EST	–	1.46	5.57	4.14
AI716045	EST	–	1.36	5.80	4.14
XM_002729840	hypothetical protein LOC100362109	LOC100362109	1.26	3.02	4.14
AI639330	EST	–	1.11	2.43	4.10
XM_001061311	Hypothetical protein LOC681335	LOC681335	1.47	3.25	4.09

NM_001017497	submaxillary gland androgen regulated protein 3A	Smr3a	1.03	2.63	4.08
BF410178	EST	–	1.15	3.81	4.06
NM_134336	neuroigin 3	Nlgn3	1.62	3.28	4.04
AI072279	EST	–	1.10	2.85	4.03
AI113047	EST	–	1.17	3.08	4.03
NM_001127659	methionine-tRNA synthetase	Mars	1.33	3.33	4.01
NM_053914	ST8 alpha-N-acetyl-neuraminide alpha-2,8-sialyltransferase 4	St8sia4	1.35	4.05	4.01
NM_001108930	pre-B lymphocyte 3	Vpreb3	1.26	3.88	4.01
BI850153	EST	–	1.26	8.13	4.00
BI298075	EST	–	0.72	3.51	3.98
XM_001076876	obscurin, cytoskeletal calmodulin and titin-interacting RhoGEF	Obscn	1.13	3.66	3.98
NM_013217	myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 4	Mllt4	0.64	2.04	3.97
BF564328	EST	–	0.77	9.47	3.97
BG380223	EST	–	1.30	2.79	3.97
AI030975	EST	–	1.61	2.59	3.96
NM_023026	ArfGAP with GTPase domain, ankyrin repeat and PH domain 2	Agap2	0.95	4.54	3.96
BF390840	EST	–	1.33	3.07	3.94
AI044004	EST	–	1.39	2.00	3.94
AI555848	EST	–	1.31	2.08	3.94
NM_173127	protease, serine, 3	Prss3	0.84	3.81	3.93
BF405026	EST	–	1.23	3.13	3.90
BM389444	EST	–	1.90	5.68	3.89
XM_001054773	ankyrin repeat domain 60	Ankrd60	1.93	3.14	3.89
BF288845	EST	–	1.86	4.18	3.88
NM_019166	synaptogyrin 1	Syngn1	1.57	2.83	3.88
NM_001191916	Bromodomain adjacent to zinc finger domain, 1B	Baz1b	1.00	4.98	3.86
NM_012996	oxytocin, prepropeptide	Oxt	0.91	4.78	3.86
NM_001107194	pogo transposable element with KRAB domain	Pogk	1.72	3.27	3.83
AI548379	EST	–	1.07	2.55	3.83
AI235868	EST	–	0.89	2.70	3.81
NM_001100514	centrosomal protein 76kDa	Cep76	0.67	3.85	3.80
NM_019183	actin, alpha, cardiac muscle 1	Actc1	1.20	2.40	3.79
NM_001042579	unc-13 homolog B	Unc13b	1.21	3.24	3.79
NM_001107153	ubiquitin specific peptidase 30	Usp30	0.93	3.11	3.78
BE098450	EST	–	1.00	2.85	3.78
NM_001082572	Tripartite motif-containing 21	Trim21	1.89	3.20	3.77
NM_001107867	ATPase, H transporting, lysosomal V1 subunit B1	Atp6v1b1	0.94	3.05	3.77
NM_138975	WW domain binding protein 2	Wbp2	1.65	3.66	3.77
NM_001039338	Zinc finger, DHHC-type containing 5	Zdhhc5	0.79	3.64	3.76
BI302053	EST	–	1.35	2.80	3.76
NM_031713	leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 3	Lilrb3	0.97	10.23	3.76
XM_002725948	hypothetical protein LOC100360513	LOC100360513	1.17	2.59	3.76
BF284722	EST	–	1.70	6.20	3.75
AW915304	EST	–	1.39	3.76	3.74
NM_001100852	zinc finger protein 746	Znf746	0.72	3.93	3.74
BF397286	EST	–	1.00	2.61	3.73
BF410377	EST	–	1.83	2.24	3.73
NM_001005877	free fatty acid receptor 2	Ffar2	1.09	5.88	3.73
BI282060	EST	–	1.21	2.19	3.73
NM_031804	cytokine inducible SH2-containing protein	Cish	1.84	2.64	3.72
BE105421	EST	–	0.61	2.84	3.72

AA956474	EST	–	0.68	3.47	3.72
NM_053789	interleukin 17B	Il17b	1.69	3.07	3.70
AI043832	EST	–	0.97	2.44	3.69
NM_001191725	Myotubularin related protein 1	Mtmr1	1.64	2.10	3.69
NM_017102	solute carrier family 2 (facilitated glucose transporter), member 3	Slc2a3	1.98	4.42	3.68
AI044912	EST	–	1.24	2.96	3.68
BF396101	EST	–	1.00	3.86	3.68
NM_001110099	ret proto-oncogene	Ret	1.67	4.48	3.67
NM_147144	cancer susceptibility candidate 3	Casc3	1.26	2.82	3.67
AA963295	EST	–	1.13	4.60	3.65
NM_031739	potassium voltage-gated channel, Shal-related subfamily, member 3	Kcnd3	0.96	3.75	3.64
NM_001191692	leucine rich repeat containing 16A	Lrrc16a	1.18	2.30	3.63
AA946053	EST	–	0.95	3.32	3.63
NM_172157	AT rich interactive domain 1B (Swi1 like)	Arid1b	1.56	5.29	3.61
XM_001075996	ring finger protein 207	Rnf207	0.93	3.31	3.60
AI555088	EST	–	1.00	2.37	3.60
NM_001107087	organic solute transporter alpha	Ostalpha	1.44	2.76	3.59
BF399291	EST	–	1.50	3.38	3.59
NM_170667	Relaxin 3	Rln3	1.06	3.19	3.58
NM_023103	murinoglobulin 1	Mug1	1.80	3.00	3.57
AW527295	EST	–	1.04	2.01	3.56
BF401303	EST	–	1.90	4.34	3.56
AI030318	EST	–	1.29	2.60	3.55
NM_001102403	seminal vesicle secretory protein 6	Svs6	1.75	3.15	3.52
NM_001011947	retinoic acid induced 14	Rai14	1.35	2.11	3.51
BE114399	EST	–	0.88	2.94	3.51
NM_012531	catechol-O-methyltransferase	Comt	0.97	3.86	3.50
XM_002726882	rCG49027-like	LOC100362836	1.71	3.52	3.50
BF396390	EST	–	0.99	3.47	3.50
NM_022684	BH3 interacting domain death agonist	Bid	1.02	2.80	3.48
BF415758	EST	–	0.86	3.96	3.47
NM_001191100	Similar to RNA binding motif protein 24	LOC690139	1.45	3.04	3.46
BE098827	EST	–	0.87	3.25	3.46
NM_031533	UDP glycosyltransferase 2 family, polypeptide B	Ugt2b	1.26	3.18	3.46
XM_001055673	Similar to SWI/SNF-related matrix-associated actin-dependent regulator of chromatin c2	LOC685179	1.53	2.29	3.44
BI296600	EST	–	1.63	2.20	3.43
NM_001108382	coiled-coil domain containing 25	Ccdc25	0.90	2.40	3.43
AFFX-TrpnX-3	EST	–	1.43	5.29	3.42
BF405278	EST	–	0.88	2.57	3.42
U39571	EST	–	0.80	2.47	3.42
BF401158	EST	–	0.71	2.19	3.41
NM_001106207	Myc target 1	Myct1	1.09	2.77	3.41
BF405941	EST	–	1.06	4.26	3.41
NM_053953	interleukin 1 receptor, type II	Il1r2	0.85	2.57	3.40
XM_001069222	Similar to KIAA1549 protein	RGD1306271	1.30	3.23	3.38
NM_012524	CCAAT/enhancer binding protein (C/EBP), alpha	Cebpa	0.95	2.63	3.36
BF386426	EST	–	0.99	2.50	3.35
NM_017254	5-hydroxytryptamine (serotonin) receptor 2A	Htr2a	1.13	3.26	3.34
BF410005	EST	–	0.52	4.13	3.33
NM_001106319	CysteinyI-tRNA synthetase	Cars	1.15	4.23	3.31
NM_017093	v-akt murine thymoma viral oncogene homolog 2	Akt2	1.07	2.91	3.30
AI072237	EST	–	1.16	2.27	3.29

NM_001107804	Molybdenum cofactor synthesis 3	Mocs3	1.12	3.19	3.28
NM_153621	disabled homolog 1	Dab1	0.45	2.85	3.28
NM_001008831	RT1 class II, locus Ba	RT1-Ba	1.25	2.26	3.27
NM_031538	CD8a molecule	Cd8a	0.99	4.37	3.27
AI144892	EST	–	1.32	4.44	3.26
NM_012849	gastrin	Gast	1.10	3.30	3.26
NR_026689	hypothetical protein LOC686120	LOC686120	1.55	2.75	3.26
NM_001008297	similar to DNA segment, Chr 14, ERATO Doi 449, expressed	RGD1305689	1.01	3.20	3.25
NM_017267	Translocase of inner mitochondrial membrane 44 homolog	Timm44	1.26	5.17	3.25
BG373668	EST	–	1.86	4.09	3.25
XR_005459	similar to inter-alpha-inhibitor H2 chain	LOC498793	1.58	2.06	3.25
NM_139085	cystatin 11	Cst11	1.26	3.27	3.24
NM_012715	adrenomedullin	Adm	1.22	2.01	3.24
AI555339	EST	–	1.48	2.29	3.23
NM_001108463	radial spokehead-like 2	Rshl2	1.75	2.87	3.22
AI638966	EST	–	0.78	2.71	3.22
NM_173135	amiloride-sensitive cation channel 3	Accn3	0.91	2.06	3.21
NM_001010970	amylase, alpha 1A	Amy1a	0.39	23.87	3.21
NM_001008758	keratin 34	Krt34	1.30	2.57	3.21
NM_001106506	nephronophthisis 1 (juvenile) homolog	Nphp1	1.50	2.07	3.20
BF404936	EST	–	1.03	2.73	3.20
NM_001013159	formin binding protein 4	Fnbp4	1.77	2.27	3.20
NM_001127652	creatine kinase, mitochondrial 2, sarcomeric	Ckmt2	1.30	2.36	3.19
BF288361	EST	–	0.93	2.41	3.19
NM_021659	synaptotagmin VII	Syt7	1.04	2.92	3.19
AI555284	EST	–	1.36	4.89	3.18
NM_001012355	Sumo1/sentrin/SMT3 specific peptidase 8	Senp8	0.95	2.42	3.18
NM_021584	doublecortin-like kinase 1	Delk1	1.11	3.98	3.18
BF410736	EST	–	0.67	3.05	3.18
NM_207605	SH2 domain protein 2A	Sh2d2a	1.19	2.84	3.17
BE110921	EST	–	1.32	2.04	3.17
BM385125	EST	–	0.73	3.06	3.17
BG378715	EST	–	0.94	2.36	3.17
NM_134327	Cd69 molecule	Cd69	1.23	2.16	3.16
AW532385	EST	–	0.82	2.97	3.16
BI273836	EST	–	1.75	2.03	3.16
NM_053291	phosphoglycerate kinase 1	Pgk1	0.56	4.09	3.15
NM_001012181	E74-like factor 2	Elf2	1.29	2.19	3.15
BF291108	EST	–	1.82	2.74	3.14
NM_012568	glycine receptor, alpha 2	Gla2	1.84	2.42	3.14
NM_022393	C-type lectin domain family 10, member A	Clec10a	1.39	3.45	3.13
AA955985	EST	–	1.12	2.45	3.13
NM_001108238	ankyrin repeat domain 12	Ankrd12	0.86	3.47	3.12
BI292034	EST	–	1.43	2.46	3.12
NR_002151	RT1 class I, locus T24, gene 2	RT1-T24-2	1.43	5.16	3.11
XM_001069306	zinc finger CCCH-type, antiviral 1-like	Zc3hav11	1.52	2.83	3.11
NM_001105727	proteasome (prosome, macropain) subunit, beta type 5	Psmb5	1.00	3.33	3.11
BF401649	EST	–	1.38	2.94	3.10
BF419542	EST	–	1.57	3.79	3.09
NM_001108880	BARX homeobox 1	Barx1	1.17	3.72	3.09
NM_001009497	killer cell lectin-like receptor, subfamily A, member 17	Klra17	0.51	3.19	3.09

BF400860	EST	–	1.14	2.48	3.09
AI013206	EST	–	1.69	2.81	3.08
NM_173300	olfactory receptor 1271	Olr1271	1.68	4.11	3.08
NM_001010970	amylase, alpha 1A	Amy1a	1.20	8.40	3.08
NM_013193	neurofibromin 2	Nf2	1.61	3.08	3.08
AI072079	EST	–	1.67	2.34	3.07
XM_001078877	genetic suppressor element 1	Gse1	0.95	2.79	3.07
BM389874	EST	–	1.38	3.03	3.06
NM_001134341	ischemia related factor NYW-1	Nyw1	0.77	2.76	3.06
NM_024401	advillin	Avil	1.76	2.74	3.06
NM_001009494	killer cell lectin-like receptor, subfamily A, member 5	Klra5	1.61	2.31	3.05
NM_022933	chromodomain helicase DNA binding protein 8	Chd8	1.16	5.84	3.04
AI169247	EST	–	0.95	2.45	3.04
AW433964	EST	–	0.90	2.16	3.03
NM_153621	Disabled homolog 1	Dab1	0.91	2.61	3.02
BF394624	EST	–	0.70	2.04	3.02
NM_001173434	hemochromatosis	Hfe	1.16	2.15	3.01
AW534610	EST	–	0.77	3.07	3.01
NM_031545	natriuretic peptide precursor B	Nppb	0.51	2.36	3.01
BM391972	EST	–	1.76	3.92	3.00
NM_173130	vomer nasal 2 receptor, 27	Vom2r27	1.25	3.70	3.00
BF396142	EST	–	1.33	2.53	2.99
NM_031140	vimentin	Vim	1.65	2.82	2.99
NM_001033958	alpha-2u globulin PGCL4	Obp3	1.68	2.79	2.99
NM_022253	Csk binding protein	Cbp	0.78	3.54	2.98
NM_024160	cytochrome b-245, alpha polypeptide	Cyba	0.93	3.21	2.98
XM_002725069	hypothetical protein LOC100364843	LOC100364843	1.11	3.10	2.97
NM_053871	signal recognition particle 54a	Srp54a	1.48	2.79	2.96
NM_001109530	SAM pointed domain containing ets transcription factor	Spdef	1.22	2.09	2.96
AI231651	EST	–	0.97	2.31	2.96
XM_001078167	Similar to CG9646-PA	RGD1304694	1.12	4.83	2.95
AA963477	EST	–	0.97	2.33	2.94
NM_001107291	inter-alpha trypsin inhibitor, heavy chain 1	Itih1	0.88	3.37	2.94
BF419668	EST	–	1.17	2.18	2.94
XM_001061208	similar to ribosomal protein S12	LOC680988	1.58	2.03	2.94
BF290383	EST	–	1.15	3.06	2.94
NM_031141	paired box 8	Pax8	1.27	2.30	2.94
NM_199107	Glycosyltransferase-like 1B	Gylt1b	1.19	2.22	2.93
NM_001109070	golgi transport 1 homolog A	Golt1a	1.60	2.23	2.93
NM_139041	mucin 13, cell surface associated	Muc13	1.03	2.36	2.92
BF393928	EST	–	0.99	2.42	2.92
NM_181365	Kv channel interacting protein 4	Kcnip4	1.78	3.43	2.92
NM_207593	prostatic steroid-binding protein C2	Psbpc2	0.75	2.80	2.91
NM_053312	dihydroipoamide branched chain transacylase E2	Dbt	0.86	2.14	2.91
NM_001009624	Spindle and kinetochore associated complex subunit 2	Ska2	1.15	4.34	2.90
BE101809	EST	–	0.94	2.30	2.89
XR_007893	similar to TICAM-1	LOC363328	1.07	2.31	2.88
NM_019321	mast cell protease 4	Mcpt4	1.06	6.31	2.88
AI072437	EST	–	0.65	2.84	2.88
NM_013097	deoxyribonuclease I	Dnase1	1.40	3.95	2.87
BF394493	EST	–	1.13	2.37	2.87
NM_001034931	polycystic kidney and hepatic disease 1-like 1	Pkhd111	0.84	2.20	2.87

NM_022235	potassium voltage-gated channel, Isk-related subfamily, gene 3	Kcne3	0.92	2.91	2.87
AA859889	EST	–	1.93	3.38	2.86
AI071526	EST	–	1.76	2.37	2.86
NM_138504	oxidative stress induced growth inhibitor 1	Osgin1	1.13	3.08	2.85
NM_001025411	DnaJ (Hsp40) homolog, subfamily A, member 4	Dnaja4	1.18	3.96	2.85
AI229471	EST	–	1.76	2.08	2.85
XR_009042	similar to RGD, leucine-rich repeat, tropomodulin and proline-rich containing protein	RGD1562390	1.85	2.37	2.84
AI763894	EST	–	1.00	2.13	2.84
NM_053640	leukotriene B4 receptor 2	Ltb4r2	1.35	2.89	2.84
XM_001055013	leucine-rich repeats and immunoglobulin-like domains 3	Lrig3	1.13	2.15	2.83
BE108754	EST	–	1.10	3.07	2.83
NM_001108559	PRP3 pre-mRNA processing factor 3 homolog	Prpf3	1.36	5.13	2.82
AI045029	EST	–	1.39	3.17	2.82
NM_020097	exostoses (multiple)-like 3	Extl3	0.97	2.02	2.82
NM_001109541	DnaJ (Hsp40) homolog, subfamily B, member 2	Dnajb2	1.03	5.85	2.81
NM_053644	cadherin 23 (otocadherin)	Cdh23	0.80	2.47	2.80
NM_017047	solute carrier family 10 (sodium/bile acid cotransporter family), member 1	Slc10a1	0.72	2.42	2.80
XR_006158	Similar to Myeloid/lymphoid or mixed-lineage leukemia protein 3 homolog (Histone-lysine N-methyltransferase, H3 lysine-4 specific MLL3)	LOC502710	0.93	2.38	2.79
NM_031588	neuregulin 1	Nrg1	0.82	2.44	2.79
AI578317	EST	–	1.49	2.10	2.79
BF400683	EST	–	0.84	5.55	2.79
AI072041	EST	–	0.95	4.60	2.78
AW921479	EST	–	1.73	2.45	2.77
BM390376	EST	–	0.98	3.45	2.76
BF403678	EST	–	1.12	3.00	2.76
AW142608	EST	–	1.01	2.76	2.75
AA957926	EST	–	1.02	3.08	2.74
NM_001126303	ubiquitin specific protease 43	Usp43_predicted	0.84	2.70	2.74
NM_212459	ADP-ribosylation factor-like 9	Arl9	0.36	2.13	2.74
AI045025	EST	–	0.68	2.28	2.74
AI071595	EST	–	0.88	2.49	2.73
BF394024	EST	–	1.24	2.80	2.72
NM_001130551	similar to secreted Ly6/uPAR related protein 2	RGD1308195	0.85	2.33	2.72
NM_017078	cholinergic receptor, nicotinic, alpha 5	Chrna5	1.18	2.19	2.72
NM_053613	reticulon 4 receptor	Rtn4r	0.99	2.88	2.72
NM_053724	glycine receptor, alpha 3	Glr3	1.15	2.05	2.71
AI600230	EST	–	1.17	3.77	2.71
BE113971	EST	–	0.96	2.44	2.71
BF410456	EST	–	1.17	2.49	2.70
AW530928	EST	–	1.02	4.05	2.70
NM_001002827	Notch homolog 4	Notch4	1.10	4.25	2.70
NM_001108652	similar to hypothetical protein DKFZp761D0211	RGD1306151	1.18	2.73	2.70
AT005664	EST	–	1.03	2.20	2.69
AW251681	EST	–	1.16	3.92	2.69
NM_012511	ATPase, Cu <sup>++</sup> transporting, beta polypeptide	Atp7b	1.34	2.84	2.69
NM_001011892	serine (or cysteine) peptidase inhibitor, clade F, member 2	Serpinf2	1.10	2.20	2.69
AI103351	EST	–	0.43	2.22	2.68
NM_001107823	growth factor independent 1B transcription repressor	Gfi1b	0.95	2.44	2.68
BF412962	EST	–	1.60	4.74	2.67



BF400135	EST	–	0.70	2.68	2.67
BE097058	EST	–	0.74	3.03	2.67
XM_002725301	TAF3 RNA polymerase II, TATA box binding protein (TBP)-associated factor	LOC100360100	1.52	3.62	2.67
NM_019359	Calponin 3, acidic	Cnn3	1.11	3.64	2.67
AI232217	EST	–	1.04	2.65	2.67
NM_012553	chymotrypsin-like elastase family, member 2A	Cela2a	0.67	2.63	2.67
NM_001004022	keratin 15	Krt15	1.17	2.17	2.67
NM_001025641	pregnancy-specific glycoprotein 29	Psg29	1.67	4.16	2.67
AI385279	EST	–	1.32	2.67	2.66
BF387484	EST	–	0.93	2.48	2.66
BF404369	EST	–	0.99	2.13	2.66
NM_001013065	gametogenetin	Ggn	1.09	2.70	2.65
NM_138861	Prolactin family 2, subfamily b, member 1	Prl2b1	0.89	2.80	2.65
NM_012944	dopamine receptor D4	Drd4	0.46	4.21	2.64
AW917217	EST	–	0.95	4.17	2.64
XM_001070946	similar to class-alpha glutathione S-transferase	RGD1562107	1.36	2.26	2.64
NM_012813	ST8 alpha-N-acetyl-neuraminidase alpha-2,8-sialyltransferase 1	St8sia1	1.02	3.22	2.64
XM_002727399	paired immunoglobulin-like type 2 receptor alpha	Pilra	1.23	2.06	2.63
NM_001013098	dehydrogenase/reductase (SDR family) member 7	Dhrs7	0.57	2.42	2.63
BE120674	EST	–	0.96	2.31	2.63
NM_001017480	homeo box B7	Hoxb7	1.18	2.74	2.62
NM_001030029	COMM domain containing 7	Comm7	1.00	2.21	2.62
NM_019292	carbonic anhydrase 3	Car3	1.15	2.01	2.61
NM_001106972	Similar to CG4768-PA	RGD1309748	1.49	2.70	2.61
NM_001108271	LOC360508	RGD1306625	1.72	2.24	2.61
BF404505	EST	–	1.00	2.79	2.60
BE106136	EST	–	0.99	2.19	2.60
NM_080768	tachykinin receptor 2	Tacr2	1.42	2.37	2.60
NM_022631	wingless-type MMTV integration site family, member 5A	Wnt5a	1.72	2.38	2.60
NM_001107422	THAP domain containing 11	Thap11	1.30	2.19	2.60
BI296703	EST	–	1.38	2.89	2.59
NM_001135173	T-kininogenase	Klk1c10	0.75	2.43	2.59
NM_019255	calcium channel, voltage-dependent, gamma subunit 1	Cacng1	1.90	4.50	2.59
AI235078	EST	–	1.17	2.16	2.59
BF413876	EST	–	0.77	2.35	2.58
NM_001135157	myosin, heavy chain 2, skeletal muscle	Myh2	1.49	3.72	2.57
XM_002727932	F-box and WD repeat domain containing 7-like	LOC100361588	0.93	3.01	2.56
BF403846	EST	–	0.95	2.98	2.56
BI300244	EST	–	0.87	2.97	2.56
AI069951	EST	–	1.08	2.06	2.55
NM_053822	S100 calcium binding protein A8	S100a8	1.74	2.01	2.55
NM_001004131	Keratin 24	Krt24	1.67	2.61	2.55
NM_022239	neuromedin U	Nmu	0.60	2.89	2.55
NM_053706	doublesex and mab-3 related transcription factor 1	Dmrt1	0.77	2.88	2.55
NM_021586	latent transforming growth factor beta binding protein 2	Ltbp2	0.88	2.30	2.54
XM_002725935	solute carrier family 25 (mitochondrial carrier; adenine nucleotide translocator), member 31-like	LOC100364813	1.00	2.49	2.54
NM_001106257	myosin binding protein C, fast-type	Mybpc2	1.00	2.53	2.54
AA859673	EST	–	0.89	3.20	2.54
NM_001109556	ovary-specific acidic protein	Osap	1.22	2.11	2.53
BF418099	EST	–	1.20	2.52	2.53

BF405181	EST	–	1.19	2.09	2.53
NM_001106196	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 2 (GalNAc-T2)	Galnt2	0.93	3.57	2.53
NM_020081	CD86 molecule	Cd86	1.23	2.34	2.53
NM_022582	lectin, galactoside-binding, soluble, 7	Lgals7	1.99	3.22	2.53
H33294	EST	–	1.11	2.78	2.53
NM_001034943	solute carrier family 22 (organic anion/urate transporter), member 12	Slc22a12	1.54	2.78	2.53
NM_001106067	glycosyltransferase 25 domain containing 1	Glt25d1	0.74	2.77	2.53
AA893234	EST	–	1.10	2.45	2.52
BF561222	EST	–	1.05	3.86	2.52
AI071607	EST	–	1.72	2.12	2.52
NM_001012047	biotinidase	Btd	1.68	2.83	2.52
NM_001024764	solute carrier family 36 (proton/amino acid symporter), member 3	Slc36a3	1.48	3.57	2.52
NM_053419	synergin, gamma	Synrg	1.49	2.38	2.52
NM_001107033	T-box 2	Tbx2	1.64	2.83	2.52
NM_001111115	brain-enriched guanylate kinase-associated	Begain	1.31	2.24	2.51
NM_012530	creatine kinase, muscle	Ckm	1.31	2.17	2.51
AI071958	EST	–	0.98	2.48	2.50
AI599545	EST	–	1.14	2.34	2.50
XM_001076431	sin3A-binding protein, SAP25	Sap25	1.33	2.28	2.50
NM_145088	mammary cancer associated protein RMT-1	Rmt1	1.68	2.23	2.50
NM_031632	caspase 9, apoptosis-related cysteine peptidase	Casp9	1.60	2.24	2.50
NM_001109912	TSC22 domain family, member 1	Tsc22d1	0.86	3.22	2.49
NM_012596	leptin receptor	Lepr	1.46	2.10	2.49
NM_138874	casein alpha s1	Csn1s1	1.33	2.15	2.49
BF393838	EST	–	1.34	2.48	2.49
NM_001006990	cell adhesion molecule JCAM	LOC304000	1.21	2.89	2.49
NM_012718	androgen regulated 20 kDa protein	Andpro	1.12	3.15	2.48
NM_001106841	Mitochondrial translation optimization 1 homolog	Mto1	1.90	2.15	2.48
NM_138977	prokineticin receptor 1	Prokr1	1.02	2.51	2.48
NM_012879	solute carrier family 2 (facilitated glucose transporter), member 2	Slc2a2	1.16	2.06	2.48
BM391896	EST	–	0.72	2.13	2.47
AA893682	EST	–	1.09	2.46	2.47
BF401574	EST	–	0.82	3.30	2.47
NM_001107133	GRB10 interacting GYF protein 1	Gigyf1	1.16	2.24	2.46
NM_001014008	asporin	Aspn	1.08	3.61	2.46
AI058733	EST	–	1.38	2.87	2.46
NM_080581	ATP-binding cassette, sub-family C (CFTR/MRP), member 3	Abcc3	0.95	2.37	2.45
AA943800	EST	–	1.14	2.95	2.44
NM_138512	cytochrome P450, family 2, subfamily c, polypeptide 22	Cyp2c22	1.30	2.09	2.43
NM_001191567	Golgi integral membrane protein 4	Golim4	0.81	2.53	2.42
XM_001070024	similar to Cystatin S precursor (LM protein)	LOC100365949	1.57	3.95	2.42
NM_001108396	ARP8 actin-related protein 8 homolog	Actr8	0.85	2.68	2.42
NM_001191609	laminin, alpha 5	Lama5	0.94	2.09	2.42
NM_017288	sodium channel, voltage-gated, type I, beta	Scn1b	0.89	2.08	2.42
BF404398	EST	–	1.59	2.05	2.42
BF390565	EST	–	1.15	2.30	2.42
AA894070	EST	–	0.69	2.40	2.42
AA955079	EST	–	1.60	2.42	2.41
BE106256	EST	–	0.92	2.71	2.41
AA875586	EST	–	1.02	2.28	2.41

NM_001108137	membrane frizzled-related protein	Mfrp	1.15	2.60	2.41
NM_001025065	angiotensin-like 3	Angptl3	0.98	3.08	2.41
NM_001013904	cytochrome P450, family 2, subfamily c, polypeptide 6	Cyp2c6	1.67	3.07	2.40
BF401148	EST	–	1.38	3.28	2.40
AA964652	EST	–	0.95	5.15	2.40
AI008432	EST	–	0.87	2.21	2.40
BF401468	EST	–	0.94	2.13	2.40
AA892778	EST	–	0.48	2.50	2.40
NM_053394	Kruppel-like factor 5	Klf5	1.60	2.46	2.39
BF283420	EST	–	0.90	2.21	2.39
BE119143	EST	–	1.02	4.14	2.39
BI296715	EST	–	1.68	2.08	2.39
NM_176074	complement component 6	C6	1.46	2.24	2.39
NM_001137645	FYVE, RhoGEF and PH domain containing 6	Fgd6	1.19	2.30	2.38
AI113302	EST	–	1.17	2.31	2.38
AI103536	EST	–	0.37	2.09	2.37
BF393042	EST	–	1.37	2.72	2.37
NM_023969	lysophosphatidic acid receptor 3	Lpar3	1.45	4.37	2.37
NM_001105776	forkhead box I1	Foxi1	1.09	2.05	2.37
NM_030873	profilin 2	Pfn2	1.02	2.79	2.37
NM_022592	transketolase	Tkt	1.25	2.85	2.36
NM_024395	5-hydroxytryptamine (serotonin) receptor 5B	Htr5b	1.75	2.13	2.36
NM_001024273	cytidine monophosphate-N-acetylneuraminic acid hydroxylase	Cmah	1.17	2.40	2.36
NM_017254	5-hydroxytryptamine (serotonin) receptor 2A	Htr2a	1.14	2.13	2.36
BF416353	EST	–	1.02	3.27	2.35
NM_144751	resection-induced TPI (rs11)	LOC246267	1.12	2.27	2.35
XM_001069443	forkhead box E3	Foxe3	0.84	2.38	2.35
NM_031982	transient receptor potential cation channel, subfamily V, member 1	Trpv1	1.22	2.56	2.35
BE111113	EST	–	1.03	2.09	2.34
NM_019241	gap junction protein, beta 5	Gjb5	1.29	3.71	2.34
NM_012790	dentin sialophosphoprotein	Dspp	1.42	2.03	2.33
AI008973	EST	–	1.39	2.31	2.33
NM_001109474	EST	–	1.26	3.86	2.32
BG376806	EST	–	1.15	2.76	2.32
XM_001077495	Nuclear receptor co-repressor 1	Ncor1	0.69	2.28	2.32
NM_023971	dystrophin related protein 2	Drp2	1.04	2.80	2.32
BI291303	EST	–	1.36	3.24	2.31
XM_001079649	Similar to KIAA2026 protein	RGD1311595	1.05	2.38	2.31
NM_022384	Achaete-scute complex homolog 1	Ascl1	1.51	2.53	2.31
NM_031705	dihydropyrimidinase	Dpys	0.89	2.03	2.31
NM_080580	RAB3D, member RAS oncogene family	Rab3d	0.95	3.17	2.31
NM_053502	ATP-binding cassette, sub-family G (WHITE), member 1	Abcg1	1.27	2.58	2.31
AW531919	EST	–	1.13	3.67	2.31
NM_017306	dodecenoyl-Coenzyme A delta isomerase (3,2 trans-enoyl-Coenzyme A isomerase)	Dci	1.01	2.92	2.30
NM_001013064	F-box protein 17	Fbxo17	0.52	2.47	2.30
BF400669	EST	–	1.11	2.02	2.30
NM_031028	gamma-aminobutyric acid (GABA) B receptor 1	Gabbr1	1.02	2.08	2.30
NM_031538	CD8a molecule	Cd8a	1.48	2.04	2.29
BF388435	EST	–	1.84	3.28	2.29
NM_001106670	complement component 8, alpha polypeptide	C8a	1.14	2.07	2.28
NM_001106711	Centromere protein A	Cenpa	1.80	3.41	2.28

BF398910	EST	–	0.98	2.36	2.28
NM_138918	synovial sarcoma translocation gene on chromosome 18-like 1	Ss18l1	1.30	3.80	2.27
NM_012877	sodium channel, voltage-gated, type II, beta	Scn2b	1.02	3.32	2.27
NM_001130510	DnaJ (Hsp40) homolog, subfamily B, member 7	Dnajb7	1.40	4.09	2.27
NM_022859	cysteine-rich secretory protein 1	Crisp1	1.19	2.36	2.26
BF561222	EST	–	1.06	3.56	2.25
AA964735	EST	–	1.52	2.05	2.25
NM_001012209	ribonuclease, RNase A family, 12 (non-active)	Rnase12	1.02	2.40	2.25
AI172180	EST	–	0.70	2.02	2.25
AW530665	EST	–	1.85	3.41	2.24
BF418127	EST	–	0.87	3.95	2.24
NM_032070	high mobility group AT-hook 2	Hmga2	0.89	2.42	2.24
NM_001134595	solute carrier family 26, member 10	Slc26a10	0.88	2.10	2.24
NM_001110345	ubiquitin-conjugating enzyme E2 variant 1	Ube2v1	0.93	3.16	2.24
NM_001107480	striatin, calmodulin binding protein 4	Strn4	1.74	4.79	2.23
NM_001015025	serine/threonine kinase 38	Stk38	0.86	2.45	2.23
NM_001191665	Rab interacting lysosomal protein-like 1	Rilpl1	1.21	2.52	2.23
NM_001106103	osteoglycin	Ogn	0.97	3.88	2.23
XM_001060972	similar to RIKEN cDNA 1500015O10	RGD1305645	1.22	2.01	2.23
BF412617	EST	–	0.85	2.23	2.22
NM_031558	steroidogenic acute regulatory protein	Star	1.70	2.07	2.22
BF288144	EST	–	1.09	2.42	2.22
BF553125	EST	–	0.74	2.57	2.22
NM_001134792	similar to complement factor H-related protein	RGD1564614	1.42	2.33	2.22
NM_001159493	glutamyl-tRNA synthetase 2 mitochondrial (putative)	Ears2	0.87	2.03	2.22
NM_001014178	family with sequence similarity 69, member B	Fam69b	1.45	3.70	2.22
AW920931	EST	–	1.78	2.57	2.22
NM_019302	v-crk sarcoma virus CT10 oncogene homolog	Crk	1.13	2.71	2.21
AI103612	EST	–	1.68	2.32	2.21
XM_002730008	translocating chain-associating membrane protein 2	LOC100361830	1.56	2.36	2.20
BF411180	EST	–	1.18	2.22	2.20
AA943548	EST	–	1.01	2.10	2.20
AI548331	EST	–	1.26	2.87	2.20
NM_024349	adenylate kinase 1	Ak1	0.79	2.01	2.19
AI548652	EST	–	0.93	2.06	2.19
NM_013028	Short stature homeobox 2	Shox2	1.99	3.78	2.19
NM_001106286	lymphatic vessel endothelial hyaluronan receptor 1	Lyve1	0.98	2.15	2.19
NM_001107250	Zinc finger protein 503	Znf503	0.87	2.02	2.18
NM_001008759	keratin complex 1, acidic, gene 5	Krt1-5	1.16	3.78	2.18
NM_001170466	mast cell peptidase 3	Mcpt3	1.01	2.23	2.18
NM_031828	potassium large conductance calcium-activated channel, subfamily M, alpha member 1	Kenma1	1.03	2.29	2.18
NM_021670	bone morphogenetic protein 15	Bmp15	0.98	4.16	2.17
NM_001106595	SET and MYND domain containing 1	Smyd1	1.00	2.16	2.17
AI556345	EST	–	1.09	2.18	2.17
NM_031017	cAMP responsive element binding protein 1	Creb1	1.21	2.01	2.17
AW251649	EST	–	0.73	2.26	2.16
NM_001107780	hydroxyacid oxidase (glycolate oxidase) 1	Hao1	1.93	2.54	2.16
NM_001034108	membrane-associated ring finger (C3HC4) 2	March2	0.89	2.37	2.16
BF417885	EST	–	1.15	3.35	2.16
AI137962	EST	–	1.63	2.87	2.16
AA860039	EST	–	1.51	2.34	2.15
NM_001107037	eosinophil peroxidase	Epx	0.85	2.01	2.15

NM_001108251	ubiquilin 2	Ubqln2	0.89	2.32	2.15
NM_001107665	Ubiquitin specific protease 13 (isopeptidase T-3)	Usp13	1.36	2.14	2.15
XM_001062926	aldehyde dehydrogenase 5 family, member A1	Aldh5a1	1.28	2.38	2.14
BG374972	EST	–	1.07	3.54	2.14
NM_001191609	laminin, alpha 5	Lama5	1.69	2.65	2.14
NM_001013998	phosphodiesterase 12	Pde12	1.41	2.05	2.14
BF416590	EST	–	1.11	2.15	2.14
NM_001011896	ATP-binding cassette, sub-family F (GCN20), member 3	Abcf3	1.22	2.42	2.14
AI145781	EST	–	1.03	2.28	2.13
BI299857	EST	–	1.55	2.62	2.13
XM_001053529	similar to CG18437-PA	LOC316460	0.76	3.95	2.13
NM_022643	histone cluster 1, H2ba	Hist1h2ba	1.52	3.41	2.12
BF390528	EST	–	1.14	2.27	2.12
AI011063	EST	–	0.82	2.33	2.12
NM_139262	cathepsin Q	Ctsq	1.46	6.03	2.12
BF394144	EST	–	1.08	2.86	2.12
NM_001106883	apolipoprotein B mRNA editing enzyme, catalytic polypeptide-like 2	Apobec2	1.19	2.07	2.12
BF387379	EST	–	0.73	2.99	2.11
NM_001195599	RGD1559980	RGD1559980	1.03	6.53	2.11
NM_001107611	RNA binding motif protein 20	Rbm20	1.95	2.17	2.10
NM_053810	synaptosomal-associated protein 29	Snap29	1.50	2.94	2.10
BF407971	EST	–	1.66	2.90	2.09
NM_030988	thyroglobulin	Tg	1.47	2.57	2.08
BM385963	EST	–	1.65	2.20	2.08
AW530922	EST	–	0.90	3.67	2.08
NM_138529	neuron navigator 2	Nav2	1.32	3.12	2.08
XM_001060344	consortin, connexin sorting protein	Cnst	1.15	2.36	2.08
AI639197	EST	–	1.39	3.27	2.08
NM_001106899	pleckstrin homology domain containing, family B (evectins) member 2	Plekhb2	1.73	2.46	2.07
NM_017269	protein tyrosine phosphatase, receptor type, J	Ptprj	1.01	2.34	2.07
BF394459	EST	–	1.84	3.07	2.07
AW252555	EST	–	0.98	2.60	2.07
AI412432	EST	–	1.05	2.14	2.07
NM_001145840	glucosidase, alpha; neutral C	Ganc	0.89	2.23	2.06
XM_001056520	Similar to cullin 7	LOC680835	1.09	2.09	2.06
AI600226	EST	–	0.87	2.49	2.06
BG376813	EST	–	1.28	2.34	2.05
AW920132	EST	–	0.92	2.22	2.05
NM_021598	mast cell protease 8	Mcpt8	0.99	5.41	2.05
BF394645	EST	–	0.96	2.74	2.05
XM_238267	RGD1560210	RGD1560210	0.96	2.31	2.05
NM_001122658	family with sequence similarity 178, member B	Fam178b	1.04	2.50	2.04
NM_001077645	protein kinase, cAMP dependent, catalytic, beta	Prkacb	1.08	2.94	2.04
BE112342	EST	–	0.74	2.67	2.04
NM_001024263	protein kinase D3	Prkd3	1.20	2.02	2.04
NM_001008854	RT1 class Ib, locus N1	RT1-N1	0.93	2.33	2.03
NM_001109006	solute carrier family 23 (nucleobase transporters), member 3	Slc23a3	1.14	2.73	2.03
XM_002726189	Sp3 transcription factor	Sp3	1.38	2.34	2.03
NM_134378	sulfatase 1	Sulf1	1.15	2.69	2.03
NM_001106664	tyrosinase-related protein 1	Tyrp1	1.14	2.02	2.03
BE112754	EST	–	1.09	2.86	2.03

NM_031343	solute carrier family 6 (neurotransmitter transporter, noradrenalin), member 2	Slc6a2	1.84	2.39	2.03
NM_022218	chemokine-like receptor 1	Cmklr1	0.90	2.46	2.03
BE119364	EST	–	1.08	2.17	2.02
NM_001107711	Moloney leukemia virus 10	Mov10	1.42	2.47	2.02
AI176826	EST	–	0.90	2.18	2.02
NM_001107965	UDP-Gal:betaGlcNAc beta 1,4-galactosyltransferase, polypeptide 2	B4galt2	1.05	3.05	2.02
NM_001109591	junctional sarcoplasmic reticulum protein 1	Jsrp1	0.94	2.88	2.02
XM_001066061	late cornified envelope 1D	Lce1d	1.36	3.96	2.02
NM_001109422	B-cell CLL/lymphoma 3	Bcl3	1.01	2.30	2.01
BF411772	EST	–	0.79	2.23	2.00
NM_173098	solute carrier family 9 (sodium/hydrogen exchanger), member 4	Slc9a4	1.37	2.17	2.00
NM_030860	myocyte enhancer factor 2D	Mef2d	0.98	3.16	2.00

Abbreviations: DBDE, decabromodiphenyl ether; EST, expressed sequence tag.