

**Supplementary Table 6**

<b>Title</b>	<b>Putative Function</b>	<b>Seq Derived From</b>	<b>Gene Symbol</b>	<b>Av FC</b>	<b>Av SLR</b>
Actin, gamma 2, smooth muscle, enteric	Structure	M22323	Actg2	2.16	1.11
cytoplasmic beta-actin	Structure	V01217	Actx	2.92	1.55
cytoplasmic beta-actin	Structure	AI070848	Actx	2.06	1.05
calreticulin	Signaling	X53363	Calr	2.25	1.17
FGF receptor-1	Signaling	D12498	Fgfr1	4.25	2.09
non-receptor protein kinase protein	Signaling	L34542	Batk	5.37	2.42
protein tyrosine phosphatase	Signaling	S74351		3.82	1.93
Protein kinase C beta	Signaling	X04139	Prkcb1	2.56	1.36
protein tyrosine phosphatase, non-receptor type 16	Signaling	U02553	Ptpn16	2.47	1.30
transforming growth factor beta-1 gene	Signaling	X52498	Tgfb1	3.92	1.97
amyloid precursor-like protein 2	signaling	X77934		2.72	1.45
amyloid precursor-like protein 3	signaling	X77934		5.53	2.47
ras-related protein p23	Signaling	X12535		5.02	2.33
3-hydroxy-3-methylglutaryl-Coenzyme A reductase	Metabolism	X55286	Hmgcr	3.58	1.84
Aldolase A, fructose-bisphosphate	Metabolism	AA924326	Aldoa	4.75	2.25
Complement component 3	Metabolism	X52477	C3	23.09	4.53
component C3	Metabolism	M29866		15.03	3.91
Glyceraldehyde-3-phosphate dehydrogenase	Metabolism	X02231	Gapd	3.33	1.74
Glyceraldehyde-3-phosphate dehydrogenase	Metabolism	X02231	Gapd	3.32	1.73
Glyceraldehyde-3-phosphate dehydrogenase	Metabolism	M17701	Gapd	1.82	0.87
iodothyronine 5'-monodeiodinase	Metabolism	M21476		6.99	2.80

NADH-cytochrome b5 reductase	Metabolism	D00636		10.10	3.34
NADPH-cytochrome P450 reductase	Metabolism	E01524		1.98	0.98
NMDA receptor glutamate-binding subunit	Metabolism	S61973		3.02	1.59
peroxisomal multifunctional enzyme type II	Metabolism	AI101743	Hsd17b4	3.80	1.92
Rattus norvegicus mRNA for mitochondrial dicarboxylate carrier	Metabolism	AJ223355		2.26	1.18
solute carrier family 25 (mitochondrial carrier; citrate transporter) member 1	Metabolism	L12016	Slc25a1	2.16	1.11
stearoyl-CoA desaturase 2	Metabolism	AA875269	Scd2	1.83	0.87
glutathione S-transferase Yc2 subunit	Metabolism	S72506		4.73	2.24
B-alpha gene for class II antigen	Immune response	X07551		2.28	1.19
butyrophilin-like 2 (MHC class II associated)	Immune response	K02815	Btnl2	2.42	1.28
MHC class II-like beta chain	Immune response	U31599		1.99	1.00
polymeric immunoglobulin receptor AATTAA-containing	Immune response	L14002		2.54	1.35
polymeric immunoglobulin receptor AATTAA-containing	Immune response	L14004		6.54	2.71
Rat mRNA for leucocyte-common antigen (L-CA)	Immune response	M10072		4.29	2.10
thymocyte L-CA	Immune response	K03039		2.96	1.57
hippocampal cholinergic neurostimulating peptide	?	E05646		1.92	0.94
ESTs	?	AA685903		2.62	1.39

ESTs	?	U31866		5.24	2.39
polypyrimidine tract binding protein	?	X74565		7.01	2.81
prostatain subunit c3	?	J00772		523.53	9.03
sulfated glycoprotein-1	?	S81353		4.25	2.09
afadin	?	D90401	AF-6	1.91	0.93
betaine-homocysteine methyltransferase		AF038870	Bhmt	24.76	4.63
caveolin		Z46614		3.10	1.63
CD24 antigen		AI171462	Cd24	9.20	3.20
CD24 antigen		U49062	Cd24	8.97	3.16
CD24 antigen		U49062	Cd24	8.58	3.10
coagulation factors X		D21215		5.29	2.40
dolichol-phosphate (beta-D) mannosyltransferase 2		AB013359	Dpm2	1.80	0.85
Epithelial membrane protein 1		Z54212	Emp1	6.02	2.59
Importin beta		L38644	Impnb	2.20	1.13
kynureninase (L-kynurenine hydrolase)		U68168	Kynu	4.59	2.20
phosphoribosylpyrophosphate synthetase-associated protein (39 kDa)		D26073	Prpsap1	1.93	0.95
polymerase alpha		M15114		1.80	0.85
R.norvegicus Pan-2 mRNA		X62323		1.76	0.82
Rattus norvegicus CORO1A protein mRNA, complete cds		AA892506		3.40	1.77
RT1 class Ib gene		M31038	RT1Aw2	1.79	0.84
transketolase		AI059508	Tkt	2.48	1.31
-	-	-	-		
Eukaryotic elongation factor 2 kinase	Translation factor	AI059692	Eef2k	3.10	-1.63
Rattus norvegicus initiation factor eIF-2B gamma subunit (eIF-2B gamma) mRNA, complete cds	Translation factor	AI639441		2.70	-1.43
transcription elongation factor B (SIII) polypeptide 2 (18kD, elongin B)	Transcription Factor	L42855	TCEB2	2.55	-1.35
DNA-damage inducible transcript 3	Transcription Factor	U30186	Ddit3	2.69	-1.43

early B-cell factor (olfactory neuronal transcription factor 1)	Transcription Factor	L24051	Ebf	3.05	-1.61
Inhibitor of DNA binding 3, dominant negative helix-loop-helix protein	Transcription Factor	AI171268	Id3	3.06	-1.62
iron-responsive element-binding protein	Transcription Factor	AI176621	Ratireb	3.09	-1.63
Kruppel-like factor 9	Transcription Factor	D12769	Klf9	2.63	-1.40
mesenchyme homeobox 2	Transcription factor	Z17223	Meox2	2.72	-1.44
NF1-X1 protein	Transcription Factor	AB012234	Nfix	3.19	-1.67
nuclear proten 1	Transcription factor	AF014503	Nupr1	3.30	-1.72
nuclear receptor subfamily 2, group F, member 1	Transcription Factor	U10995	Nr2f1	2.73	-1.45
RaIA binding protein 1	Transcription Factor	U82623	Ralbp1	3.03	-1.60
alpha 7A integrin	Structure	X65036	Itga7	3.66	-1.87
dynein, cytoplasmic, light chain 1	Structure	AI009806	Pin	2.72	-1.44
Laminin chain beta 2	Structure	AI104225	Lamb2	3.78	-1.92
Lysosomal associated membrane protein 1 (120 kDa)	Structure	AI233362	Lamp1	2.54	-1.34
Lysosomal-associated membrane protein 2	Structure	M32016	Lamp2	2.87	-1.52
membrane interacting protein of RGS16	Structure	AA891916	Mir16	3.27	-1.71
membrane interacting protein of RGS16	Structure	AA891916	Mir16	3.27	-1.71
microtubule-associated proteins 1A/1B light chain 3	Structure	U05784	MPL3	2.74	-1.46
myosin regulatory light chain	Structure	X54617		2.78	-1.48
myosin regulatory light chain isoform C	Structure	S77900		2.37	-1.24

Myosin heavy chain 11	Structure	X16262	Myh11	2.80	-1.49
Rat agrin mRNA, complete cds	Structure	M64780		3.11	-1.64
Rat agrin mRNA, complete cds	Structure	M64780		2.59	-1.37
Rat mRNA for trans-Golgi network integral membrane protein TGN38	Structure	AA926242		2.79	-1.48
Tenascin X	Structure	U24489	Tnxa	3.83	-1.94
Extracellular matrix protein 2	Structure	U27562	Ecm2	2.82	-1.50
Integrin, beta 1	Structure	A1177366	Itgb1	3.17	-1.67
LIC-2 dynein light intermediate chain 53/55	Structure	U15138	Dncli2	2.92	-1.55
sperm membrane protein (YWK-II)	Structure	M31322	LOC64312	2.69	-1.43
S-100 beta subunit	signaling	S53527		2.76	-1.46
type II cAMP-dependent protein kinase regulatory subunit	signaling	M12492		2.93	-1.55
14-3-3 protein beta-subtype	signaling	AA924084	Ywhab	3.13	-1.65
Activin receptor like kinase 1	signaling	L36088	Acvr11	2.78	-1.48
Adenylyl cyclase 4	signaling	M80633	Adcy4	2.59	-1.37
ADP-ribosylation-like 4	signaling	X77235	Arl4	2.82	-1.50
annexin VI	signaling	X86086	Anxa6	2.68	-1.42
ATPase, Ca <sup>++</sup> transporting, cardiac muscle, slow twitch 2	signaling	AA799276	Atp2a2	2.54	-1.35
BH3 interacting (with BCL2 family) domain, apoptosis agonist	signaling	AA925846	Bid3	15.55	-3.96
cerebellar Ca-binding protein, spot 35 protein	signaling	A1102839	Calb1	3.50	-1.81
cerebellar Ca-binding protein, spot 35 protein	signaling	A1102839	Calb1	2.69	-1.43
cysteine rich protein 61	signaling	AA800784	Cyr61	3.02	-1.59
cysteine-rich protein 2	signaling	D17512	Csrp2	2.53	-1.34

cysteine-sulfinate decarboxylase	signaling	M64755	Csad	3.48	-1.80
Drosophila polarity gene (frizzled) homologue	signaling	L02529	Fzd1	2.97	-1.57
ephrin A1	signaling	AA892417	Efna1	3.00	-1.59
Farnesyltransferase, subunit alpha	signaling	M81225	Fnta	2.61	-1.38
fibulin 5	signaling	AA875033	Fbln5	3.26	-1.70
Follistatin	signaling	AA858520	Fst	4.14	-2.05
Follistatin	signaling	AA858520	Fst	3.44	-1.78
follistatin-related protein precursor	signaling	AA849769	Fstl	3.21	-1.68
G protein-coupled receptor kinase 5	signaling	AA946251	Gprk5	2.99	-1.58
GTP-binding protein	signaling	M17526	RATBPGTPC	3.31	-1.73
GTP-binding protein gamma subunit (Ggamma8)	signaling	L35921		3.05	-1.61
guanylyl cyclase A/atrial natriuretic peptide receptor	signaling	J05677		2.83	-1.50
Heat shock 27 kDa protein	signaling	AA998683	Hsp27	3.77	-1.91
insulin receptor substrate-3	signaling	U93880	Irs3	5.34	-2.42
insulin-like growth factor II	signaling	X17012		3.96	-1.99
Insulin-like growth factor binding protein 2	signaling	J04486	Igfbp2	3.58	-1.84
jagged 1	signaling	L38483	Jag1	3.54	-1.82
jagged 1	signaling	AA900503	Jag1	3.26	-1.71
Phospholipase C , beta4	signaling	L15556	Plcb4	3.27	-1.71
phosphatidylinositol transfer protein, beta	signaling	D21132	Pitpnb	2.65	-1.40
Platelet-derived growth factor receptor alpha	signaling	AI232379	Pdgfra	2.67	-1.42
potassium voltage gated channel, Shal-related family, member 2	signaling	AI230211	Kcnd3	7.21	-2.85

Proprotein convertase subtilisin/kexin type 3 (paired basic amino acid cleaving enzyme, furin, membrane associated receptor protein)	signaling	X55660	Pcsk3	2.34	-1.23
protein phosphatase 1, regulatory (inhibitor) subunit 1A	signaling	J05592	Ppp1r1a	4.50	-2.17
protein phosphatase 1, regulatory (inhibitor) subunit 1A	signaling	J05592	Ppp1r1a	4.01	-2.00
protein tyrosine phosphatase, receptor type, R	signaling	D64050	Ptprr	3.55	-1.83
Rab3B protein	signaling	AA799389	Rab3b	3.61	-1.85
Rattus norvegicus protein tyrosine phosphatase 2E (PTP2E) mRNA, complete cds	signaling	U17971		2.78	-1.47
Rattus norvegicus ryk mRNA for tyrosine kinase-related protein, partial cds	signaling	AA892268		2.84	-1.51
Rattus norvegicus thyroid hormone responsive protein mRNA, complete cds	signaling	U94904		2.85	-1.51
S100A1	signaling	U26356		2.57	-1.36
TGFB inducible early growth response	signaling	AI172476	Tieg	2.97	-1.57
thyroid hormone receptor alpha	signaling	M31174	Thra1	2.95	-1.56
Transforming growth factor beta stimulated clone 22	signaling	L25785	Tgfb1i4	2.98	-1.57
transforming growth factor-beta (TGF-beta) masking protein large subunit	signaling	AI232078	Ltbp1	2.66	-1.41
unr protein	signaling	X52311	unr	2.87	-1.52
upregulated by 1,25-dihydroxyvitamin D-3	signaling	AI014169	Vdup1	2.52	-1.33

v-ral simian leukemia viral oncogene homolog B (ras related)	signaling	L19699	Ralb	2.52	-1.33
phosphoprotein phosphatase 2A beta	signaling	M23591		2.89	-1.53
ATP synthase, H+ transporting, mitochondrial F0 complex, subunit F6	Oxidoredox	X54510	Atp5j	2.44	-1.29
Rat NADH-cytochrome b-5 reductase mRNA, complete cds	Oxidoredox	AI229440		3.09	-1.63
lysyl oxidase	Other	S77494		3.35	-1.74
lysyl oxidase	Other	S66184		3.25	-1.70
Aquaporin 1 (aquaporin channel forming integral protein 28 (CHIP))	Other	X67948	Aqp1	3.15	-1.66
Arg/Abl-interacting protein ArgBP2	Other	AA891194	Argbp2	3.10	-1.63
bcl-2 associated death agonist	Other	AA818072	Bad	2.56	-1.35
calpain 4	Other	U53859	Capn4	2.65	-1.41
calpain 4	Other	U53859	Capn4	2.54	-1.35
carbonic anhydrase IV	Other	S68245		3.54	-1.82
carbonic anhydrase IV	Other	S68245		3.23	-1.69
carboxypeptidase E	Other	L07281		2.96	-1.57
Carboxypeptidase D precursor	Other	U62897	Cpd	2.75	-1.46
CD59 antigen	Other	AA818025	Cd59	3.12	-1.64
clathrin, light polypeptide (Lcb)	Other	M15883	Cltb	2.79	-1.48
dipeptidase 1	Other	L07315	Dpep1	4.14	-2.05
ectonucleotide pyrophosphatase/phosphodiesterase 2	Other	D28560	Enpp2	3.10	-1.63
Heat shock protein 70-1	Other	L16764	Hspa1a	4.44	-2.15
latexin	Other	X76985	Lxn	2.99	-1.58
Lysyl oxidase	Other	AA875582	Lox	3.18	-1.67
metallothionein-2 and metallothionein-1	Other	M11794		2.54	-1.35
MIPP65 protein	Other	AB000098	Mipp65	2.98	-1.57
neuronatin	Other	U08290	Nnat	3.12	-1.64
prothymosin beta 10	Other	M58404	Ptmb10	2.59	-1.37

R.norvegicus hsp70.2 mRNA for heat shock protein 70	Other	Z75029		3.56	-1.83
Rat metallothionein-2 and metallothionein-1 genes, complete cds	Other	AI176456		3.64	-1.86
Rattus norvegicus late gestation lung protein 1 (Lgl1) mRNA, complete cds	Other	AA859581		2.93	-1.55
Retinoblastoma-related gene	Other	AI230602	Rb2	2.77	-1.47
transcobalamin II precursor	Other	AA893702	Tcn2p	2.64	-1.40
Tropoelastin	Other	J04035	Eln	3.03	-1.60
ubiquitin C	Other	D17296	Ubc	2.66	-1.41
phosphatidate phosphohydrolase type 2	Metabolism/lipogenesis	AA818593	Ppap2	3.34	-1.74
adipsin	Metabolism	M92059		3.14	-1.65
aldolase A	Metabolism	M12919		2.64	-1.40
antigen identified by monoclonal antibodies 4F2	Metabolism	AB015433	Mdu1	2.47	-1.31
carboxylesterase 1	Metabolism	AA800851	Ces1	21.55	-4.43
CD36 antigen (collagen type I receptor, thrombospondin receptor)	Metabolism	AA946368	Cd36	3.22	-1.69
CD36 antigen (collagen type I receptor, thrombospondin receptor)	Metabolism	AA925752	Cd36	3.18	-1.67
cytochrome P450, 2b19	Metabolism	K01721	Cyp2b15	3.87	-1.95
enoyl hydratase-like protein, peroxisomal	Metabolism	U08976	Ech1	3.54	-1.82
Flavin-containing monooxygenase 1	Metabolism	M84719	Fmo1	2.91	-1.54
for proteasomal ATPase (SUG1)	Metabolism	AB000491	LOC81827	2.84	-1.51
glutamine synthetase	Metabolism	M91652		2.88	-1.52
guanine deaminase	Metabolism	AA859837	Gda	3.09	-1.63
guanine nucleotide binding protein gamma subunit 11	Metabolism	AA860043	Gng11	3.22	-1.69
guanylate cyclase 1, soluble, alpha 3	Metabolism	AA849036	Gucy1a3	3.04	-1.61

heart fatty acid binding protein	Metabolism	J02773	Fabp3	2.78	-1.47
Lactate dehydrogenase B	Metabolism	U07181	Ldhb	3.09	-1.63
Lactate dehydrogenase B	Metabolism	U07181	Ldhb	2.54	-1.34
Lipoprotein lipase	Metabolism	L03294	Lpl	2.62	-1.39
Peptidylglycine alpha-amidating monooxygenase	Metabolism	AA891068	Pam	2.65	-1.41
peroxisomal membrane protein 2, 22 kDa	Metabolism	X70223	Pxmp2	3.13	-1.65
Phosphoribosylpyrophosphate synthetase, subunit II	Metabolism	X16555	Prps2	3.03	-1.60
phosphatidate phosphohydrolase type 2	Metabolism	AA818593	Ppap2	3.42	-1.77
plasma glutathione peroxidase precursor	Metabolism	AI172411	Gpxp	2.79	-1.48
R.norvegicus mRNA for 2'5' oligoadenylate synthetase	Metabolism	Z18877		3.95	-1.98
Rat mitochondrial 3-hydroxy-3-methylglutaryl-CoA synthase mRNA, complete cds	Metabolism	M33648		3.30	-1.72
Rattus norvegicus mRNA for endothelial receptor for oxidized low-density lipoprotein, complete cds	Metabolism	AI071531		3.68	-1.88
Rattus norvegicus mRNA for endothelial receptor for oxidized low-density lipoprotein, complete cds	Metabolism	AB005900		3.41	-1.77
Solute carrier family 11 member 2 (natural resistance-associated macrophage protein 2)	Metabolism	AF008439	Slc11a2	2.79	-1.48
tricarboxylate carrier	Metabolism	S70011		3.15	-1.65
tricarboxylate carrier	Metabolism	S70011		3.13	-1.65
tricarboxylate carrier-like protein	Metabolism	AA891880	Loc65042	2.60	-1.38
type II hexokinase	Metabolism	D26393		2.81	-1.49
xanthine dehydrogenase	Metabolism	AI172247	Xdh	3.33	-1.74

angiotensin receptor (AT1)	Inflammation	M86912		3.00	-1.59
bradykinin B2 receptor	Inflammation	X80187		5.84	-2.55
granulin	Inflammation	X62322	Grn	2.34	-1.23
Phospholipase D gene 2	Inflammation	AA998338	Plid2	2.68	-1.42
R.norvegicus trg mRNA	Inflammation	X68101		2.49	-1.32
Complement component 4	Immune response	U42719	C4a	2.61	-1.38
T16	Immune response	D89730		3.04	-1.60
Fc receptor, IgG, alpha chain transporter	Immune response	AI180013	Fcgrt	2.52	-1.34
ESTs, Moderately similar to FINC RAT FIBRONECTIN PRECURSOR [R.norvegicus]	Coagulation	AA891769		2.56	-1.36
fibromodulin	Coagulation	X82152	Fmod	3.77	-1.92
Plasminogen activator, tissue	Coagulation	M23697	Plat	2.97	-1.57
platelet endothelial tetraspan antigen-3	Coagulation	AA800005	Cd151	3.24	-1.69
alpha B-crystallin	?	X60351		2.84	-1.50
alpha-crystallin B chain	?	M55534		2.96	-1.57
Cbp/p300-interacting transactivator, with Glu/Asp-rich carboxy-terminal domain, 2	?	AA900476	Cited2	2.54	-1.34
crp-ductin	?	U32681	Crpd	3.94	-1.98
EST	?	AI639205		4.49	-2.17
ESTs	?	AA800840		5.86	-2.55
ESTs	?	AI639101		4.13	-2.04
ESTs	?	AA893244		4.01	-2.00
ESTs	?	AA866444		3.76	-1.91
ESTs	?	AI639185		3.76	-1.91
ESTs	?	AI639402		3.75	-1.91
ESTs	?	AI639162		3.69	-1.88
ESTs	?	AI639294		3.66	-1.87
ESTs	?	H31550		3.64	-1.86

ESTs	?	AI639072		3.62	-1.86
ESTs	?	AA892578		3.62	-1.86
ESTs	?	AA800882		3.58	-1.84
ESTs	?	AA800786		3.58	-1.84
ESTs	?	AA799832		3.57	-1.84
ESTs	?	AA800711		3.55	-1.83
ESTs	?	AA684631		3.53	-1.82
ESTs	?	AA894233		3.51	-1.81
ESTs	?	AA800844		3.48	-1.80
ESTs	?	AA800771		3.47	-1.79
ESTs	?	AA800790		3.47	-1.79
ESTs	?	AA892319		3.44	-1.78
ESTs	?	AA799641		3.42	-1.77
ESTs	?	AA799560		3.41	-1.77
ESTs	?	AA892754		3.38	-1.76
ESTs	?	AA799663		3.35	-1.74
ESTs	?	AA799494		3.35	-1.74
ESTs	?	AA800849		3.35	-1.74
ESTs	?	AA799666		3.34	-1.74
ESTs	?	AA799732		3.32	-1.73
ESTs	?	AI228548		3.26	-1.71
ESTs	?	AA799740		3.21	-1.68
ESTs	?	AI228548		3.21	-1.68
ESTs	?	AA800686		3.21	-1.68
ESTs	?	AA875633		3.18	-1.67

ESTs	?	AA800243	3.18	-1.67
ESTs	?	AA800678	3.16	-1.66
ESTs	?	AA875032	3.13	-1.64
ESTs	?	AA893000	3.11	-1.64
ESTs	?	AI639001	3.08	-1.62
ESTs	?	AA800822	3.08	-1.62
ESTs	?	AA892918	3.06	-1.61
ESTs	?	AA894304	3.06	-1.61
ESTs	?	AA893770	3.04	-1.61
ESTs	?	AA800737	3.04	-1.61
ESTs	?	AA799641	3.04	-1.60
ESTs	?	AA866426	3.01	-1.59
ESTs	?	AA859688	3.00	-1.58
ESTs	?	AA800850	2.99	-1.58
ESTs	?	U75404	2.99	-1.58
ESTs	?	AI638989	2.98	-1.58
ESTs	?	H31144	2.97	-1.57
ESTs	?	AA799515	2.95	-1.56
ESTs	?	AA893869	2.95	-1.56
ESTs	?	AA685175	2.93	-1.55
ESTs	?	AA892520	2.93	-1.55
ESTs	?	AA799340	2.92	-1.55
ESTs	?	AI639058	2.91	-1.54
ESTs	?	AA859519	2.90	-1.54
ESTs	?	AA894318	2.90	-1.53
ESTs	?	AA866439	2.89	-1.53

ESTs	?	AA891724	2.86	-1.52
ESTs	?	AI007824	2.86	-1.52
ESTs	?	AA859519	2.81	-1.49
ESTs	?	AA800735	2.80	-1.49
ESTs	?	AI175959	2.80	-1.48
ESTs	?	AI639149	2.79	-1.48
ESTs	?	AA892005	2.79	-1.48
ESTs	?	H33629	2.78	-1.48
ESTs	?	AA891681	2.78	-1.48
ESTs	?	AA858617	2.77	-1.47
ESTs	?	AA874889	2.76	-1.47
ESTs	?	AA859877	2.76	-1.46
ESTs	?	H31118	2.74	-1.46
ESTs	?	AA799964	2.74	-1.45
ESTs	?	AA859562	2.74	-1.45
ESTs	?	AA891797	2.73	-1.45
ESTs	?	S55223	2.73	-1.45
ESTs	?	AA858590	2.72	-1.44
ESTs	?	AA800029	2.72	-1.44
ESTs	?	AA892248	2.71	-1.44
ESTs	?	AA799537	2.69	-1.43
ESTs	?	AA891220	2.67	-1.42
ESTs	?	AA875500	2.67	-1.41
ESTs	?	AA892010	2.66	-1.41
ESTs	?	AA800576	2.63	-1.40
ESTs	?	AA859616	2.63	-1.39
ESTs	?	AI639167	2.61	-1.38

ESTs	?	AA799396		2.61	-1.38
ESTs	?	AA892248		2.60	-1.38
ESTs	?	AA800795		2.59	-1.37
ESTs	?	AI639501		2.59	-1.37
ESTs	?	AI007824		2.59	-1.37
ESTs	?	AA891171		2.55	-1.35
ESTs	?	AA894200		2.53	-1.34
ESTs	?	AA893065		2.52	-1.33
ESTs	?	AA875552		2.51	-1.33
ESTs	?	AA800034		2.49	-1.31
ESTs	?	AA891669		2.43	-1.28
ESTs	?	H31887		2.39	-1.26
ESTs	?	AA875523		2.37	-1.24
Lost on transformation 1	?	U72620	Lot1	4.22	-2.08
minoxidil sulfotransferase	?	L19998	PST-1	4.09	-2.03
minoxidil sulfotransferase	?	L19998	PST-1	3.52	-1.82
multiple PDZ domain protein	?	AJ001320	Mpdz	4.08	-2.03
presenilin-2	?	X99267	Psen2	2.53	-1.34
reticulon 1	?	U17604	Rtn1	3.71	-1.89
Rgc32 protein	?	AA859967	Rgc32	3.74	-1.90
Secreted acidic cystein-rich glycoprotein (osteonectin)	?	AA946313	Sparc	2.90	-1.54
Secreted acidic cystein-rich glycoprotein (osteonectin)	?	AA891204	Sparc	2.78	-1.48
Secreted acidic cystein-rich glycoprotein (osteonectin)	?	Y13714	Sparc	2.56	-1.35
sialyltransferase 5	?	AA859911	Siat5	2.92	-1.55
synuclein, gamma	?	X86789	Sncg	2.95	-1.56