Genes expressed in all microglia cultures

GB code	Title	Mean Expression value	Std
AA799406	EST188903	547.9	163.9
AA799452	transaldolase 1	134.1	57.3
AA799501	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex 5	120	26.5
AA799520	DD6G4-2	298.2	54.6
AA799645	FXYD domain-containing ion transport regulator 1	85.7	36.9
AA799672	ribosomal protein L6	354.5	105.1
AA799804	Transcribed sequences	279.9	48.5
AA799899	Transcribed sequence with strong similarity to protein pir:R5RT18 (R.norvegicus) R5RT	854.2	146.6
AA800054	ribosomal protein L19	1154	187
AA800669	protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform	228.4	95.6
AA800814	EST190311 Transcribed sequence with strong similarity to protein ref:NP_076006.1 (M.	199	102.1
AA800849	EST190346	261	84.6
AA800881	EST190378	202.2	88.7
AA801130	growth factor receptor bound protein 2	385.5	78.5
AA817887	profilin	696.5	28.7
AA817892	guanine nucleotide binding protein, beta polypeptide 2	242	34.9
AA817997	ribosomal protein L24	933.2	142.2
AA817997	ribosomal protein L24	725.2	113.1
AA818025	CD59 antigen	263.1	66.8
AA818025	CD59 antigen	131.5	48.4
AA818069	polyubiquitin	1938.4	152.2
AA818152	UI-R-A0-am-b-09-0-UI /clone	565.4	237.9
AA818843	postsynaptic protein Cript	67.4	21.5
AA818858	Peptidylprolyl isomerase A (cyclophilin A)	338.1	128.4
AA819338	signal sequence receptor 4	90.4	19.4
AA849038	ribosomal protein L31	1000.3	68.5
AA849648	ribosomal protein L21	170.3	37.5
AA850885	unknown Glu-Pro dipeptide repeat protein	3336.7	538
AA852004	glutamine synthetase 1	142.4	39.2
AA858590	Transcribed sequences	291.3	107.2
AA858607	Transcribed sequence with strong similarity to protein ref:NP_035934.1 (M.musculus) c	294.8	70
AA859919	UI-R-E0-cg-c-01-0-UI.s1 Rattus norvegicus	330.4	103.8
AA860030	tubulin, beta 5	384	51.4
AA866460	coated vesicle membrane protein	162.1	31.7
AA874999	Transcribed sequence with strong similarity to protein ref:NP_006799.1 (H.sapiens) pro	147.7	11.9
AA875225	GTP-binding protein (G-alpha-i2)	1169.9	333.7
AA875225	GTP-binding protein (G-alpha-i2)	301.9	54.9
AA875233	prosaposin	7750.4	488.1
AA875286	Transcribed sequence with strong similarity to protein ref:NP_059128.1 (H.sapiens) pro	311.7	89.5
AA875523	Transcribed sequence with strong similarity to protein sp:Q64119 (R.norvegicus) MLES_	2738.7	620.7
AA891171	EST194974 Transcribed sequence with moderate similarity to protein ref:NP_004540.1	151.2	29.8
AA891651	EST195454 Transcribed sequence with moderate similarity to protein pir:T00741 (H.sap	120.5	53.4
AA891729	ribosomal protein S27a	2000.1	198.4
AA892086	Transcribed sequence with strong similarity to protein pir:KJHUGU (H.sapiens) KJHUGU	491.8	104.4
AA892123	ribosomal protein L36	172.5	67.3
AA892248	EST196051	360.1	134
AA892248	EST196051	172.8	31.3
AA892332	cystatin C	252.4	69.8
AA892333	alpha-tubulin	1332.6	300.5
AA892367	Transcribed sequence with strong similarity to protein sp:P21531 (R.norvegicus) RL3_R	1264.2	222.5
AA892373	syntenin	403.9	141.8
AA892414	EST196217 Transcribed sequences	137	33.9
AA892462	Transcribed sequence with strong similarity to protein ref:NP_079926.1 (M.musculus) R	522.9	84.7
AA892557	spermidine synthase	148.3	39.2
AA892582	aldehyde dehydrogenase family 3, member A1	698.4	131.3
AA892649	gamma-aminobutyric acid receptor associated protein	690.6	106.7
AA892775	Lysozyme	2217.9	455.1
AA892776	solute carrier family 25 (mitochondrial carrier; adenine nucleotide translocator), member	302.5	100.6
AA892797	phosphoglycerate kinase 1	250.5	97.9
AA892814	calpain, small subunit 1	168.7	41.7
AA892832	fatty acid elongase 1	106.1	32.7
AA892854	EST196657 Transcribed sequences	116.8	31.9
AA893007	Transcribed sequences	94.3	32.6
AA893082	Transcribed sequences	274.7	37.9
AA893217	EST197020 Transcribed sequences	145.1	37.7
AA893246	EST197049 Transcribed sequence with strong similarity to protein sp:P57746 (M.muscu	160.3	30.1
AA893443	RAP1B, member of RAS oncogene family	466.6	89.6
AA893485	EST197288	5738.7	1210.1
AA893870	EST197673	118	37.5
AA894004	Transcribed sequence with weak similarity to protein ref:NP_077377.1 (R.norvegicus)	370.5	73.4
AA894029	Transcribed sequences	166.8	40.6
AA894200	Transcribed sequence with strong similarity to protein sp:Q64119 (R.norvegicus) MLES_	2364	242.4
AA942751	Tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, theta poly	80.1	26.6
AA944007	nucleobindin	319.4	76.1
AA944856	RAP1B, member of RAS oncogene family	310.3	32.5
AA945152	EST200651	2252.6	406.4
AA945611	ribosomal protein L10	743.3	98.4

AA945806	ribosomal protein S14	470.3	95.3
AA946040	EST201539 Transcribed sequence with strong similarity to protein ref:NP_079904.1 (M.	116.5	20.6
AA956114	Transcribed sequence with strong similarity to protein ref:NP_112400.1 (R.norvegicus)	83.7	12.3
AA963674	eukaryotic translation elongation factor 2	646	337
AA964320	Transcribed sequence with strong similarity to protein ref:NP_080337.1 (M.musculus) F	171.4	59.2
AA965264	Ribosomal protein S29	4339	561.2
AA998683	heat shock 27kDa protein 1	187.3	73.3
AB013454	Rat amino acid starvation-induced protein mRNA, 3' end	85.7	17.5
AB022209	ribonucleoprotein F	73.9	25.3
AF001898	aldehyde dehydrogenase family 1, member A1	95.7	52.7
AF009656	hypoxanthine guanine phosphoribosyl transferase	220.6	81.6
AF029357	olfactory receptor-like protein gene	157.4	75.8
AF031642	solute carrier family 14, member 2 kidney urea transporter (UT4)	152.2	53.7
AF048687	UDP-Gal:betaGlcNAc beta 1,4-galactosyltransferase, polypeptide 6	141.3	40.8
AF065438	peptidylprolyl isomerase C-associated protein	183.3	50.5
AF067795	C-terminal binding protein 1	523	117.4
AF083269	actin related protein 2/3 complex, subunit 1B	409	64.7
AF083269	actin related protein 2/3 complex, subunit 1B	316.7	53.1
AF087943	CD14 antigen	430.3	84.2
AF087944	CD14 antigen	453.1	51.8
AF089817	regulator of G-protein signaling 19 interacting protein 1	830.4	336.4
AI007820	heat shock protein 90 beta	622.1	121.4
AI007824	EST202275	197.4	87.5
AI008641	ribosomal protein L22	329.5	153
AI008852	eukaryotic translation elongation factor 1 alpha 1	294.6	59.9
AI008888	cystatin B	795.6	97.3
AI009132	Transcribed sequence with strong similarity to protein ref:NP_079900.1 (M.musculus) F	187.9	35
AI009605	Ras homolog enriched in brain	91.2	21.1
AI010083	peroxiredoxin 1	467.6	91.6
AI010292	EST204743	420	190.3
AI010632	Transcribed sequence with strong similarity to protein sp:Q62931 (R.norvegicus) GS28_	387.3	94.2
AI012805	ribosomal protein L13A	550.3	92.4
AI013627	defender against cell death 1	96.4	15.3
AI013853	ribosomal protein S23	585.7	77.6
AI014087	ribosomal protein S26	829	187.3
AI014135	EST207690	334	79.6
AI070848	actin, beta	2279.3	316.4
AI071866	Rattus norvegicus Nclone10 mRNA	112.6	13.8
AI102044	Transcribed sequences	273.2	145
AI102505	cytochrome c oxidase, subunit VIIIa	1017.2	239.9
AI103074	ribosomal protein S12	1519.8	272.4
AI103236	ribosomal protein L35a	219.9	54.3
AI103396	EST212685	2594.2	860.4
AI103957	CD 81 antigen	596.5	132.4
AI104035	Transcribed sequence with strong similarity to protein ref:NP_079904.1 (M.musculus) F	380.5	174.2
AI104520	cytochrome c oxidase, subunit VIa, polypeptide 1	678	86.7
AI104544	ribosomal protein S17	302	42.2
AI105050	ATP synthase, H+ transporting, mitochondrial F1 complex, beta polypeptide	208.6	58.9
AI112173	ATPase Na+/K+ transporting beta 1 polypeptide	449.4	140.7
AI169104	Transcribed sequence with strong similarity to protein sp:P06765 (R.norvegicus) PLF4_	1048	174.6
AI169370	alpha-tubulin	1457.1	187.4
AI169802	ferritin, heavy polypeptide 1	2480.8	142.2
AI170268	Beta-2-microglobulin	760	106
AI170776	growth factor receptor bound protein 2	102.3	37
AI171085	Ribosomal protein L39	305.3	51.1
AI171355	EST217310	628	156.7
AI171844	ATP synthase, H+ transporting, mitochondrial F1 complex, epsilon subunit	210.4	38.1
AI172162	proteasome (prosome, macropain) subunit, beta type 4	483	112.9
AI172411	glutathione peroxidase 3	194.3	34
AI175486	ribosomal protein S7	149.1	31.8
AI176546	heat shock protein 86	277.8	99.1
AI176589	ribosomal protein L27	825.4	53.4
AI176589	ribosomal protein L27	372.8	52.4
AI176595	Cathepsin L	962.5	186.1
AI178207	ribosomal protein S21	451.8	25.1
AI178750	eukaryotic translation elongation factor 2	606.9	37.5
AI179012	actin, beta	9899	876
AI179150	EST222834	2946.6	465
AI179610	Heme oxygenase	669.1	65.4
AI180013	Fc receptor, IgG, alpha chain transporter	542.2	130.8
AI180424	Tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, zeta polyp	342.1	45.6
AI227887	cell division cycle 42 homolog (S. cerevisiae)	350.2	73.9
AI228674	Peptidylprolyl isomerase A (cyclophilin A)	868.6	193.2
AI229440	diaphorase 1	243.7	115.5
AI230406	EST227101	211.4	65.9
AI230748	tumor protein, translationally-controlled 1	2096.2	313.3
AI231292	cystatin C	694.3	243
AI231292	cystatin C	352.8	60.8
AI231807	ferritin light chain 1	2736	266

AI231807	ferritin light chain 1	977.8	217.8
AI232783	glutamine synthetase 1	299	94.6
AI233362	Lysosomal associated membrane protein 1 (120 kDa)	799.1	107.7
AI233749	ribosomal protein L30	263.2	100.4
AI234604	heat shock protein 8	884.4	103.2
AI235364	ribosomal protein S15a	290.1	45.2
AI235585	cathepsin D	1163.5	157.3
AI236795	heat shock protein 90 beta	356.1	44.8
AI237731	Lipoprotein lipase	394.5	160.1
AI237836	GNAS complex locus	1074.5	159.4
AI237836	GNAS complex locus	351.6	27.8
AI639012	Transcribed sequence with weak similarity to protein ref:NP_076947.1 (H.sapiens) hyp	111.1	20.4
AI639076	cDNA clone rx04025 3'	1034	313.9
AI639107	cDNA clone rx04084 3	333.9	84.8
AI639255	Transcribed sequences	212.7	35.5
AJ004912	integral membrane protein Tmp21-I (p23)	668.2	111.9
AJ009698	embigin	266.8	39.9
AJ009698	embigin	256.7	50.8
D10706	ornithine decarboxylase antizyme 1	451.8	42.2
D10706	ornithine decarboxylase antizyme 1	359.5	53.8
D10706	ornithine decarboxylase antizyme 1	241.4	60.8
D10755	proteasome (prosome, macropain) subunit, alpha type 6	105.4	34.2
D10854	aldo-keto reductase family 1, member A1	393.1	109.7
D10874	ATPase, H+ transporting, lysosomal (vacuolar proton pump) 16 kDa	1923.7	239.3
D10874	ATPase, H+ transporting, lysosomal (vacuolar proton pump) 16 kDa	764	100.3
D12771	solute carrier family 25 (mitochondrial carrier; adenine nucleotide translocator), member	410	138.4
D13127	ATP synthase, H+ transporting, mitochondrial F1 complex, O subunit (oligomycin sensit	143.1	60.7
D13966	insulin receptor-related receptor	217.2	107.7
D14688	myosin regulatory light chain	280.3	36.8
D16102	glycerol kinase	68.4	51.4
D16309	Cyclin D3	70.6	17.7
D16554	polyubiquitin	1234.1	246
D17445	Tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, eta polype	607.5	61.7
D25224	laminin receptor 1 (67kD, ribosomal protein SA)	606.8	75.1
D38380	Transferrin	147.6	44
D42116	complement receptor related protein	237.2	60.1
D45247	proteasome subunit RCX	131	36.7
D45249	protease (prosome, macropain) 28 subunit, alpha	220.6	28.3
D78308	calreticulin	596.6	140.9
D78613	Protein tyrosine phosphatase, receptor type, epsilon polypeptide	76.4	23.6
D84450	ATPase, Na+/K+ transporting, beta 3 polypeptide	202.3	78.1
D87840	Mucosal vascular addressin cell adhesion molecule 1	71	10.7
D90404	cathepsin C	196.6	39
D90404	cathepsin C	126.1	29.9
E01524	P450 (cytochrome) oxidoreductase	54.4	17.2
E01534	DNA sequence expressed especially in rat insulinoma	356.1	81.3
E02315	calmodulin	242.6	55.4
J00797	alpha-tubulin	155.1	46.6
J01435	mitochondrial cytochrome oxidase subunits I,II, III genes, ATPase subunit 6 gene	7775.6	1714.9
J01435	mitochondrial cytochrome oxidase subunits I,II, III genes, ATPase subunit 6 gene	4243	428.5
J01435	mitochondrial cytochrome oxidase subunits I,II, III genes, ATPase subunit 6 gene,	265.7	213.4
J01436	mitochondrial cytochrome B	1125.2	458
J02780	tropomyosin 4	283.3	43
J02962	lectin, galactose binding, soluble 3	730.9	129.8
J03969	nucleophosmin 1	122	21.9
K02816	pR-ET2 encoded oncodevelopmental protein	166.2	44.2
K02816	pR-ET2 encoded oncodevelopmental protein	83.7	29.6
K03250	ribosomal protein S11	489.1	29.2
L01624	serum/glucocorticoid regulated kinase	102	27.3
L03201	cathepsin S	1767.1	160.8
L12383	ADP-ribosylation factor 4	102.8	16.7
L13039	annexin II	278.4	44.8
L19927	ATP synthase, H+ transporting, mitochondrial F1 complex, gamma polypeptide 1	107.4	38.9
L19998	sulfotransferase family 1A, phenol-preferring, member 1	55.4	35.6
L28801	general transcription factor III C 1	79.7	34.9
L33869	ceruloplasmin	100.8	61.2
L38437	NADH dehydrogenase (ubiquinone) Fe-S protein 6	92.5	31.4
M10068	P450 (cytochrome) oxidoreductase	120.1	24.1
M11071	MHC RT1 class Ib gene	615.2	141.5
M11942	heat shock protein 8	397.4	58.3
M12672	GTP-binding protein (G-alpha-i2)	170.5	28.7
M12919	aldolase A	998.3	111.6
M12919	aldolase A	607.1	131.7
M13100	long interspersed repetitive DNA sequence LINE3	2109	793
M13100	long interspersed repetitive DNA sequence LINE3 (L1Rn)	107.2	14.3
M13100	long interspersed repetitive DNA sequence LINE3 (L1Rn)	75.5	20.8
M14050	heat shock 70kD protein 5	125	22.3
M14656	osteopontin	337.4	138.9
M15481	insulin-like growth factor 1	183.2	51.3

M15768	CD4 antigen	195.5	18.7
M15882	clathrin, light polypeptide (Lca)	583.3	117.9
M17419	ribosomal protein L5	597.5	61
M17701	Glyceraldehyde-3-phosphate dehydrogenase	564.1	131.3
M19936	prosaposin	782.6	79.4
M20156	ribosomal protein L18	441.2	77.9
M23591	Protein phosphatase 2 (formerly 2A), catalytic subunit, beta isoform	101.2	28.3
M24026	RT1 class Ib gene	425.1	52.4
M25758	phosphatidylinositol transfer protein	141.6	12.4
M27905	ribosomal protein L21	491.7	67.3
M28255	cytochrome c oxidase, subunit VIIIa	697.6	64.9
M29358	ribosomal protein S6	531.6	87
M29358	ribosomal protein S6	337	94.1
M29866	Complement component 3	275.4	34.8
M31322	sperm membrane protein (YWK-II)	132.4	52.4
M31788	phosphoglycerate kinase 1	217.5	40.5
M32016	Lysosomal-associated membrane protein 2	64.7	18.9
M32062	Fc receptor, IgG, low affinity III	479.6	126
M32062	Fc receptor, IgG, low affinity III	295.3	49.4
M34043	thymosin beta-4	13017.2	0
M34331	ribosomal protein L35	334.3	34.1
M34331	ribosomal protein L35	330	29.1
M38135	Cathepsin H	154.6	31.8
M54926	lactate dehydrogenase A	424.1	40.7
M58404	thymosin, beta 10	240.1	62.6
M81088	eukaryotic translation elongation factor 1 alpha 1	518.7	81.6
M83143	Sialyltransferase 1 (beta-galactoside alpha-2,6-sialytransferase)	119.6	53.8
M84716	ribosomal protein S3a	518.3	101.9
M86564	prothymosin alpha	196.5	10.8
M89646	ribosomal protein S24	530.5	83.1
M91597	nucleoside diphosphate kinase	261	116.5
M91652	glutamine synthetase 1	638.1	150.6
M91652	glutamine synthetase 1	207.7	67.2
M93257	catechol-O-methyltransferase (COMT)	296.6	98
M98820	Interleukin 1 beta	746.8	131.3
M98820	Interleukin 1 beta	163.6	71.6
S45392	heat shock protein 90	190.3	63.5
S45663	synaptic glycoprotein SC2	99.1	20.2
S46798	ND5, ND6 [rats, adult and senescent cerebral hemispheres, Mitochondrial, 954 nt]	368.4	124.3
S55223	tyrosine 3-monooxgenase/tryptophan 5 monooxgenase activation protein, beta polypep	296.8	68.4
S59893	La=autoantigen SS-B/La	93.3	44.1
S61973	NMDA receptor glutamate-binding chain	251	81.2
S61973	NMDA receptor glutamate-binding chain	90.7	36.9
S63233	phosphoglycerate mutase type B subunit	289.5	131.1
S71021	ribosomal protein L6	334.9	53.9
S72594	tissue inhibitor of metalloproteinase 2	450.6	66.1
S76779	apolipoprotein E	4595.2	336
S77858	non-muscle myosin alkali light chain	798.5	95.8
S79304	cytochrome oxidase subunit I,	3738.1	246.9
S81353	prosaposin	1751.9	115.6
S81497	lipase A, lysosomal acid	865.5	156.5
S82383	slow-twitch alpha TM/hTMnm homolog	718.4	101.2
S83025	TSH receptor suppressor element-binding protein-1=Y-box protein	403.7	117.4
S87522	leukotriene A4 hydrolase	225.4	47.6
U03390	guanine nucleotide binding protein, beta polypeptide 2-like 1	285.1	54.6
U10894	allograft inflammatory factor 1	254.9	99.3
U11071	polyadenylate-binding protein-related protein	811.7	237.3
U15138	LIC-2 dynein light intermediate chain 53/55	117.6	12.5
U17919	allograft inflammatory factor 1	251.9	75.2
U18729	cytochrome b558 alpha-subunit	781.3	232.2
U20643	aldolase A	503.9	65.4
U31598	MHC class II-like alpha chain RT1.Ma	220.6	53.1
U47315	brain protein 44-like	142	42.5
U49099	Transcribed sequence with strong similarity to protein sp:Q62931 (R.norvegicus) GS28_	136.7	39.9
U53184	LPS-induced TNF-alpha factor	232.8	32.2
U59184	bcl2-associated X protein	497.5	56.1
U77829	growth arrest specific 5	158.2	31.7
U77931	unknown mRNA	580.3	107.8
U95052	eukaryotic translation initiation factor 4 gamma, 2	218.1	65.3
U95178	disabled homolog 2, mitogen-responsive phosphoprotein (Drosophila)	355.4	35
V01217	actin, beta	5960.6	1537.5
X02610	enolase 1, alpha	435.6	68.2
X02918	Protein disulfide isomerase (Prolyl 4-hydroxylase, beta polypeptide)	273.3	57
X04979	apolipoprotein E	10319.7	557.4
X05472	RNREP24R Rat 2.4 kb repeat DNA right terminal region	359.1	152.2
X06423	ribosomal protein S8	583.5	76.9
X06423	ribosomal protein S8	425.7	40.1
X06483	ribosomal protein L32	1040.4	229
X14181	ribosomal protein L18a	1349.3	110.4

X14181	ribosomal protein L18a	698.2	68.1
X14210	ribosomal protein S4	1713.7	304.5
X14210	ribosomal protein S4	330.6	168
X14671	ribosomal protein L26	431.6	99.1
X14848	MIRNXX Rattus norvegicus mitochondrial genome	1297.7	268.5
X15013	ribosomal protein L7a	417.3	58.3
X15030	cytochrome c oxidase, subunit Va	432.8	94
X15096	acidic ribosomal protein P0	1116.3	141
X15216	ribosomal protein L21	83.9	18.5
X16145	Fucosidase, alpha-L-1, tissue	187.8	24.5
X17053	small inducible cytokine A2	256.8	98.3
X17053	small inducible cytokine A2	120.2	36
X17665	ribosomal protein S16	1279.9	225.9
X51536	ribosomal protein S3	873.4	79.7
X51536	ribosomal protein S3	243.1	44.7
X51706	ribosomal protein L9	540.7	109.6
X51706	ribosomal protein L9	342.7	70.2
X51707	ribosomal protein S19	648.3	192.8
X52619	ribosomal protein L28	681.8	30.5
X52733	ribosomal protein L27a	1529.5	188.9
X52815	cytoplasmic-gamma isoform of actin	2436.8	470.3
X53377	ribosomal protein S7	310.9	64.3
X53378	ribosomal protein S13	306.9	97.6
X53504	ribosomal protein L12	297.2	33.6
X54081	cytochrome c oxidase subunit IV	338.5	103.8
X54467	cathepsin D	2184.5	352.2
X54617	myosin regulatory light chain	197.3	43.9
X55153	ribosomal protein P2	1240.8	180.7
X57432	ribosomal protein S2	1777.9	167.5
X57529	ribosomal protein S18	1433.5	267.3
X58200	ribosomal protein L23	1653.8	478.2
X58200	ribosomal protein L23	662.4	63.7
X58389	Rat amino acid starvation-induced protein mRNA, 3' end	318.7	72.2
X58465	ribosomal protein S5	477.1	109.6
X58465	ribosomal protein S5	302.9	73.9
X59051	Ribosomal protein S29	2830.5	423.9
X59375	ribosomal protein S27	295.8	63.3
X60212	Rat amino acid starvation-induced protein mRNA, 3' end	1505.3	212.7
X61295	L1 retroposon, ORF2	74	17.5
X61654	CD63 antigen	1878.9	134.1
X62145	ribosomal protein L8	459.2	144.2
X62146	ribosomal protein L11	493.1	42.5
X62146	ribosomal protein L11	463.3	43.1
X62166	ribosomal protein L3	363.6	84.7
X62322	granulin	2421	421.4
X62322	granulin	771	96
X62482	ribosomal protein S25	177.1	51.3
X62671	hybrid protein (ubiquitin-like protein/rps30)	607.8	118.7
X62952	vimentin	241.4	27
X63434	Urinary plasminogen activator, urokinase	165	37
X66369	ribosomal protein L37	378.5	123.1
X66370	ribosomal protein S9	549.3	94.9
X70871	Cyclin G1	436.3	154.6
X71127	complement component 1, q subcomponent, beta polypeptide	507.6	83.2
X78167	ribosomal protein L15	280.1	85.9
X82396	cathepsin B	1595	132.7
X84047	XLas protein	205	28
X93352	ribosomal protein L10a	219.2	63.1
X94242	ribosomal protein L14	94.5	28.2
X95850	R.norvegicus mRNA for novel gene expressed in circadian manner, clone SCN8	174.1	59.4
X96437	PRG1 gene	71.7	19.9
Y00404	superoxide dismutase 1	734	270.4
Y00497	superoxide dismutase 2	202.1	271.5
Y12635	ATPase, H+ transporting, lysosomal (vacuolar proton pump), beta 56/58 kDa, isoform 2	160.5	53.2