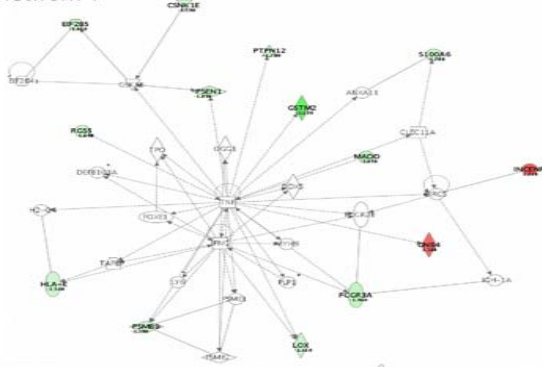
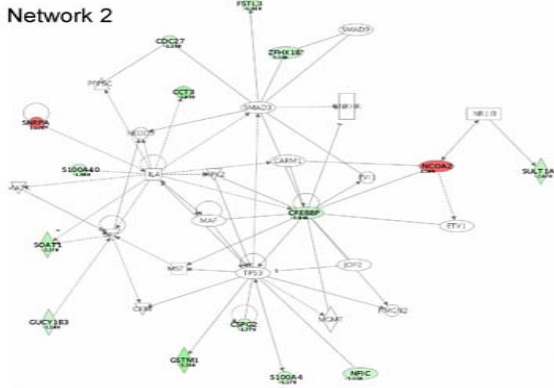


Supplemental Figure 1

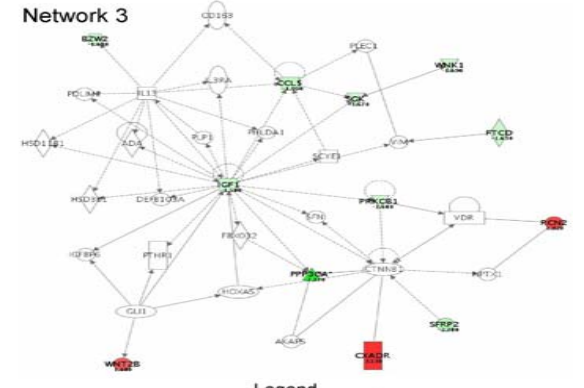
Week 4-Network 1



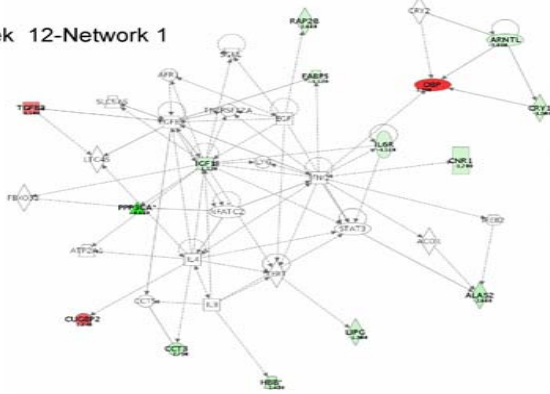
Network 2



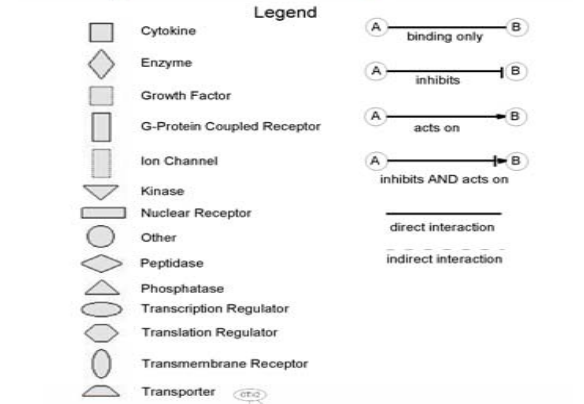
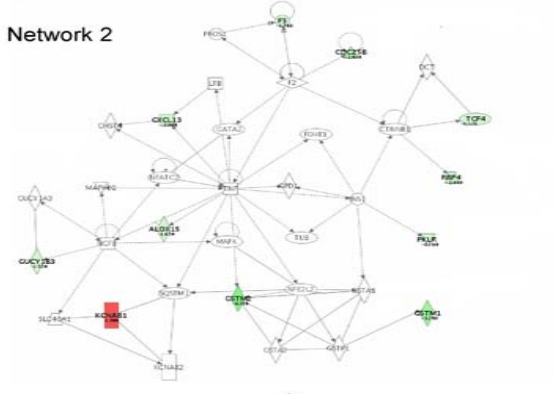
Network 3



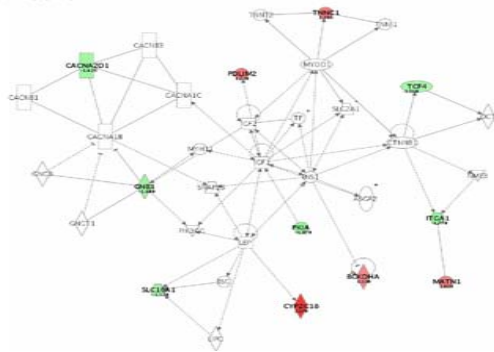
Week 12-Network 1



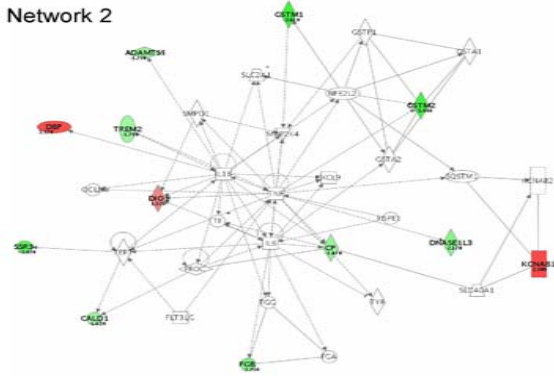
Network 2



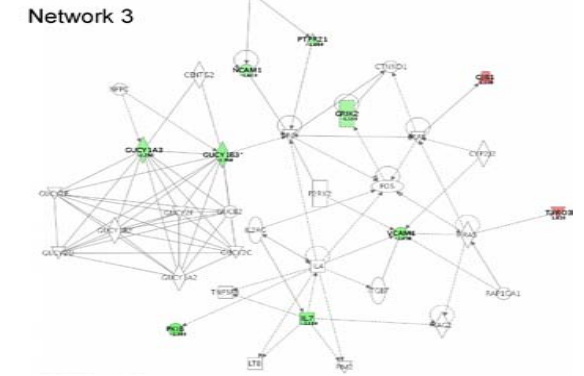
Week 20-Network 1



Network 2

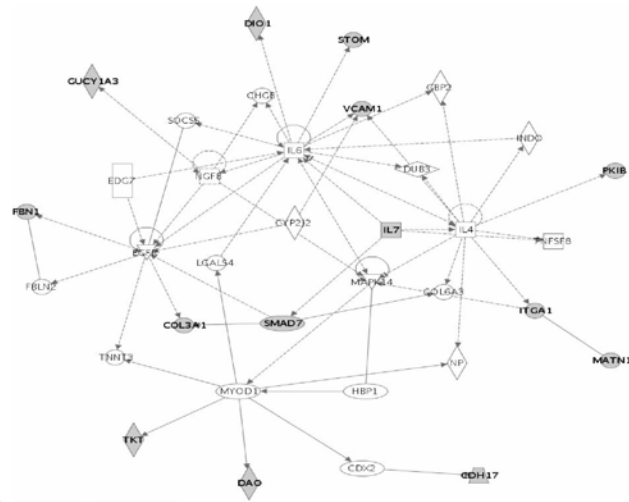


Network 3



Supplemental Figure 2

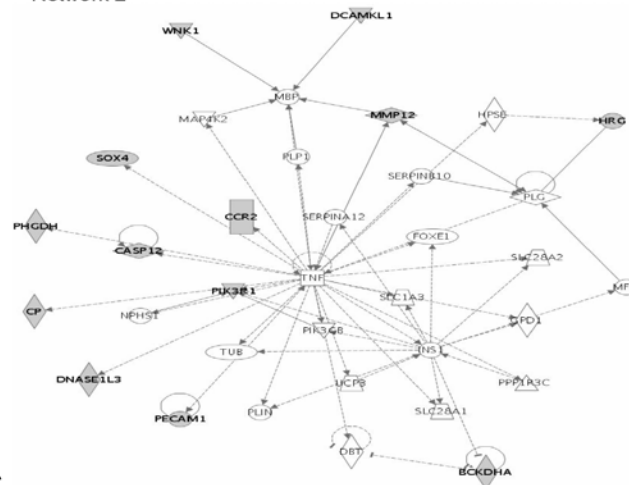
A Network 1



Top Function (All Networks)	Significance
Cellular Assembly and Organization	7.15E-5 - 4.86E-2
Cellular Movement	1.47E-4 - 4.86E-2
Immune and Lymphatic Development and Function	1.48E-4 - 4.86E-2
Immune Response	1.48E-4 - 4.86E-2

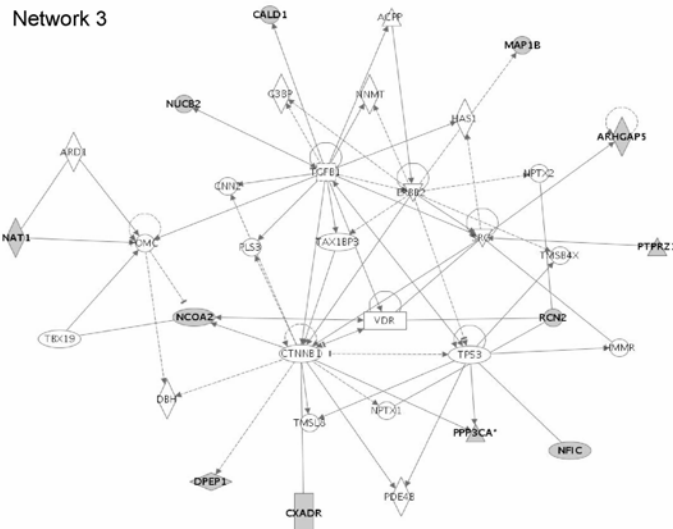
High Level Functions	Significance
Cellular Function and Maintenance	1.76E-9 - 6.33E-4
Cancer	7.32E-9 - 3.76E-3
Inflammatory Disease	2.85E-7 - 2.36E-3
Cell Morphology	1.80E-6 - 2.36E-3

B Network 2



High Level Functions	Significance
Molecular Transport	2.07E-7 - 3.84E-3
Carbohydrate Metabolism	2.42E-7 - 3.84E-3
Small Molecular Biochemistry	2.42E-7 - 3.84E-3
Cell Death	1.80E-6 - 3.84E-3

C Network 3



High Level Functions	Significance
Cellular Movement	2.07E-7 - 3.36E-3
Cellular Growth and Proliferation	2.42E-7 - 3.76E-3
Reproductive System Disease	2.42E-7 - 3.84E-3
Cancer	2.42E-7 - 3.84E-3

Supplemental Figure 1. Functional network analysis of genes differentially expressed between S and S.SHR(2) congenic at week 4, 12 , and 20. Nodes represent genes, with their shape representing the functional class of the gene product (see legend). The lines connecting the nodes indicate the biological relationship between each node (see legend). Nodes are color coded based on degree and direction of fold change [red, S.SHR(2) is overexpressed compared to S; and green, S.SHR(2) is underexpressed compared to S]. Note: each network is numbered (e.g. 1, 2...) based on the number of focus genes it contains. So, network 1 at week 4 and network 1 at week 12 need not be related.

Supplemental Figure 2. Functional network analysis of genes differentially expressed between S and S.SHR(2) and over time (strain X time). Nodes represent genes, with their shape representing the functional class of the gene product (see figure legend , supplemental Fig. 1). The lines connecting the nodes indicate the biological relationship between each node. Shaded nodes are genes found to be differentially expressed between strains and over time. High level functions for each network and significance level are shown. Top functions in the entire data set are also shown.

Type of file: table

Label: Tables S1 - S4

Filename: Copy of SupplementalTables1-4.xls

Table 1

Chromosome	Gene Identifier	Gene ID	Other ID	Position (Mb)	Direction	Week4		Week12		Week20		Gene Description		
						Ratio	p-value	Ratio	p-value	Ratio	p-value			
1	BP399429	-	1389026	149425862	Up	1.6	0.0157	0.1555				Rattus norvegicus similar to RIKEN cDNA 4633417L02 (LOC293117), mRNA		
1	BP399429	-	1389252	212326940	Down	1.7	0.0427	0.2103				Rattus norvegicus similar to inner centromere protein-8 (LOC293733), mRNA		
1	BI289559	-	1393376	173556971	Up	1.5	0.1441	0.1468				Rattus norvegicus similar to SOXB6 (LOC293165), mRNA		
1	AI230728	Snrpa	1388436	82295660	Up	1.5	0.1193	0.1677				Rattus norvegicus similar to U1 small nuclear ribonucleoprotein A (U1 snRNP A protein) (LOC292729), mRNA		
1	BP399445	-	1373465	207398410	Down	2.6	0.0021	0.0024				Rattus norvegicus similar to nuclear cap-binding protein 3, subunit 10 (ncaps 10) (trans. 15011702a)		
1	AF304763	Sufl1a1	1370019	185829205	Down	2.1	0.0298	0.1894				sulfotransferase family 1A, phenol-polyferin, member 1		
1	AI408286	-	1378193	213828662	Down	1.8	0.0481	0.2103				Rattus norvegicus similar to RIKEN cDNA AA430103C15 (LOC293744), mRNA		
1	NM_019232	Sgk	1367802	23401262	Down	1.7	0.1181	0.1640				serine/threonine-specific protein kinase		
1	AI145746	-	1392249	20355444	Down	1.6	0.0007	0.0593				Rattus norvegicus transcribed sequences		
1	AA964142	-	1398888	167532994	Down	2.1	0.0011	0.0973			1.7	0.0044	0.0372	Rattus norvegicus similar to RIKEN cDNA 2610209A2 (LOC368121), mRNA
1	YX4443	Prxcb3	1370585	151118102	Down	1.7	0.0027	0.0711			2.4	0.0039	0.0217	protein kinase C, beta 1
1	AA858962	Rbp4	1371762	242443798	Down	2.4	0.0039	0.0217					retinol binding protein 4	
1	NM_033234	Hbb	1367553	161590568	Down	2.4	0.0165	0.0461					hemoglobin beta chain complex	
1	BI287200	Hbb	1371245	161518261	Down	2.4	0.0074	0.0292					Rat hemoglobin beta-chain mRNA	
1	U13263	Fabp5	1370281	218829748	Down	1.6	0.0059	0.0241					fatty acid binding protein 5, epidermal	
1	AI230048	Dbp	1387874	96173573	Up	2.3	0.0252	0.0574			2.3	0.0007	0.0216	D site albumin promoter binding protein
1	AB012660	Amitl	1370510	114132014	Down	1.6	0.0221	0.0538			3.0	0.0294	0.0639	aryl hydrocarbon receptor nuclear translocator-like
1	AI189562	RSK-4	1371050	94412893	Down	1.6	0.0221	0.0538			3.0	0.0294	0.0639	kinase
1	NM_018184	Cyp2c2	1387328	243281320	Down	2.6	0.0034	0.0327			2.8	0.0051	0.0392	Cytochrome P450, subfamily IIC (mephentoin 4-hydroxylase)
1	BM385272	-	1393139	76971904	Up	2.0	0.0051	0.0392			1.8	0.0117	0.0323	Rattus norvegicus similar to Adiponectin O2 (LOC292697), mRNA
1	BI981129	-	1373369	197054477	Up	2.0	0.0145	0.0554					Cytochrome P450 monooxygenase CYP11B	
1	BM390571	-	1378248	96197898	Up	1.9	0.0410	0.0888					Rattus norvegicus similar to cytosolic sulfotransferase (LOC292915), mRNA	
1	AW522786	-	1378867	140157974	Up	1.9	0.0062	0.0477					Rattus norvegicus transcribed sequences	
1	BP286052	-	1393474	96197898	Up	1.7	0.0122	0.0526					Rattus norvegicus similar to cytosolic sulfotransferase (LOC292915), mRNA	
1	AI060133	-	1382476	207556536	Up	1.6	0.0242	0.0700			2.3	0.0047	0.0190	similarity to protein pdb 1L6G (E. coli) B Chain B, Lactose Operon Repressor Bound To 21-Base Pair Symmetric Operator Dna, Alpha Carbons Only
1	NM_033781	Cyp21	1390596	233410939	Up	1.6	0.0308	0.0780					Rattus norvegicus similar to Melanoma antigen recognized by T-cells 1, (MAST1-1) (Melan-A protein) (Antigen L839-AA) (Antigen SC29-AA), mRNA	
1	NM_143436	Cyp21	1390596	82229569	Up	1.6	0.0308	0.0780					Cytochrome P450 monooxygenase CYP21	
1	BF561222	-	1382424	87626241	Up	1.6	0.0002	0.0173					Rattus norvegicus transcribed sequence with moderate similarity to protein pI60307 (E. coli) 850307 beta-galactosidase, alpha peptide - Escherichia coli	
1	NM_033711	Arf2	1368166	208909458	Up	1.6	0.0283	0.0742					AHP-ribosylase-like 2	
1	AA054379	Arp1	1390017	85390213	Down	1.6	0.0458	0.0937					Rattus norvegicus transcribed sequence with strong similarity to protein ref NP_005157 (H.sapiens) amyloid beta (A4) precursor-like protein 1 Homo sapiens1	
1	BE098025	-	1374924	22263990	Up	1.5	0.0093	0.0493					Rattus norvegicus transcribed sequences	
1	BE112882	-	1383008	201997982	Up	1.5	0.0284	0.0723					Rattus norvegicus similar to TOLLUP protein (LOC361677), mRNA	
1	AI287507	Bochkha	1317981	60837907	Up	1.5	0.0098	0.0448					branched chain keto acid decarboxylase subunit E1, alpha polypeptide	
1	BI283664	-	1377209	131370265	Up	1.5	0.0029	0.0327					Rattus norvegicus transcribed sequences	
1	AI072405	Aap11	1384877	164973796	Up	1.5	0.0100	0.0493					scaevola 11	
1	BI099174	-	1382118	245725403	Down	1.5	0.0051	0.0327					Rattus norvegicus similar to RIKEN cDNA 4303521E07 (LOC309486), mRNA	
1	AI008432	-	1392843	31940826	Down	2.2	0.0174	0.0610					Rattus norvegicus transcribed sequences	
1	BM389922	-	1380768	14816855	Down	2.1	0.0082	0.0477					Rattus norvegicus transcribed sequences	
1	BM382448	-	1383688	168011643	Down	1.9	0.0119	0.0522					Rattus norvegicus transcribed sequences	
1	BF397243	-	1395047	241156989	Down	1.9	0.0007	0.0216					Rattus norvegicus similar to tankyrase, TRF1-interacting ankyrin-related ADP-ribose polymerase 2, tankyrase 2 (LOC309512), mRNA	
1	AW920054	Ctsc	1397808	144840775	Down	1.9	0.0415	0.0888					Rattus norvegicus transcribed sequences	
1	BE091460	-	1380760	463994401	Down	1.8	0.0068	0.0448					Rattus norvegicus transcribed sequences	
1	AI114002	-	1374775	195310301	Down	1.8	0.0134	0.0546					Rattus norvegicus similar to pC-F7 (LOC291234), mRNA	
1	AI511069	-	1378391	109033934	Down	1.7	0.0070	0.0448					Rattus norvegicus transcribed sequences	
1	AI170304	Mbeat1	1389006	215387912	Down	1.7	0.0069	0.0493					macrophage-expressed gene 1	
1	AA909491	-	1385351	46385119	Down	1.7	0.0431	0.0933					Rattus norvegicus transcribed sequences	
1	BI289045	-	1381557	219904685	Down	1.6	0.0379	0.0866					Rattus norvegicus transcribed sequences	
1	AI008409	-	1389986	208427238	Down	1.6	0.0267	0.0712					Rattus norvegicus transcribed sequence with strong similarity to protein spP00722 (E. coli) B GAL_ECOLI Beta-galactosidase (Lactase)	
1	BM389792	-	1397463	245429376	Down	1.6	0.0048	0.0162					Rattus norvegicus similar to pC-F7 (LOC291234), mRNA	
1	AI171480	-	1390909	162170846	Down	1.6	0.0007	0.0126					similarity to protein pdb 1J25 (E. coli) B Chain B, E. Coli (LacZ) Beta-Galactosidase In Complex With D-Galactopyranosyl-1-On	
1	BE115281	Qki	1372542	45026384	Down	1.5	0.0326	0.0788					Rattus norvegicus transcribed sequences	
1	AI245172	Qki	1387659	224517843	Down	1.5	0.0418	0.0888					guanine deaminase	
1	BG670294	-	1390107	147022234	Down	1.5	0.0095	0.0493					similarity to protein pdb 1L6G (E. coli) B Chain B, Lactose Operon Repressor Bound To 21-Base Pair Symmetric Operator Dna, Alpha Carbons Only	
2	AI501417	-	1386831	140736218	Up	2.4	0.0312	0.1809					Rattus norvegicus transcribed sequences	
2	BE282916	-	1392728	100709895	Up	1.9	0.0092	0.0493					Rattus norvegicus transcribed sequences	
2	BE112158	-	1374640	189287027	Up	1.7	0.0426	0.2103					Rattus norvegicus transcribed sequences	
2	BF543355	Isgf10	1390901	148704637	Up	1.6	0.0070	0.0363					bone-specific C/EBP-beta	
2	AI573368	-	1396636	223039449	Up	1.6	0.0034	0.0264					Rattus norvegicus transcribed sequences	
2	BI284800	-	1371071	118791451	Up	1.6	0.0226	0.1748					Rattus norvegicus guanine nucleotide binding protein beta 4 subunit mRNA, partial cds	
2	BF406099	-	1384824	138616556	Up	1.5	0.0018	0.0604					Rattus norvegicus transcribed sequences	
2	BF398545	Sh2c2	1396614	175479529	Down	2.4	0.0047	0.0327					secreted frizzled-related protein 3	
2	AF140232	S100a6	1367661	182895080	Down	1.8	0.0481	0.2103					calcium binding protein A6 (calpalcin)	
2	NM_020538	Ariad	1369462	149348935	Down	1.7	0.0205	0.1716					invariant (amide) diacylglycerol esterase	
2	AA891529	-	1382954	192441767	Down	1.6	0.0021	0.0604					Rattus norvegicus transcribed sequence with weak similarity to protein sp Q92664 (H.sapiens) TF3A, HUMAN Transcription factor IIIA (Factor A) (TFIIIA)	
2	NM_012618	S100a4	1367846	182885070	Down	1.6	0.0276	0.1858					S100 calcium-binding protein A4	
2	AI407385	-	1371480	112227120	Down	1.6	0.0219	0.1738					Rattus norvegicus transcribed sequence with strong similarity to protein pI36670 (H.sapiens) A36670 cell division control protein Cks1 - human	
2	BF390321	-	1392282	191399944	Down	1.5	0.0110	0.1393					Rattus norvegicus similar to hypothetical protein MGC46719 (LOC295283), mRNA	
2	BF553729	Wnt2b	1397620	200219630	Up	1.7	0.0012	0.0593			2.2	0.0000	0.0109	S100 calcium-binding protein A4
2	AI554644	Wnt2b	1394901	200215823	Up	2.4	0.0055	0.0410					Rattus norvegicus transcribed sequences	
2	AI175140	-	1384307	161953228	Up	2.0	0.0040	0.0217					Rattus norvegicus similar to histone protein Hist23C1 (LOC310679), mRNA	
2	NM_033114	S100a10	1388890	188645674	Down	2.0	0.0470	0.2103					S-100 related protein, clone A2C	
2	AF084544	Capo2	1371232	18712829	Down	1.6	0.0424	0.2103					chondroitin sulfate proteoglycan 2 (versican)	
2	AI556018	-	1376996	231018070	Down	1.6	0.0140	0.1496			1.6	0.0196	0.0632	Rattus norvegicus similar to KIAA1246 protein (LOC310859), mRNA
2	NM_012769	Gucy1b3	1389097	173684385	Down	1.6	0.0151	0.1530			2.0	0.0026	0.0151	guanylate cyclase 1, soluble, beta 3
2	BF399387	Gucy1b3	1374389	173684385	Down	1.5	0.0070	0.0279			1.7	0.0028	0.0327	guanylate cyclase 1, soluble, beta 3
2	BI5374101	-	1381262	181637899	Up	3.6	0.0081	0.0977			2.7	0.0008	0.0099	Rattus norvegicus transcribed sequences
2	AA909892	-	1389811	152104473	Up	2.6	0.0238	0.1768			2.2	0.0003	0.0072	Rattus norvegicus transcribed sequences
2	AA900057	Snx2	1379804	189474923	Up	1.9	0.0062	0.0977			1.8	0.0015	0.0134	PDZ protein Mrt1
2	AI176320	-	1373416	113315728	Up	1.8	0.0012	0.0593			1.6	0.0048	0.0225	Rattus norvegicus transcribed sequences
2	BF522863	-	1386768	212890950	Down	2.1	0.0057	0.0977			2.1	0.0033	0.0214	similarity to protein pdb 4GTU (H.sapiens) G Chain G, Ligand-Free Homodimeric Human Glutathione S-Transferase M4-4 (E.C.2.5.1.18)
2	BI285782	-	1375909	203060404	Down	1.9	0.0020							

Table 1

Chromosome	Gene Identifier	Gene ID	Other ID	Position (Mb)	Direction	Ratio	p-value	adj-p-value	Ratio	p-value	adj-p-value	Direction	Ratio	p-value	adj-p-value	Gene Description
2	AA816268	-	1362274 at	157331763	Up		1.8	0.0008	0.0097	2.3	0.0031	0.0327				Retinoic acid receptor responder protein 1 (Tazartolene-induced gene 1 protein) (RAK-responsive protein TIG1) (LOC310486), mRNA
2	AI111940	-	13757594 at	160737614	Up		1.7	0.0265	0.0616	1.9	0.0203	0.0327				Rattus norvegicus similar to BCR4 aminesac-binding protein (LOC26213), mRNA
2	NM_021584	-	1387276 at	144686423	Down		2.0	0.0456	0.0840	2.8	0.0022	0.0327				actin and neurotransmitter-induced early gene protein 4 (ania-4)
2	BE102803	-	1383867 at	118197860	Down		1.9	0.0146	0.0432	1.9	0.0013	0.0292				Rattus norvegicus similar to BIF-5A2 protein (LOC310281), mRNA
2	BF490207	-	1393263 at	118199119	Down		1.7	0.0259	0.0241	0.303	0.0327	0.0327				Rattus norvegicus transcribed sequences
2	BE120508	-	1319781 at	118931483	Down		1.6	0.0209	0.0528	1.6	0.0443	0.0515				Rattus norvegicus similar to BAF-53a (LOC381925), mRNA
2	NM_017020	Irf7	1386987 at	182078051	Down		1.5	0.0104	0.0351	1.8	0.0138	0.0549				interleukin 6 receptor
2	BF411765	Celr2	1371018 at	203960524	Up		2.1	0.0002	0.0173		0.0002	0.0173				cadherin EGF-LAG seven-pass C-type receptor 2
2	BF549971	-	1377706 x at	913063	Up		1.9	0.0212	0.0847			0.0847				---
2	AI112346	-	1393952 at	89037914	Up		1.8	0.0030	0.0327			0.0327				Rattus norvegicus transcribed sequences
2	AI137898	-	1393330 at	84724179	Up		1.8	0.0153	0.0574			0.0574				Rattus norvegicus transcribed sequences
2	BI284261	-	1377869 at	140084501	Up		1.7	0.0336	0.0802			0.0802				Rattus norvegicus similar to carbon catabolite repression 4 protein homolog (LOC310395), mRNA
2	BQ371895	-	1383499 at	203756191	Up		1.7	0.0098	0.0438			0.0438				Rattus norvegicus transcribed sequences
2	AI137642	-	1393164 at	90341592	Up		1.7	0.0311	0.0783			0.0783				Rattus norvegicus similar to mitochondrial ribosomal protein S27 (LOC361883), mRNA
2	BI284747	-	1390973 at	189597223	Up		1.7	0.0490	0.0966			0.0966				Rattus norvegicus similar to tripartite motif-containing 45 (LOC295323), mRNA
2	BQ374446	-	1374760 at	80380796	Up		1.6	0.0009	0.0232			0.0232				Rattus norvegicus transcribed sequences
2	AB021517	-	1384653 at	218712747	Down		1.6	0.0293	0.0740			0.0740				Rattus norvegicus similar to ATP-binding cassette transporter (LOC310836), mRNA
2	BF284054	Mccc1	1395844 at	122437985	Up		1.6	0.0104	0.0493			0.0493				Rattus norvegicus similar to methylcrotonoyl-Coenzyme A carboxylase 1 (alpha) (LOC294972), mRNA
2	BE116137	-	1378929 at	180446723	Up		1.6	0.0493	0.0966			0.0966				Rattus norvegicus transcribed sequences
2	NM_130747	rACH	1394945 at	21984949	Up		1.6	0.0144	0.0554			0.0554				cytosolic acyl-CoA hydrolase
2	AI228548	S100a1	1388459 at	182784683	Up		1.6	0.0155	0.0575			0.0575				Rattus norvegicus similar to S-100 protein, alpha chain (LOC295214), mRNA
2	NM_024187	Clf	1386205 at	227308841	Down		4.0	0.0077	0.0463			0.0463				complement factor 1
2	NM_031120	Srs3	1360971 at	155149234	Down		2.9	0.0062	0.0428			0.0428				IRAP-complex gamma subunit
2	BE109107	-	1384969 at	243721308	Down		2.5	0.0468	0.0946			0.0946				Rattus norvegicus transcribed sequences
2	NM_012932	Cp	1368420 at	105086278	Down		2.3	0.0170	0.0608			0.0608				catenoplagmin
2	AF202115	Cp	1368418 at	105086278	Down		1.9	0.0529	0.0529			0.0529				catenoplagmin
2	AF202115	Cp	1368419 at	105086278	Down		1.9	0.0090	0.0493			0.0493				catenoplagmin
2	UP5675	Fab	1373511 at	174767192	Down		2.3	0.0106	0.0493			0.0493				fibronectin, beta polypeptide
2	BE290410	-	1388052 at	581005566	Down		2.2	0.0033	0.0327			0.0327				Rattus norvegicus transcribed sequences
2	AF367210	Irf7	1389208 at	96356083	Down		2.1	0.0041	0.0364			0.0364				interleukin 7
2	AI170450	-	1391587 at	77391890	Down		2.1	0.0218	0.0653			0.0653				Rattus norvegicus transcribed sequences
2	NM_012889	Vcam1	1388474 at	212227654	Down		2.1	0.0051	0.0364			0.0364				vascular cell adhesion molecule-1
2	BE120370	-	1384303 at	202489628	Down		2.1	0.0058	0.0417			0.0417				Rattus norvegicus transcribed sequences
2	NM_059772	Plk1	1389892 at	96824399	Down		2.0	0.0426	0.0893			0.0893				protein kinase inhibitor, alpha
2	AA803743	Plk1	1382176 at	96821745	Down		2.0	0.0024	0.0327			0.0327				protein kinase inhibitor, alpha
2	AW528452	-	1381688 at	236389754	Down		2.0	0.0275	0.0740			0.0740				Rattus norvegicus transcribed sequences
2	BF283984	-	1379444 at	171388285	Down		1.9	0.0111	0.0501			0.0501				Rattus norvegicus transcribed sequences
2	BE109744	-	1392852 at	180627289	Down		1.9	0.0028	0.0327			0.0327				Rattus norvegicus similar to transcription repressor p66 (LOC310814), mRNA
2	BM385061	-	1391106 at	56981705	Down		1.9	0.0124	0.0528			0.0528				Rattus norvegicus transcribed sequences
2	BF282631	C7	1383291 at	54144754	Down		1.9	0.0273	0.0736			0.0736				Rattus norvegicus similar to Complement component C7 precursor (LOC310126), mRNA
2	BI27436	-	1393169 at	219714168	Down		1.9	0.0292	0.0792			0.0792				similarity to protein ar-RGECDW (E. coli) RGECDW transcription activator of D-serine dehydratase - Escherichia coli
2	BI27472	-	1390404 at	213164026	Down		1.8	0.0426	0.0946			0.0946				Rattus norvegicus transcribed sequences
2	BI282029	-	1395388 at	88552849	Down		1.8	0.0336	0.0802			0.0802				Rattus norvegicus transcribed sequences
2	AA957364	Ibaa1	1372728 at	47172364	Down		1.8	0.0072	0.0448			0.0448				microRNA-1
2	NM_030964	Ibaa1	1387744 at	47107864	Down		1.8	0.0166	0.0596			0.0596				microRNA-1
2	M67405	Galcyl1a3	1387079 at	173750007	Down		1.8	0.0003	0.0173			0.0173				quadrifate cyclase 1, soluble, alpha 3
2	NM_019225	Slic1a3	1389895 at	98270236	Down		1.8	0.0286	0.0747			0.0747				solute carrier family 1, member 3
2	AW528269	-	1379691 at	383837	Down		1.8	0.0093	0.0327			0.0327				Rattus norvegicus retroviral-like C-Ha-ras proto-oncogene mRNA, partial sequence
2	AI237532	-	1385872 at	152125382	Down		1.8	0.0002	0.0173			0.0173				Rattus norvegicus transcribed sequences
2	UP7887	Ania4	1398985 at	144267748	Down		1.7	0.0080	0.0475			0.0475				actin and neurotransmitter-induced early gene protein 4 (ania-4)
2	BF284922	-	1393985 at	118026379	Down		1.7	0.0108	0.0493			0.0493				Rattus norvegicus similar to osteoblast specific factor 2 precursor (LOC381945), mRNA
2	BM389026	-	1373911 at	143531710	Down		1.7	0.0149	0.0565			0.0565				Rattus norvegicus similar to osteoblast specific factor 2 precursor (LOC381945), mRNA
2	AW52618	-	1378133 at	139240746	Down		1.7	0.0369	0.0845			0.0845				Rattus norvegicus transcribed sequences
2	AA858564	-	1379942 at	100989394	Down		1.7	0.0045	0.0376			0.0376				Rattus norvegicus transcribed sequences
2	BE108581	-	1382746 x at	224929644	Down		1.7	0.0211	0.0647			0.0647				Rattus norvegicus transcribed sequences
2	BI281702	Map1b	1373363 at	30415255	Down		1.7	0.0341	0.0803			0.0803				microtubule-associated protein 1b
2	AA851404	-	1391501 at	158802615	Down		1.7	0.0029	0.0327			0.0327				Rattus norvegicus similar to RIKEN cDNA 4921524P20 (LOC298106), mRNA
2	AI409635	-	1375029 at	179487312	Down		1.7	0.0172	0.0609			0.0609				similarity to protein dcD1 (E. coli) B Chain B. Lactose Operon Repressor Bound to 21-Base Pair Symmetric Operator Dna. Alpha Carbons Only
2	BI282167	Cdh6	1378480 at	62543102	Down		1.6	0.0497	0.0966			0.0966				Rattus norvegicus transcribed sequences
2	AA864378	Cdh6	1378614 at	62544084	Down		1.6	0.0154	0.0574			0.0574				Rattus norvegicus transcribed sequences
2	AI574991	-	1379356 at	204567499	Down		1.6	0.0019	0.0325			0.0325				Rattus norvegicus transcribed sequences
2	NM_019243	Platn	1387986 at	18601505	Down		1.6	0.0228	0.0687			0.0687				crocataldin F2 receptor, negative regulator
2	BM38077	-	1393146 at	144831076	Down		1.6	0.0075	0.0376			0.0376				Rattus norvegicus transcribed sequences
2	AA859029	-	1393751 at	93469787	Down		1.6	0.0531	0.127			0.127				similarity to protein gp-P0072 (E. coli) [GAL, ECOLI] Beta-galactosidase (Lactase)
2	AI178422	-	1381771 at	148600812	Down		1.6	0.0491	0.0966			0.0966				Rattus norvegicus transcribed sequences
2	AW191545	-	1392254 at	27599983	Down		1.6	0.0016	0.0323			0.0323				---
2	BM385126	-	1394545 at	4899	Down		1.6	0.0456	0.0912			0.0912				---
2	BM389035	-	1378453 at	47107864	Down		1.6	0.0477	0.0957			0.0957				Rattus norvegicus transcribed sequences
2	AA851345	-	1391979 at	56891395	Down		1.6	0.0494	0.0966			0.0966				Rattus norvegicus transcribed sequences
2	BI282008	-	1389003 at	2911103	Down		1.6	0.0005	0.0182			0.0182				Rattus norvegicus transcribed sequences
2	BE102821	-	1378153 at	140408472	Down		1.5	0.0197	0.0632			0.0632				Rattus norvegicus transcribed sequences
2	AW528269	-	1386100 at	11606188	Down		1.5	0.0198	0.0632			0.0632				similar to misshapen/NIK-related kinase isoform 2; GCK family kinase MINK; serine/threonine protein kinase (LOC294917), mRNA
2	NM_012716	Sic1b1	1386681 at	159860337	Down		1.5	0.0001	0.0011			0.0011				solute carrier family 16, member 1
2	AI102620	Map3k1	1375673 at	43062252	Down		1.5	0.0177	0.0614			0.0614				

Table 1

Chromosome	Gene Identifier	Gene ID	Other ID	Position (Mb)	Direction	Ratio	Week4		Week12		Week20		Gene Description		
							p-value	adj-p-value	Ratio	p-value	adj-p-value	Ratio		p-value	adj-p-value
3	BF415891	-	1362441 at	25564295	Down	1.5	0.0021	0.0804					similarity to protein sp.06015 (H. sapiens) SIF1_HUMAN Zinc finger homeobox protein 1b (Smad interacting protein 1) (SMAD1P1)		
3	BF54887	-	1365153 at	96911095	Down				1.7	0.0303	0.0635		Rattus norvegicus transcribed sequences		
3	NM_133572	Cdc25b	1370034 at	118893716	Down				3.0	0.0040	0.0217	1.5	0.0142	0.0551	cell division cycle 25B
3	AJ234249	-	1374630 at	36223883	Up				1.7	0.0375	0.0846				Rattus norvegicus similar to chloride intracellular channel 3 (LOC366566). mRNA
3	NM_017092	Twp3	1375793 at	110339871	Up				1.8	0.0343	0.0372				Tyrosin protein kinase kinase 3
3	AJ071620	-	1385580 at	142009065	Up				1.6	0.0476	0.0957				Rattus norvegicus transcribed sequences
3	BF75292	-	1377088 at	142260679	Up				1.6	0.0202	0.0327				Rattus norvegicus similar to RIKEN cDNA 2310046K01 (LOC311536). mRNA
3	NM_134403	Ahr2	138111 at	86853338	Down				1.6	0.0029	0.0327				Cytochrome P450
3	AF-067650	Sardh	1370573 at	6075853	Up				1.5	0.0211	0.0647				sarcosine dehydrogenase
3	BE105446	Fnbp1	1377342 at	10091465	Up				1.5	0.0275	0.0736				racalmin
3	BC06106	-	1383307 at	75309098	Down				1.5	0.0343	0.0654				Rattus norvegicus similar to nicotinic acetylcholine receptor-associated 46K protein - mouse (LOC362161). mRNA
3	NM_031055	Mmp9	1398275 at	155995473	Up				1.5	0.0208	0.0646				matrix metalloproteinase 9 (gelatinase B, 92-kDa type IV collagenase)
3	BI294934	-	1380612 at	169762453	Down				1.6	0.0174	0.0610				Rattus norvegicus transcribed sequences
3	BF543871	-	1384382 at	44547884	Down				1.6	0.0167	0.0598				Rattus norvegicus transcribed sequences
3	BF565756	-	1388695 at	42002053	Down				1.6	0.0201	0.0639				Rattus norvegicus transcribed sequences
3	BF542618	-	1394845 at	100002831	Down				1.5	0.0068	0.0448				Rattus norvegicus similar to mKIAA9560 protein (LOC366183). mRNA
3	BF552077	-	1372307 at	112303545	Down				1.5	0.0364	0.0837				Rattus norvegicus similar to Na-Ca exchanger 5 (LOC313387). mRNA
3	AJ012697	-	1377072 at	94611907	Up	1.6	0.0159	0.1557							similarity to protein ref.NP_002680.1 (H. sapiens) polymerase (DNA-directed), alpha (TK01) [Homo sapiens]
4	BF397628	-	1396422 at	121746928	Up	1.6	0.0228	0.1748							protein kinase, transmembrane type 12
4	NM_057115	Phn12	1369496 at	9617633	Down	1.6	0.0601	0.0372							protein kinase (phosphatase, non-receptor) type 12
4	AJ011345	Prknk1	1391140 at	156297841	Down	1.7	0.0372	0.2071							CASA transcription factor 1
4	BE116205	-	1384795 at	118125137	Down	1.7	0.0283	0.1877							Rattus norvegicus similar to nuclear protein, NP220 (LOC312491). mRNA
4	BE109884	-	1379651 at	133722289	Down	1.7	0.0306	0.1906							Rattus norvegicus similar to forkhead-related transcription factor 1A (LOC287480). mRNA
4	BC070960	MGC95152	1395919 at	83981797	Down	1.6	0.0443	0.2103							Rattus norvegicus similar to R32020.0L019K protein (LOC297100). mRNA
4	BE115521	-	1391170 at	148981770	Down	1.6	0.0472	0.2103							Rattus norvegicus similar to mKIAA1757 protein (LOC297514). mRNA
4	AJ714037	-	1384729 at	156334911	Down	1.6	0.0088	0.1249							Rattus norvegicus transcribed sequences
4	NM_017335	Sicr12	1397295 at	577151292	Down	1.5	0.0414	0.2103							CASA transcription factor 2
4	AJ529483	-	1378365 at	117432385	Down	1.9	0.0138	0.1498	1.6	0.0218	0.0538				Rattus norvegicus transcribed sequences
4	AJ385379	-	1398107 at	83462077	Down	1.9	0.0455	0.0840							Rattus norvegicus similar to RIKEN cDNA A030007L17; EST: AA673177 (LOC362370). mRNA
4	BI211290	-	1394963 at	43309017	Down				1.8	0.0124	0.0395				Rattus norvegicus transcribed sequences
4	AJ548449	-	1384851 at	94549854	Up							1.7	0.0131	0.0540	similarity to protein ref.NP_064544.1 (H. sapiens) hypothetical protein DKFZ6762K2015 (Homo sapiens)
4	BF400819	Gain11	1396602 at	5272176	Up				1.6	0.0069	0.0448				Rattus norvegicus transcribed sequences
4	AJ181344	Onb5	1370961 at	14995495	Up				1.6	0.0204	0.0648				camelinolase 3
4	AF023090	-	1372293 at	8964935	Up				1.6	0.0178	0.0614				Rattus norvegicus transcribed sequences
4	AJ235658	-	1383023 at	57124082	Up				1.6	0.0380	0.0846				similar to Ubiquitin-conjugating enzyme E2 H (Ubiquitin-protein ligase H) (Ubiquitin carrier protein H) (UBCH2) (E2-20K) (LOC296956). mRNA
4	AJ232716	-	1373913 at	83934016	Down				1.6	0.0270	0.0846				Rattus norvegicus transcribed sequences
4	AJ230054	-	1373902 at	158801835	Up				1.5	0.0288	0.0749				Rattus norvegicus similar to hypothetical protein MGC47816 (LOC302038). mRNA
4	AA848944	-	1379584 at	168128286	Down	2.1	0.0267	0.0712							Rattus norvegicus transcribed sequences
4	AA690113	-	1382083 at	80865142	Down	2.1	0.0147	0.0512							similar to protein ref.AJ23284 (H. sapiens) A57384 multimerin, endothelial cell, precursor - human
4	AJ915454	-	1376394 at	186697703	Down	1.9	0.0188	0.0629							Rattus norvegicus transcribed sequences
4	UJ68883	Kfra5	1370740 at	168840017	Down				1.6	0.0110	0.0500				Ly-49.9 antigen
4	AJ716184	Fgfr2	1372884 at	91783133	Down	1.7	0.0030	0.0330							fibroblast growth factor receptor 2
4	NM_013080	Hhr23	1368350 at	43534202	Down	1.7	0.0003	0.0173							Rattus norvegicus similar to theother 5-methyltransferase (LOC297114). mRNA
4	BM392106	Cald1	1368824 at	62062872	Down	1.6	0.0069	0.0448							Rattus norvegicus similar to talin 2 (LOC315776). mRNA
4	BI291848	Gnid1	1371969 at	62062872	Down	1.6	0.0143	0.0564							calcaldinin 1
4	AJ140952	Cfrcan2d1	1396849 at	153398718	Down	1.6	0.0108	0.0467							calcium channel, voltage-dependent, alpha2delta subunit 1
4	BF281550	-	1390810 at	25065233	Down				1.6	0.0197	0.0632				Rattus norvegicus transcribed sequences
4	AA943147	-	1382802 at	87279682	Down				1.6	0.0043	0.0372				Rattus norvegicus similar to hypothetical protein FLJ20637 (LOC362376). mRNA
4	AJ694045	-	1398933 at	171879448	Up	1.7	0.0028	0.0711							Rattus norvegicus similar to R6511 homolog (LOC298878). mRNA
4	BF401989	Neoa2	1396254 at	5248682	Up	1.6	0.0442	0.2103							Rattus norvegicus transcribed sequences
4	AA909447	-	1382715 at	96407867	Up	1.6	0.0118	0.1401							Rattus norvegicus similar to cysteine-rich protein NF-X-1 (LOC313166). mRNA
4	AJ111618	MGC105681	1373063 at	56658101	Down	2.0	0.0062	0.2103							Rattus norvegicus similar to C10 G chain (LOC362854). mRNA
4	BM386775	-	1379228 at	152821172	Down	1.6	0.0368	0.2071							Rattus norvegicus similar to absent in melanoma 1 (LOC298543). mRNA
4	NM_053718	Mll3	1388279 at	107070479	Down	1.5	0.0431	0.2103							(myeloid/lymphoid or mixed-lineage leukemia (fms/retroviral) (Drosophila) homolog; translocated to 3
4	BI211980	-	1371291 at	168128286	Down				1.5	0.0294	0.0629				Rattus norvegicus transcribed sequences
4	XJ5812	Cnr1	1369677 at	50427173	Down				1.6	0.0151	0.0433				canabinoid receptor 1
4	BI278779	Slc30a2	1378896 at	163087073	Up	2.0	0.0032	0.0327							solute carrier family 30, member 2
4	NM_012980	Slc30a2	1380264 at	163087073	Up	1.9	0.0192	0.0630							solute carrier family 30, member 2
4	BM384214	Maih1	1380579 at	148905644	Up	1.6	0.0005	0.0182							Rattus norvegicus similar to Matrilin 1, cartilage matrix protein 1 (LOC297894). mRNA
4	BF284337	-	1373963 at	79494958	Up	1.6	0.0099	0.0493							Rattus norvegicus similar to RIKEN cDNA 281043D12 (LOC298097). mRNA
4	NM_021883	Ox1	1390259 at	129385247	Up	1.6	0.0025	0.0327							deoxianilic acid oxidase, type 1
4	AJ145487	-	1393945 at	40708424	Down	2.2	0.0072	0.0448							Rattus norvegicus transcribed sequences
4	NM_053977	Cohn17	1389224 at	26047160	Down	1.9	0.0126	0.0532							cadherin 17
4	AJ209669	Gnb1	1382962 at	172386934	Down	1.7	0.0035	0.0331							Rattus norvegicus transcribed sequences
4	BF396154	-	1378142 at	129921739	Down				1.6	0.0198	0.0632				Rattus norvegicus transcribed sequences
4	AJ103530	-	1382431 at	70490031	Down				1.6	0.0041	0.0365				similarity to protein sp.O95477 (H. sapiens) ABC1_HUMAN ATP-binding cassette, sub-family A, member 1 (ATP-binding cassette transporter 1)
4	BF397628	-	1391139 at	105474702	Down				1.6	0.0387	0.0857				Rattus norvegicus transcribed sequences
4	BE115159	-	1384059 at	102264948	Down				1.6	0.0000	0.0109				Rattus norvegicus transcribed sequences
4	AA874945	-	1382967 at	12379423	Down	1.5	0.0128	0.0535							Rattus norvegicus transcribed sequences
4	AA958428	-	1381895 at	96509955	Down	1.6	0.0062	0.1626							Rattus norvegicus transcribed sequences
4	BF396286	-	1397015 at	57765343	Down	2.1	0.0309	0.1906							Rattus norvegicus transcribed sequences
4	BE116294	-	1396263 at	51807151	Down	1.8	0.0098	0.1324							Rattus norvegicus transcribed sequences
4	BI303268	Bzw2	1370298 at	94817272	Down	1.7	0.0444	0.2103							HbZ protein
4	BI378236	-	1372564 at	40265862	Down	1.6	0.0330	0.1971							Rattus norvegicus similar to Kiso0575 (LOC313977). mRNA
4	BM384008	-	1382864 at	72885228	Down	1.6	0.0020	0.0604							Rattus norvegicus similar to p190-B (LOC299012). mRNA
4	NM_019163	Psen1	1388782 at	107725733	Down	1.6	0.0091	0.2103							presenilin 1
4	BM385731	-	1385609 at	42217273	Up	2.2	0.0034	0.0764	2.1	0.0118	0.0383				presenilin 1
4	BM383749	-	1378128 at	42262080	Up	2.0	0.0183	0.1640	1.6	0.0386	0.0746				

Table 1

Chromosome	Gene Identifier	Gene ID	Other ID	Position (Mb)	Direction	Ratio	Week4		Week12		Week20		Gene Description		
							p-value	adjp-value	Ratio	p-value	adjp-value	Ratio		p-value	adjp-value
11	AA989697	-	1384122 a.t	70386039	Down						1.6	0.0204	0.0639	Rattus norvegicus transcribed sequences	
11	AA650765	-	1384724 a.t	25263390	Down						1.6	0.0216	0.047	Rattus norvegicus transcribed sequences	
11	BG669292	-	1388879 a.t	44854445	Down						1.6	0.0313	0.0764	Rattus norvegicus similar to 503341B22Rik protein (LOC363767), mRNA	
11	BF561284	-	1393285 a.t	2390377	Down						1.6	0.0251	0.0709	---	
11	BE105691	-	1390443 a.t	17441730	Down						1.6	0.0343	0.0834	Rattus norvegicus transcribed sequences	
11	U31584	Laamp	1370550 a.t	60165324	Down						1.6	0.0003	0.0173	limb3-associated membrane protein	
11	BM339685	-	1378937 a.t	43764257	Down						1.6	0.0497	0.0986	Rattus norvegicus similar to downregulated in ovarian cancer 1 (LOC304020), mRNA	
11	BI303277	-	1382025 a.t	2390377	Down						1.6	0.0406	0.0726	Rattus norvegicus transcribed sequences	
11	BE109290	Digh1	1397135 a.t	70775877	Down						1.5	0.0192	0.0630	Rattus norvegicus transcribed sequences	
11	BF368632	-	1385742 a.t	86161377	Down						1.5	0.0400	0.0876	---	
12	NM_053962	Ssa	1389884 a.t	137281250	Down	1.9	0.0278	0.1860						serine dehydratase	
12	BE103748	-	1392941 a.t	22421151	Down	1.6	0.0297	0.1894						Rattus norvegicus transcribed sequences	
12	AW531730	-	1394764 a.t	10757471	Up						1.7	0.0108	0.0588	Rattus norvegicus similar to kinase phosphatase inhibitor 2 (LOC304286), mRNA	
12	AI235340	-	1384634 a.t	43273320	Down						1.7	0.0309	0.0640	Rattus norvegicus transcribed sequences	
12	BF288115	-	1376688 a.t	19885349	Down						1.6	0.0148	0.0432	Rattus norvegicus similar to paired immunoglobulin-like type 2 receptor alpha, cell surface receptor FDF03 (LOC288552), mRNA	
12	BF521655	-	1385670 a.t	37302260	Up						1.5	0.0184	0.0622	Rattus norvegicus similar to serine dehydratase related sequence 1 (LOC360816), mRNA	
12	AI995117	-	1381431 a.t	84144	Down						1.5	0.0336	0.0802	Rattus norvegicus transcribed sequences	
13	NM_031118	Seaf1	1399002 a.t	71420882	Down	2.3	0.0402	0.2103						Rattus norvegicus transcribed sequences	
13	NM_053843	Fcgr3	1398246 a.t	86809011	Down	2.0	0.0370	0.2071						Rattus norvegicus similar to Fc receptor, IgG, low affinity III	
13	AI231626	Ras5	1393167 a.t	85445325	Down	1.6	0.0342	0.2009						regulator of G-protein signaling 5	
13	AA894335	-	1375590 a.t	46545034	Down	1.6	0.0048	0.0932						Rattus norvegicus similar to RIKEN cDNA 5730454B08 (LOC310649), mRNA	
13	AI704885	-	1381542 a.t	40948540	Down	1.5	0.0062	0.0977						Rattus norvegicus similar to RIKEN cDNA 1300013G12 (LOC304766), mRNA	
13	BF402061	Capon	1398843 a.t	86360648	Down				1.8	0.0132	0.0407			Rattus norvegicus transcribed sequences	
13	AI717113	FS	1374300 a.t	79992013	Down				1.8	0.0237	0.0563			coagulation factor 5	
13	AI406858	-	1390987 a.t	86139707	Up						1.5	0.0142	0.0551	Rattus norvegicus transcribed sequences	
13	AI407002	-	1373908 a.t	86360648	Down				2.2					Rattus norvegicus transcribed sequences	
13	BF288190	Ptprc	1390789 a.t	51247016	Down				1.9	0.0293	0.0742			protein tyrosine phosphatase, receptor type, C	
13	AA803579	-	1392794 a.t	86131914	Down				1.8	0.0091	0.0493			Rattus norvegicus transcribed sequences	
13	BF417626	-	1393187 a.t	89545247	Down				1.8	0.0416	0.0898			Rattus norvegicus transcribed sequences	
13	AI339193	-	1379387 a.t	58927898	Down				1.7	0.0204	0.0609			Rattus norvegicus transcribed sequences	
13	BF406841	-	1377301 a.t	48378522	Down				1.6	0.0032	0.0327			Rattus norvegicus transcribed sequences	
13	BE108956	-	1392489 a.t	86360648	Down				1.6	0.0184	0.0622			Rattus norvegicus transcribed sequences	
13	AA943075	-	1382388 a.t	57588620	Down				1.6	0.0028	0.0327			Rattus norvegicus transcribed sequences	
13	BI275261	-	1379497 a.t	237646	Down				1.6	0.0003	0.0173			Rattus norvegicus transcribed sequences	
13	BM390782	-	1384972 a.t	57588620	Down				1.6	0.0051	0.0392			Rattus norvegicus transcribed sequences	
13	NM_031170	Sae	1381763 a.t	89112728	Down				1.5					serum amylid D-component	
13	AI012859	Rnasase	1380425 a.t	68822011	Down				1.5	0.0419	0.0888			ribonuclease L (2, E_colloplasmid-encoded symbiote-dependent)	
13	BC369581	-	1371472 a.t	85484906	Down				1.5	0.0031	0.0327			Rattus norvegicus transcribed sequences	
14	AA826001	-	13882181 a.t	118862198	Down	1.5	0.0014	0.0604						Rattus norvegicus transcribed sequence with weak similarity to protein sc008116 (R.norvegicus) WASL, RAT Neural Wiskott-Aldrich syndrome protein (N-WASP)	
14	BI290769	-	1382063 a.t	81988783	Down	1.6	0.0062	0.0977						Rattus norvegicus hypothetical LOC305452 (LOC305452), mRNA	
14	AA828254	-	1383930 a.t	14995498	Down				2.1	0.0176	0.0484			small inducible cytokine B subfamily (Cys-X-Cys motif), member 13 (B-cell chemokine/tractant); B-cell-homolog chemokine (knakin for Burkitts lymphoma re	
14	BI285076	-	1377286 a.t	10049664	Down				1.6	0.0054	0.0406			Rattus norvegicus transcribed sequences	
14	BI285078	-	1376870 a.t	83909631	Up				1.6	0.0157	0.0575			Rattus norvegicus similar to transcription factor MAZR2 (LOC305471), mRNA	
14	BI289418	Cd38	1393025 a.t	72320479	Down				1.7	0.0489	0.0966			CD38 antigen	
14	AB001382	Sept1	1387581 a.t	84423008	Down				1.7	0.0428	0.0893			recruited phosphoinositid	
14	AA848699	-	1384044 a.t	39959936	Down				1.6	0.0163	0.0544			Rattus norvegicus similar to beta-galactosidase (LOC305305), mRNA	
14	BF523573	-	1388061 a.t	110484799	Down				1.6	0.0307	0.0760			Rattus norvegicus transcribed sequences	
14	BE110655	-	1388317 a.t	87224018	Down				1.5	0.0431	0.0893			Rattus norvegicus transcribed sequences	
14	BF391447	-	1392162 a.t	56994709	Down	1.7	0.0148	0.1530						Rattus norvegicus transcribed sequences	
15	BM387164	-	1378087 a.t	56803906	Down				1.6	0.0243	0.0572			Rattus norvegicus transcribed sequences	
15	AI111212	Pdim2	1375367 a.t	90594108	Up				1.7	0.0093	0.0493			Rattus norvegicus similar to PDZ and LIM domain 2 (LOC290354), mRNA	
15	AI15544	-	1383392 a.t	39495179	Down				1.7	0.0466	0.0946			Rattus norvegicus transcribed sequences	
15	AA824588	-	1392725 a.t	108030444	Up				1.7	0.0466	0.0946			Rattus norvegicus similar to cDNA sequence BC006662 (LOC290500), mRNA	
15	NM_012770	Gucy1b2	1387719 a.t	41819400	Up				1.6	0.0284	0.0723			guanylate cyclase, soluble, beta 2	
15	AA892699	-	1392967 a.t	87382731	Down				1.6	0.0187	0.0629			Rattus norvegicus transcribed sequences	
15	NM_053907	Dnase1l3	1368204 a.t	18900363	Down				2.2	0.0058	0.0417			deoxyribonuclease Like 3	
15	BI378310	-	1385099 a.t	87547816	Down				1.8	0.0073	0.0448			similarity to protein ref NP_033844 (H.sapiens) sciiellin (Homo sapiens)	
15	BI303023	-	1383708 a.t	108785601	Down				1.7	0.0031	0.0327			strong similarity to protein ref NP_000782.1 (H.sapiens) integrin, beta-like 1 (with EGF-like repeat domains) [Homo sapiens]	
15	AF314657	Clu	1367784 a.t	45399691	Down				1.6	0.0338	0.0802			clustarin	
15	BE104961	-	1379538 a.t	35487815	Down				1.5	0.0365	0.0837			Rattus norvegicus similar to centromere protein J, centrosomal P4.1-associated protein, LYST-interacting protein LIP1, LAG-3-associated protein (LOC305908), mRNA	
15	AI137275	-	1375115 a.t	10333818	Up				1.6	0.0256	0.1821			Rattus norvegicus similar to wingless apart-like CG3707-PA (LOC290577), mRNA	
16	AF209114	LOC192253	1388040 a.t	83812189	Down	1.6	0.0256	0.1821						myosin heavy chain Myr 8	
16	LIJ2323	Nrq1	1370667 a.t	63053630	Down	1.5	0.0468	0.2103						neurexin 1	
16	AI190253	-	1379471 a.t	20463328	Up				1.9	0.0266	0.0582			Rattus norvegicus similar to otupulin protein family member (XC177) (LOC296386), mRNA	
16	AI710682	-	1371354 a.t	6639336	Up				1.7	0.0028	0.0327			Rattus norvegicus similar to troponin C, cardiac muscle - mouse (LOC290961), mRNA	
16	AI030212	-	1383031 a.t	62307869	Up				1.6	0.0096	0.0493			---	
16	AA892204	-	1386382 a.t	5892391	Up				1.5	0.0372	0.0846			Rattus norvegicus transcribed sequences	
16	BE118080	LOC207125	1371091 a.t	83402975	Up				1.5	0.0178	0.0614			unknown protein	
16	BI303853	-	1381566 a.t	30959054	Down				1.8	0.0416	0.0888			Rattus norvegicus transcribed sequences	
16	NM_131328	Cpe	1386921 a.t	52049664	Down				1.9	0.0557	0.0831			carboxypeptidase E	
16	AI237688	-	1382601 a.t	56307360	Down				1.7	0.0050	0.0392			Rattus norvegicus transcribed sequences	
17	BF388256	-	1381316 a.t	41225418	Up	1.9	0.0346	0.2016						Rattus norvegicus transcribed sequences	
17	BF397269	-	1388111 a.t	8748412	Down	2.3	0.0380	0.2095						Rattus norvegicus LOC361197 (LOC361197), mRNA	
17	BF553179	-	1383759 a.t	11962085	Down	2.1	0.0112	0.1393						Rattus norvegicus similar to nuclear ATP/GTP-binding protein (LOC290986), mRNA	
17	AA979591	-	1386180 a.t	47529887	Down	1.7	0.0019	0.0604						Rattus norvegicus similar to leucine rich repeat containing 16 (LOC369841), mRNA	
17	AA997590	-	1385248 a.t	20881962	Down	1.6	0.0487	0.2103		1.7	0.0495	0.0886			Rattus norvegicus transcribed sequence with moderate similarity to protein prc35272 (H.sapiens) B35272 osteoinductive factor - human
17	AV140475	Cuob2	1388169 a.t	83023864	Down				1.6	0.0148	0.0614			CUO-1 related RNA-binding protein 2	
17	AI029126	-	1379518 a.t	33137357	Up				1.6	0.0009	0.0232			Rattus norvegicus transcribed sequences	
17	AI007775	-	1393348 a.t	36881748	Up				1.6	0.0059	0.0419			Rattus norvegicus transcribed sequences	
17	AA920109	-	1393489 a.t	57290713	Down				3.0	0.0183	0.0622			similarity to protein pdb 1LBG (E. coli) B Chain B, Lactose Operon Repressor Bound To 21-Base Pair Symmetric Operator Dna, Alpha Carbon Only	
17	BI294018	-	1392648 a.t	86361701	Down				2.1	0.0449	0.0923			Rattus norvegicus similar to macrophage mannose receptor precursor (LOC2	

Table 1

Chromosome	Gene Identifier	Gene ID	Other ID	Position (Mb)	Direction	Ratio	Week4 p-value	adjp-value	Ratio	Week12 p-value	adjp-value	Direction	Week20 p-value	adjp-value	Gene Description
19	BE108555	ZD106	1386661	40365883	Up							1.6	0.0123	0.0528	Rattus norvegicus similar to Ddx19 protein (LOC292022). mRNA
19	BE108538	-	1379079	40389933	Up							1.6	0.0408	0.0886	Rattus norvegicus transcribed sequences
19	AI575458	-	1383083	52213643	Up							1.5	0.0105	0.0493	Rattus norvegicus similar to SMART1 (LOC292064). mRNA
19	BI289813	-	1372800	at	Up							1.5	0.0028	0.0327	---
19	AI071216	-	1383973	at	Up							1.5	0.0215	0.0647	Rattus norvegicus similar to hypothetical protein E330010G16 (LOC307816). mRNA
19	BF414702	-	1383641	at	Down							1.6	0.0420	0.0888	endothelin receptor type A
19	BI275887	-	1360733	at	Down							1.6	0.0431	0.0893	Rattus norvegicus transcribed sequences
19	BF448891	-	1394145	at	Down							1.6	0.0062	0.0428	endothelin receptor type A
20	AI012393	Bmi7	1376373	at	Up	1.6	0.0446	0.2103							similarity to protein sp-P14373 (H.sapiens) RFP_HUMAN Zinc-finger protein RFP (Ret finger protein) (Tripartite motif protein 27)
20	BE096729	Fhd	1387877	at	Down	1.6	0.0040	0.0809							forminotransferase cyclodeaminase
20	AI599350	Phm59	1370188	at	Down	1.6	0.0048	0.0922							proteasome (prosome, macropain) subunit, beta type 9
20	AW252296	Col18a1	1381431	at	Down	1.6	0.0000	0.0216							Rattus norvegicus transcribed sequences
20	BE108058	Col18a1	1388394	at	Down							1.6	0.0260	0.0709	Rattus norvegicus transcribed sequences
20	AJ249375	RT1.S9	1388213	at	Down	1.6	0.0489	0.2103							RT1 class Ib RT1.S9
20	AI548626	-	1388128	at	Down	1.5	0.0176	0.1640							Rattus norvegicus transcribed sequences
20	BI290161	-	1382041	at	Up										---
20	AI113218	Sagf1	1382963	at	Up							1.7	0.0019	0.0325	Rattus norvegicus similar to 1-acylglycerol-3-phosphate-gamma (LOC294324). mRNA
20	AF413572	Pkb	1389105	at	Down							2.4	0.0091	0.0493	serine/threonine-1-phosphatase lyase 1
20	AI411352	Gia1	1372002	at	Down							1.8	0.0299	0.0760	protein kinase (GAMF-dependent, catalytic) inhibitor beta
20	AW326939	-	1382736	at	Down							1.8	0.0136	0.0546	gap junction membrane channel protein alpha 1
20	BM383464	-	1382437	at	Down							1.7	0.0124	0.0528	RT1 class Ib gene(Aw2)
20	NM_019309	Grik2	1369036	at	Down							1.5	0.0013	0.0282	glutamate receptor, ionotropic, kainate 2
X	AI189241	-	1372637	at	Up	1.9	0.0459	0.2103							Rattus norvegicus transcribed sequence with weak similarity to protein pir138488 (H.sapiens) 138488 trophinin - human
X	BG371620	-	1383675	at	Up	1.5	0.0481	0.2103							---
X	NM_053428	Fgf13	1368114	at	Up	1.5	0.0351	0.2016							fibroblast growth factor 13
X	BF396777	MGC94181	1395201	at	Up	1.5	0.0227	0.1748							Rattus norvegicus transcribed sequences
X	AI170324	-	1373862	at	Down	1.5	0.0057	0.0977							similarity to protein sp-P97946 (M.musculus) VEGD_MOUSE Vascular endothelial growth factor D precursor (VEGF-D) (c-fos induced growth factor) (FIGF)
X	NM_013197	Aaa2	1367985	at	Down				2.7	0.0201	0.0523	1.9	0.0341	0.0803	aminolevulinic acid synthase 2
X	NM_021870	Bmp16	1387734	at	Up							1.9	0.0425	0.0893	bone morphogenetic protein 15
X	NM_012521	Cals3	1369339	at	Up							1.8	0.0247	0.0794	calpain 3
X	BF389476	MGC94465	1380835	at	Up							1.6	0.0133	0.0546	Rattus norvegicus similar to hypothetical protein FLJ12671 (LOC317205). mRNA
X	BG379491	-	1374218	at	Up							1.8	0.0007	0.0216	Rattus norvegicus transcribed sequences
X	BI274139	-	1375276	at	Up							1.6	0.0019	0.0325	Rattus norvegicus similar to mitochondrial inner membrane translocase component Tim17b (LOC317374). mRNA
X	NM_017251	Gp1	1387145	at	Up							1.6	0.0050	0.0392	gap junction membrane channel protein beta 1
X	BG373822	-	1376706	at	Down							1.6	0.0486	0.0986	Rattus norvegicus transcribed sequences
X	BF413976	-	1389693	at	Down							1.6	0.0482	0.0982	---
UN	BM388500	-	1383977	at	Up	1.9	0.0252	0.1818							---
UN	BF551160	-	1394056	at	Down	1.7	0.0299	0.1884							---
UN	LI15600	Nr1d2	1370560	at	Up				2.2	0.0086	0.0318				nuclear receptor subfamily 1, group D, member 2
UN	LI20796	Nr1d2	1370541	at	Up				1.7	0.0149	0.0432				nuclear receptor subfamily 1, group D, member 2
UN	BF284190	Nr1d2	1390430	at	Up				1.6	0.0380	0.0742				---
UN	AI237240	-	1395238	at	Down							2.7	0.0253	0.0674	---
UN	BE119819	Rap2b	1392622	at	Down				1.7	0.0259	0.0574				Rattus norvegicus transcribed sequences
UN	AF361355	Pr1	1388103	at	Up		1.5	0.0048	0.0225	1.7			0.0026	0.0327	voltage-dependent calcium channel gamma subunit-like protein
UN	AF121791	-	1384303	at	Up							1.7	0.0376	0.0846	---
UN	AI717791	-	1384302	at	Up							1.6	0.0400	0.0391	---
UN	BF418508	Ahp10	1389441	at	Down							1.5	0.0010	0.0238	annexin V-binding protein ABP-10

Table2

Week	Total Networks	Top Networks	High Level Functions	Significance	Top Function	Significance	Canonical Pathway (p<0.05)	Node
4	10	1	Immune Response	3.87E-8 - 2.88E-2	Development Disorder	3.46E-15 - 3.15E-5	Wnt/B-catenin Signaling	Crebbp(2), Csnk1e(1), Sfrp2(3), Wnt2b(3)
			Cell-to-Cell Signaling and Interaction	1.27E-6 - 2.88E-2	Cellular Growth and Proliferation	2.09E-14 - 5.88E-5	Amyloid Processing	Csnk1e(1), Psen(1)
			Immune and Lymphatic Development and Function	1.27E-6 - 2.88E-2	Cell Death	1.87E-13 - 5.88E-5	Antigen Presentation	HLA-E(1), Psmb9(1)
					Cellular Development	8.90E-13 - 6.36E-5		
		2	Gene Expression	1.15E-7 - 3.84E-3				
			Viral Function	6.30E-6 - 3.84E-3				
			Cancer	1.43E-5 - 3.84E-3				
		3	Cellular Development	8.04E-8 - 3.84E-3				
			Developmental Disorder	2.64E-7 - 3.84E-3				
			Cellular Growth and Proliferation	2.87E-7 - 3.84E-3				
12	6	1	Gene Expression	2.68E-10 - 6.60E-5	Gene Expression	8.19E-13 - 1.04E-5	Wnt/B-catenin Signaling	Tcf4(2), Tgfb3(1)
			Cell-to-Cell Signaling and Interaction	9.15E-10 - 8.57E-5	Cell-to-Cell Signaling and Interaction	2.71E-11 - 8.91E-6	Glutathione Metabolism	Gstm1(2), Gstm2 (2)
			Cellular Growth and Proliferation	9.15E-10 - 8.57E-5	Cellular Growth and Proliferation	2.71E-11 - 1.25E-5	Glycine, Serine, and Threonine Metabolism	Alas2(1), Sars
		2	Cardiovascular Disease	5.17E-7 - 3.67E-3	Connective Tissue Development and Function	2.71E-11 - 3.42E-6		
			Drug Metabolism	5.17E-7 - 1.46E-3				
			Hematological Disease	5.17E-7 - 3.84E-3				
			Small Molecule Biochemistry	5.17E-7 - 3.84E-3				
20	10	1	Cell Morphology	2.07E-7 - 3.84E-3	Cellular Growth and Proliferation	3.40E-11 - 4.16E-5	Nitric Oxide Signaling in Cardiovascular System	Gucy1a3(3), Gucy1b3(3)
			Cell Cycle	1.03E-6 - 3.84E-3	Connective Tissue Development and Function	7.21E-11 - 4.11E-5		
			Endocrine System Development and Function	1.03E-6 - 2.36E-3	Cellular Movement	8.58E-11 - 3.60E-5		
			Cellular Development	2.18E-6 - 3.49E-3	Immune and Lymphatic Development and Function	1.49E-10 - 4.18E-5		
		2	Gene Expression	1.99E-8 - 9.24E-4				
			Cell Morphology	5.20E-8 - 1.43E-5				
			Cell-to-Cell Signaling and Interaction	2.07E-7 - 9.54E-4				
		3	Cellular Movement	1.31E-8 - 3.16E-3				
			Cellular Development	1.31E-7 - 3.84E-3				
			Immune and Lymphatic Development and Function	1.31E-7 - 3.84E-3				

Table3

Chromosome	Gene Identifier	Gene ID	Other ID	Position (Mb)	Gene Description
1	NM_021663	NUC52_RAT	1370000_at	174624096	NEFA precursor
1	I02827	LOC8A_RAT	1370897_at	90337907	branched chain keto acid dehydrogenase subunit E1, alpha polypeptide
1	A1169562	Q8374_RAT	1371080_at	94412803	kallikrein
1	BF389640	QOAG20_RAT	1373485_at	267369410	Rattus norvegicus similar to eukaryotic translation initiation factor 3, subunit 10 (theta, 150/170kDa; eukaryotic translation initiation factor 3, subunit 10, 170kD); Eukaryotic translation initiation factor 3, subunit 10, 170kD; eukaryotic tr
1	BE099904		1381264_at	264689833	Rattus norvegicus transcribed sequences
1	AA020133	XP_218291_3	1381118_x_at	12076570136	Rattus norvegicus similar to protein pb; ILBG (E. coli) B Chain B, Lactose Operon Repressor Bound To 21-Base Pair Symmetric Operator Dna, Alpha Carbons Only
1	AA956308	XP_218291_3	1381118_x_at	72699408	Rattus norvegicus similar to RIKEN cDNA 2810439M05 (LOC208361), mRNA
1	BF39429	XP_215000_3	1380226_at	149425862	Rattus norvegicus similar to RIKEN cDNA 4933417L02 (LOC293117), mRNA
1	A023296	Q2V057_RAT	1389845_at	85494812	Rattus norvegicus similar to proline dehydrogenase (oxidase) 2 (LOC361538), mRNA
1	AA064142	XP_344925_2	1389889_at	1576372994	Rattus norvegicus similar to RIKEN cDNA 2810209A20 (LOC368121), mRNA
1	AA121274	C071_RAT	1390801_at	228097480	Rattus norvegicus transcribed sequences
1	AA145746		1392249_at	20355444	Rattus norvegicus transcribed sequences
1	AA893579		1392794_at	203379419	Rattus norvegicus transcribed sequences
1	AA819959		1392982_at	101810286	Rattus norvegicus transcribed sequences
1	BF285952	NP_001034754.1	1393474_at	98197898	Rattus norvegicus similar to cytosolic sulfotransferase (LOC292915), mRNA
1	AA045155	NP_001031363.1	1393929_at	47115584	Rates norvegicus similar to G protein-coupled receptor KY411 (LOC308163), mRNA
1	BF294064		1395544_at	103984540	Rattus norvegicus similar to methylcrotonyl-Coenzyme A carboxylase 1 (alpha) (LOC294972), mRNA
1	BE097460		1398700_at	40889901	Rattus norvegicus transcribed sequences
2	NM_031620	SERA_RAT	1367811_at	124600017	3-phosphoglycerate dehydrogenase
2	NM_017090	GCYA3_RAT	1368154_at	173755007	guanylate cyclase 1, soluble, alpha 3
2	AF202115	ICERU_RAT	1368419_at	10508278	Juvenile hemin
2	NM_012889	VICAM1_RAT	1368474_at	212277654	vascular cell adhesion molecule 1
2	AF367210	IL7_RAT	1369208_at	96356083	interleukin 7
2	U78897		1369886_at	144267748	activity and neurotransmitter-induced early gene protein 4 (ania-4)
2	BE411765	CELRF2_RAT	1371018_at	203905254	cadherin EGF LAG seven-pass C-type receptor 2
2	BI281702	MAP1B_RAT	1373263_at	30415255	microtubule-associated protein 1b
2	AA176320	XP_226988.3	1373416_at	113315728	Rattus norvegicus transcribed sequences
2	BE107853		1373497_at	142404903	Rattus norvegicus transcribed sequences
2	AA087384		1373778_at	47172864	Rattus norvegicus transcribed sequences
2	BF388224		1379175_at	234240301	Rattus norvegicus transcribed sequences
2	BG374101		1381262_at	181637899	Rattus norvegicus transcribed sequences
2	AA819288		1382274_at	157332263	Rattus norvegicus similar to Retinoid acid receptor responder protein 1 (Tazarotene-induced gene 1 protein) (RAR-responsive protein TIG1) (LOC310486), mRNA
2	BM334664		1382437_at	208517037	RIT class B gene/ha2
2	AA511405	XP_227217.3	1385407_at	155315259	Rattus norvegicus similar to hypothetical protein FLJ22693 (LOC310467), mRNA
2	A0237532	XP_227203.2	1385872_at	162126352	Rattus norvegicus transcribed sequences
2	BF280410	XP_226824.3	1386952_at	58105666	Rattus norvegicus transcribed sequences
2	BF294922		1388857_at	119296979	Rattus norvegicus transcribed sequences
2	BI282008		1389003_at	2911603	Rattus norvegicus transcribed sequences
2	BF402235		1392262_at	204000799	Rattus norvegicus transcribed sequences
2	BF409007		1393563_at	116159619	Rattus norvegicus transcribed sequences
2	BF411556		1394363_at	224263853	Rattus norvegicus transcribed sequences
2	BF553729	QOQXK5_RAT	1397820_at	200218630	Rattus norvegicus transcribed sequences
3	NM_032066	NP_114455.1	1398051_at	79436634	smooth muscle-specific 17 beta-hydroxysteroid dehydrogenase type 3
3	NM_053498	ENP8_RAT	1398315_at	141384540	ectonucleoside triphosphate diphosphotriphosphate 6
3	BM369019	NP_114013.1	1398829_at	112608480	fibrin-1
3	BI295949	NP_001011965.1	1373590_at	14393734	Rattus norvegicus transcribed sequences
3	BF284124	XP_230325.3	1377018_at	87795505	Rattus norvegicus similar to E43000205Rik protein (LOC311252), mRNA
3	AA273762	XP_216007.3	1379394_at	4180013	Rattus norvegicus similar to G333139-PA (LOC296580), mRNA
3	BI294934		1380612_at	169762453	Rattus norvegicus transcribed sequences
3	AA527286		1380899_at	143411116	Rattus norvegicus similar to Protein C20orf160 (LOC311550), mRNA
3	BF553538	CXA9_RAT	1386123_at	99861095	Rattus norvegicus transcribed sequences
3	BF563766		1386995_at	4206895	Rattus norvegicus transcribed sequences
4	AA011345	WNK1_RAT	1398140_at	156297841	protein kinase, vaine deficient 1
4	NM_013080	PTPRZ_RAT	1388350_at	49334202	protein tyrosine phosphatase, receptor-type, Z polypeptide 1
4	BI291846		1371969_at	52062872	caldesmon 1
4	BF537890	MET_RAT	1374665_at	43154101	Rattus norvegicus transcribed sequences
4	AA012630	PTGD2_RAT	1377012_at	94611607	Rattus norvegicus transcribed sequences
4	AA529483	NP_997677.1	1378365_at	117432385	Rattus norvegicus transcribed sequences
4	BF389527		1381723_at	152898769	---
4	BF289769	NP_001034433.1	1383401_at	42700914	Rattus norvegicus transcribed sequences
4	BE116205	XP_242556.3	1384795_at	118125137	Rattus norvegicus similar to nuclear protein, NP220 (LOC312491), mRNA
4	AA524822		1390965_at	49659289	Rattus norvegicus transcribed sequence with moderate similarity to protein ref:NP_036191.1 (M.musculus) Ca2+-dependent activator protein for secretion; Ca2+-dependent activator protein for secretion [Mus musculus]
4	AA639113		1392553_at	88666142	Rattus norvegicus transcribed sequence with moderate similarity to protein pir:AS7384 (H.sapiens) A57384 multimerin, endothelial cell, precursor - human
4	BF283618		1392610_at	62792942	Rattus norvegicus similar to hypothetical protein B290314D19 (LOC626236), mRNA
4	BF400819		1396602_at	5272676	Rattus norvegicus transcribed sequences
5	NM_053977	CAD17_RAT	1369224_at	26047160	cadherin 17
5	NM_021653	ID01_RAT	1369259_at	123835247	iodotransin, iodothyronine, type I
5	BF283766	NP_852143.1	1373023_at	2286373	Rattus norvegicus transcribed sequences
5	BF284337		1373063_at	79494958	Rattus norvegicus similar to RIKEN cDNA 2810435D12 (LOC298097), mRNA
5	BE109524		1390487_at	61961151	Rattus norvegicus transcribed sequences
5	BM394214	NP_001069980.1	1138059_at	144905644	Rattus norvegicus similar to Matrin 1, cartilage matrin protein 1 (LOC297894), mRNA
5	AA028979	QSBK25_RAT	1383395_at	160648328	Rattus norvegicus similar to Argmatinase, mitochondrial precursor (Argmatine urechydrosase) (AUH) (LOC298607), mRNA
5	AA547423	BSND_RAT	1393209_at	127542219	bartin
5	BF401989		1396254_at	5249162	Rattus norvegicus transcribed sequences
5	GC378899		1399992_at	148615030	Rattus norvegicus similar to hypothetical protein FL110315 (LOC362608), mRNA
6	BG378238		1372524_at	40251362	Rattus norvegicus similar to Kiaa0575 (LOC313977), mRNA
6	BE107255	XP_234275.3	1374480_at	94024421	Rattus norvegicus similar to hypothetical protein E130308H01 (LOC314212), mRNA
6	BG667451		1377185_at	101599793	Rattus norvegicus transcribed sequences
6	AA103284	XP_216791.3	1381376_at	136186929	Rattus norvegicus similar to Trifid2 protein (LOC299340), mRNA
6	BM391545		1392733_at	77959323	Rattus norvegicus transcribed sequences
6	BM384008	XP_216688.3	1392864_at	72685228	Rattus norvegicus similar to p190-B (LOC299012), mRNA
6	AA819288		1371776_at	53801895	phosphatidylinositol 3-kinase, regulatory subunit, polypeptide 1
7	BF351206	QOQY87_RAT	1176542_at	1376542	nuclear factor I2
7	BI298372		1375467_at	140961650	Rattus norvegicus similar to orly1 TD-element binding protein 7 (LOC366994), mRNA
7	BM384525	XP_218859.1	1395045_at	16321201	Rattus norvegicus similar to NADH oxidoreductase (LOC299643), mRNA
8	NM_052963	MMP12_RAT	1398530_at	4249938	matrix metalloproteinase 12
8	BI291272		1373712_at	6644972	Rattus norvegicus transcribed sequences
8	BF405592		1377531_at	32289929	Rattus norvegicus transcribed sequences
8	AA997421	XP_343612.2	1379721_at	21058321	Rattus norvegicus similar to 2310047B19Rik protein (LOC363029), mRNA
8	BF542912	SCN5B_RAT	1383435_at	42321453	Rattus norvegicus similar to sodium ion channel protein, type I, form 1 (LOC367298), mRNA
8	NM_139422	NP_569108.1	1387305_at	2039306	caspase 12
8	NM_021866	CCR2_RAT	1387742_at	128892784	chemokine receptor CCR2 gene
8	AA893518	UP3_RAT	1389270_x_at	36892614	Rattus norvegicus similar to Urinary protein 3 precursor (RUP-3) (LOC300504), mRNA
8	AA896918	QR9R2_RAT	1390710_x_at	44869380	Rattus norvegicus similar to gp250 precursor (LOC300652), mRNA
8	BF408557		1391021_at	76091646	Rattus norvegicus similar to KIAA1749 protein (LOC315756), mRNA
9	BI275716	CODA1_RAT	1370959_at	44281582	collagen, type III, alpha 1
9	AA527151		1374635_at	62252515	Rattus norvegicus transcribed sequence with strong similarity to protein ref:NP_003003.1 (H.sapiens), a disintegrin and metalloproteinase domain 23 preproprotein [Homo sapiens]
9	AA385237		1378346_at	110452096	Rattus norvegicus transcribed sequences
9	AA639159	XP_340662.2	1386454_at	74409235	Rattus norvegicus similar to YSP-1 form 1 (LOC367298), mRNA
9	BM384831		1393653_at	83412793	Rattus norvegicus transcribed sequences
9	BF523991		1396268_at	8425233	Rattus norvegicus similar to putative nuclear protein (LOC301570), mRNA
9	BE094410		1396170_at	73814131	Rattus norvegicus transcribed sequences
9	AA08959		1396942_at	13460537	Rattus norvegicus hypothetical LOC301287 (LOC301287), mRNA
10	AA10414		1371545_at	96056298	Rattus norvegicus transcribed sequences
10	BI285135	HCPI1_RAT	1374073_at	64576051	Rattus norvegicus similar to DNA segment, Chr 11, ERATO Doi 18, expressed (LOC303333), mRNA
10	BE114669	XP_340743.2	1374208_at	4372981	Rattus norvegicus similar to RIKEN cDNA AA30104C18 (LOC364668), mRNA

Table3

10	B1298463		1374356_at	15207723	Rattus norvegicus similar to tripartite motif protein 32 (LOC302999), mRNA
10	BC390279		1375672_at	62782723	Rattus norvegicus transcribed sequence with weak similarity to protein pR:RGECDW (E. coli) RGECDW transcription activator of D-serine dehydratase - Escherichia coli
10	B1294949	NP_001008361.1	1375674_at	10911559	Rattus norvegicus similar to chromosome 16 open reading frame 5 (LOC360480), mRNA
10	BE100978		1379644_at	40248088	Rattus norvegicus transcribed sequences
10	AW528010		1381233_at	97298015	Rattus norvegicus transcribed sequences
10	BF408881	XP_340881.2	1382957_at	82931188	Rattus norvegicus similar to cisplatin resistance-associated overexpressed protein (LOC360602), mRNA
10	BE112523		1383454_a_at	1110674434	Rattus norvegicus transcribed sequence with moderate similarity to protein ref:NP_071441.1 (H.sapiens) fructosamine-3-kinase [Homo sapiens]
10	AI180413	APOH_RAT	1389350_at	97725096	Rattus norvegicus similar to beta-2-glycoprotein I (LOC287774), mRNA
10	BF419586		1389982_at	72806191	LIM homeobox protein 1
10	BC532455		1390117_at	78207038	Rattus norvegicus transcribed sequences
10	BC671264		1390793_at	99474536	Rattus norvegicus similar to ATP-binding cassette transporter sub-family A member 8a (LOC303638), mRNA
10	AI703949		1397956_at	15129199	---
11	NM_133428	NP_596919.1	1398583_a_at	80230659	histidine-rich glycoprotein
11	BF283405		1397979_at	28558401	Rattus norvegicus transcribed sequence with moderate similarity to protein sp:P00722 (E. coli) BGAL_ECOLI Beta-galactosidase (Lactase)
11	AI909448		1398049_at	89871055	Rattus norvegicus transcribed sequences
11	AI137518	CXAR_RAT	1384816_at	17193505	Rattus norvegicus transcribed sequences
11	AA891032	XP_341011.2	1388503_at	84811466	---
11	BC692292		1388879_at	44854446	---
11	BI303277		1392025_x_at	23905377	Rattus norvegicus similar to 5033411B22Rik protein (LOC363767), mRNA
12	NM_053626	OXDA_RAT	1369491_at	43628160	D-amino acid oxidase
12	AI103040		1376840_at	9688106	Rattus norvegicus transcribed sequences
12	BF553979	QFOI4_RAT	1383446_at	36233484	Rattus norvegicus transcribed sequences
12	BF283000	XP_347254.1	1390983_at	22777345	Rattus norvegicus similar to Williams-Beuren syndrome critical region protein 21 (LOC368083), mRNA
13	BE099979	XP_213921.3	1375428_at	81491327	Rattus norvegicus similar to cellular repressor of E1A-stimulated genes CREG (LOC289185), mRNA
13	BF406641		1377301_at	48379022	Rattus norvegicus transcribed sequences
13	BF380956		1377807_a_at	81083440	Rattus norvegicus transcribed sequence with strong similarity to protein sp:P00722 (E. coli) BGAL_ECOLI Beta-galactosidase (Lactase)
13	AI704885	NP_001012025.1	1381542_at	40948540	Rattus norvegicus similar to RIKEN cDNA 1300013G12 (LOC304766), mRNA
13	BF542890		1394400_at	96950732	Rattus norvegicus transcribed sequences
14	A2325294	XP_223611.1	1376161_at	86999069	Rattus norvegicus similar to RIKEN cDNA 1110014L17 (LOC305502), mRNA
14	BI291600		1377262_at	110170474	Rattus norvegicus similar to KIAA2010 protein (LOC309993), mRNA
14	BI299600		1378313_at	18662686	Rattus norvegicus transcribed sequence with weak similarity to protein sp:O08816 (R.norvegicus) WASL_RAT Neural Wiskott-Aldrich syndrome protein (N-WASP)
14	BF401106	FURB_RAT	1383753_at	110139395	Rattus norvegicus transcribed sequences
14	AI012358		1391303_at	87231825	Rattus norvegicus transcribed sequence with strong similarity to protein ref:NP_150093.1 (H.sapiens) purine-rich element binding protein B [Homo sapiens]
15	NM_052907	DNSL3_RAT	1388294_at	10890283	lecithinase 1-like 3
15	BI303923	ITGB1	1383708_at	108785601	Rattus norvegicus transcribed sequence with strong similarity to protein ref:NP_004782.1 (H.sapiens) integrin, beta-like 1 (with EGF-like repeat domains) [Homo sapiens]
15	A058292	XP_224283.1	1384415_at	43208046	Rattus norvegicus transcribed sequences
15	BC378310		1386509_at	87547816	Rattus norvegicus transcribed sequence with strong similarity to protein ref:NP_003834.1 (H.sapiens) scellin [Homo sapiens]
15	AI861175	NP_001012040.1	1389332_at	2868684	Rattus norvegicus similar to RIKEN cDNA 1700010P07 (LOC305884), mRNA
16	BF283341	TKT_RAT	1376635_at	5908757	transketolase
16	BE114427		1377333_at	3758163	Rattus norvegicus transcribed sequences
16	AI180253		1379411_at	20483828	---
16	AI17260	ARY1_RAT	1367187_a_at	23845894	N-acetyltransferase 1 (arylamine N-acetyltransferase)
17	AI029126		1379518_at	33137857	Rattus norvegicus transcribed sequences
17	AA926109		1383486_at	57240213	Rattus norvegicus transcribed sequence with moderate similarity to protein pdb:1LBG (E. coli) B Chain B, Lactose Operon Repressor Bound To 21-Base Pair Symmetric Operator Dna, Alpha Carbons Only
17	BF560932		1384000_at	42175820	Rattus norvegicus similar to sox-4 protein - mouse (LOC364712), mRNA
18	NM_030858	SMAD7_RAT	1368896_at	72204803	MAD homolog 7 (Drosophila)
18	BM391907		1373692_at	29303561	Rattus norvegicus transcribed sequence with moderate similarity to protein ref:NP_003723.1 (H.sapiens) eukaryotic translation initiation factor 4E binding protein 3; eukaryotic initiation factor 4E-binding protein 3 [Homo sapiens]
18	BI281497	GRAM3_RAT	1384177_at	52202307	Rattus norvegicus similar to RIKEN cDNA 9130427A09 (LOC307288), mRNA
18	A231438	NP_001007688.1	1390569_at	81160881	Rattus norvegicus transcribed sequences
18	AW920883		1394907_at	56545579	Rattus norvegicus transcribed sequences
19	L07316	MDP1_RAT	1388281_at	53503490	dipeptidase 1
19	AY081218	NP_612513.1	1370336_at	49882142	pregnanancy-induced growth inhibitor
19	AIWS3228	IQOIQ3_RAT	1374167_at	35719490	Rattus norvegicus hypothetical LOC291982 (LOC291982), mRNA
19	BF548891	EDNRA_RAT	1393415_at	32048233	endothelin receptor type A
19	BE108555		1396661_at	40906383	Rattus norvegicus similar to Ddx19 protein (LOC292022), mRNA
19	AF413572	IPKB_RAT	1369105_a_at	36655515	protein kinase (cAMP dependent, catalytic) inhibitor beta
20	AB047324	NP_620186.1	1370545_at	44211846	solute carrier family 16, member 10
20	BM383785		1376432_at	8450329	Rattus norvegicus similar to Tex27 protein (LOC361816), mRNA
20	BI290161	XP_215367.2	1382041_at	10731287	Rattus norvegicus similar to 1-acvlqlvorol-3-phosphate-gamma (LOC294324), mRNA
20	BI289995	VPS26_RAT	1382999_at	29709095	Rattus norvegicus similar to vacuolar protein sorting 26; vacuole protein sorting 26; beta 58 (LOC361846), mRNA
20	BE090729	ITCD_RAT	1387877_at	12470290	formin/nucleation factor cytoskeletonase
20	AA901176	XP_215372.3	1398511_at	13435983	Rattus norvegicus similar to Sushi domain (SCR repeat) containing (LOC294335), mRNA
X	AI170324	VEGFD_RAT	1373882_at	50829626	Rattus norvegicus transcribed sequence with moderate similarity to protein sp:P97946 (M.musculus) VEGD_MOUSE Vascular endothelial growth factor D precursor (VEGF-D) (c-fos induced growth factor) (FIGF)
UN	AIWS35280		1370986_s_at	---	Rattus norvegicus retroviral-like c-Ha-ras proto-oncogene mRNA, partial sequence
UN	AI716098		1376132_at	---	Rattus norvegicus transcribed sequences
UN	AI145015		1381418_at	---	Rattus norvegicus transcribed sequences
UN	BE096005		1383665_at	---	---
UN	AI169075		1388526_at	---	---
UN	BF416508		1389441_at	---	annexin V-binding protein ABP-10
UN	BM383122		1389500_at	---	---
UN	BI289762		1390174_at	---	---
UN	A232305		1392001_at	---	---
UN	AW915435		1392554_a_at	---	Rattus norvegicus transcribed sequence with moderate similarity to protein ref:NP_036135.1 (M.musculus) cofactor required for Sp1 transcriptional activation subunit 2 (150 kDa) [Mus musculus]
UN	AI008432		1392843_at	---	Rattus norvegicus transcribed sequences
UN	AA943800		1394127_at	---	Rattus norvegicus transcribed sequences
UN	AI239935		1394526_at	---	Rattus norvegicus transcribed sequences
UN	A232740		1395238_at	---	---
UN	BE116471		1395994_at	---	Rattus norvegicus transcribed sequences
UN	BF298545		1396127_at	---	---
UN	BI281813		1397744_at	---	Rattus norvegicus transcribed sequences

Table4

Number	Gene Identifier	Gene ID	Position (Mb)	Gene Description
1	XM_579199.1	Fga	174737640	174753599 fibrinogen, alpha polypeptide
2	NM_020071.1	Fgb	174768160	174775103 fibrinogen, B beta polypeptide
3	NM_021757.2	Plrg1	174780415	174796746 pleiotropic regulator 1 homolog (Arabidopsis)
4	XM_227313.3	LOC310550	174836071	175056529 similar to protocadherin 16 precursor; protocadherin 16; fibroblast cadherin FIB1; fibroblast cadherin 1; dachous homologue
5	XM_227314.3	Sfrp2	175479310	175486882 secreted frizzled-related protein 2
6	NM_198769.2	Tlr2	175613992	175613992 toll-like receptor 2
7	XM_579995.1	LOC499646	175800216	175801906 LOC499646
8	XM_215595.3	RGD1309007_predicted	175841812	175906450 similar to RIKEN cDNA 2610034E18 gene (predicted)
9	XM_342268.2	Trim2_predicted	175913283	175987825 tripartite motif protein 2 (predicted)
10	XM_215594.3	RGD1311955_predicted	176220891	176253825 similar to CG32384-PA (predicted)
11	NM_021763.1	LOC60382	176275425	176353675 arfaptin 1
12	XM_227318.2	Tigd4_predicted	176369448	176371430 tigger transposable element derived 4 (predicted)
13	XM_579996.1	LOC499647	176437962	176444681 LOC499647
14	XM_342269.2	LOC361972	176445662	176485867 similar to RIKEN cDNA 9930117H01 gene
15	XM_579140.1	Dear	176736639	176739891 dual endothelin 1, angiotensin II receptor
16	XM_574968.1	LOC361973	177242730	177243851 LOC361973
17	XM_342270.2	Pet112l_predicted	177408100	177433793 PET112-like (yeast) (predicted)
18	XM_345227.2	LOC365834	177632475	177669341 similar to KIAA1759 protein
19	XM_227324.3	LOC295167	177769873	177774358 similar to S-ANTIGEN PROTEIN PRECURSOR
20	XM_215597.3	RGD1304885_predicted	177970585	178081504 similar to SH3 domain protein D19 (predicted)
21	NM_017153.1	Rps3a	178092351	178096720 ribosomal protein S3a
22	XM_342271.2	Lrba_predicted	178267690	178787634 LPS-responsive beige-like anchor (predicted)
23	XM_579311.1	RGD1308384_predicted	178793765	178923600 similar to RIKEN cDNA 6330415M09 (predicted)
24	NM_017079.1	Cd1d1	179011165	179014672 CD1d1 antigen
25	XM_574969.1	LOC499649	179088285	179089078 similar to high mobility group protein
26	NM_207606.1	Neph1	179114757	179169185 nephrin 1
27	XM_227485.3	Msr2_predicted	179278607	179293565 macrophage scavenger receptor 2 (predicted)
28	XM_227484.3	Cd5l_predicted	179363734	179425297 CD5 antigen-like (predicted)
29	XM_574970.1	LOC499650	179450738	179452834 LOC499650
30	XM_227483.3	Spap1_predicted	179483570	179576639 SH2 domain containing phosphatase anchor protein 1 (predicted)
31	XM_574971.1	LOC499651	179591714	179603714 similar to FLJ16478 protein
32	XM_227480.3	LOC295295	179625973	179625738 similar to phosphoglycerate mutase (EC 5.4.2.1) B chain - rat
33	NM_023982.1	Arhgef11	179738870	179796785 Rho guanine nucleotide exchange factor (GEF) 11
34	XM_227479.3	RGD1309453_predicted	179798792	179808808 similar to hypothetical protein FLJ32884 (predicted)
35	XM_215648.3	RGD1305653_predicted	179810529	179834374 similar to MEGF12 (predicted)
36	NM_021589.1	Ntrk1	179838740	179855545 neurotrophic tyrosine kinase, receptor, type 1
37	XM_574972.1	Insr	179857574	179875876 insulin receptor-related receptor
38	NM_207605.3	Sh2d2a	179916108	179924623 SH2 domain protein 2A
39	XM_227476.3	Prcc_predicted	179931946	179957221 papillary renal cell carcinoma (translocation-associated) (predicted)
40	NM_053707.2	Hdgp	179981346	179991892 hepatoma-derived growth factor
41	NM_001007637.1	mrpl24	179995713	180001745 mitochondrial ribosomal protein L24
42	XM_342272.2	RGD1311265_predicted	180001177	180009023 similar to CGI-41 protein (predicted)
43	NM_001007741.1	MGC94465	180009276	180018762 similar to hypothetical protein FLJ12671
44	NM_017244.1	Crabp2	180029801	180034147 cellular retinoic acid binding protein 2
45	NM_012987.1	Nes	180051907	180060947 nestin
46	XM_579358.1	LOC497759	180072888	180080749 hypothetical gene supported by NM_012916
47	NM_022285.1	Hapln2	180104684	180110115 hyaluronan and proteoglycan link protein 2
48	XM_215614.3	RGD1311086_predicted	180117561	180131693 similar to RIKEN cDNA 2610029K21 (predicted)
49	XM_215635.3	Apoa1bp_predicted	180132556	180144393 apolipoprotein A-I binding protein (predicted)
50	XM_227396.3	Iqgap3_predicted	180155674	180198814 IQ motif containing GTPase activating protein 3 (predicted)
51	NM_030860.2	Mef2d	180221078	180246917 myocyte enhancer factor 2D
52	NM_183054.1	Rhbg	180320787	180333050 Rh type B glycoprotein
53	XM_342276.2	RGD1304953_predicted	180371109	180380696 similar to SSTK-interacting protein (predicted)
54	NM_199091.1	Cct3	180381044	180434156 chaperonin containing TCP1, subunit 3 (gamma)
55	NM_001004226.1	0610031j06rik	180438070	180441660 kidney predominant protein NCU-G1
56	XM_227398.3	RGD1309886_predicted	180442734	180446076 similar to RIKEN cDNA 2310042N02 (predicted)
57	NM_013414.1	Bglap2	180482313	180483290 bone gamma-carboxylglutamate protein 2
58	XM_345231.2	RGD1307244_predicted	180514428	180526708 similar to CG5805-PA (predicted)
59	XM_578027.1	LOC502539	180537560	180545828 similar to IQ motif containing GTPase activating protein 3
60	XM_574973.1	Sema4a_predicted	180552405	180573625 sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4A (predicted)
61	NM_001002016.1	Lmna	180595724	180616354 lamin A
62	XM_227403.3	LOC310631	180645534	180653100 similar to KIAA2031 protein
63	XM_227404.3	Rab25_predicted	180657133	180662995 RAB25, member RAS oncogene family (predicted)
64	XM_215630.3	LOC295234	180665358	180668716 similar to late endosomal/lysosomal MP1 interacting protein

Area shown in yellow denotes region of overlap between the rat QTL and all human kidney disease located on human chromosome 1q21

Genes shown in bold denote that these gene were differentially expressed between the S and S.SHR(2) congenic (see supplemental Table 1)