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### Publication Date

2011

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UNIVERSITY OF CALIFORNIA, SAN DIEGO

**The Proteome of Mouse Brain Microvessel Membranes and Basal Lamina**

A thesis submitted in partial satisfaction of the requirements for the degree

Master of Science

in

Biology

by

Hyun Bae Chun

Committee in charge:

Professor Brian P. Eliceiri, Chair  
Professor Andrew D. Chisholm, Co-chair  
Professor Steven P. Briggs

2011



The Thesis of Hyun Bae Chun is approved and it is accepted in quality and form for publication on microfilm and electronically:

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University of California, San Diego

2011

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## LIST OF ABBREVIATIONS

ABC	ATP binding cassette family
Aqp4	Aquaporin-4
BBB	Blood-brain barrier
DAPI	4',6-diamidino-2-phenylindole
ECM	Extracellular matrix
FITC	Fluorescein isothiocyanate
MudPIT	Multi-dimensional Protein Identification Technology
NVU	Neurovascular Unit
Pgp	P-glycoprotein (ABCB1)
S0	Whole brain
S1	Crude microvessel preparation
S5	Purified microvessel preparation
SLC	Solute Carrier family of membrane transport proteins
Syn	Synaptophysin

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## ACKNOWLEDGEMENTS

I sincerely thank my thesis supervisor, Dr. Brian Eliceiri, for the interesting research project that I enjoyed working on. Also, thank you for the opportunity to conduct my thesis work at UCSD Medical Center – Hillcrest. It was an invigorating, inspiring, and humbling time of my college experience. This experience of being able to work in an excellent research environment with great staff and personnel will have a great impact on my future academic career.

My gratitude also goes to Dr. Bruce Torbett, Mike Scott, Dr. John Yates III, Dr. Sherry Niessen, and Heather Hoover for the collaboration in this project. I am also very thankful to Dr. Sergio Catz for his suggestions and support concerning cell lysis and the use of relaxation buffer with cell membranes and to Dr. Gabriel Simon for providing dtarray2.pl.

I would also like to thank the members of my defense committee Dr. Andrew Chisholm and Dr. Steven Briggs.

Certainly, this thesis would not have been possible without the support from the members of Eliceiri, Baird, and Coimbra lab, especially Dr. Andrew Baird, Dr. Raul Coimbra, Dr. Vishal Bansal, Dr. Jisook Lee for their guidance and moral support.

Special thanks goes to my father Sang Jin Chun for being the best dad.

**ABSTRACT OF THE THESIS**

**The Proteome of Mouse Brain Microvesel Membranes and Basal Lamina**

**By**

**Hyun Bae Chun**

**Master of Science in Biology**

**University of California, San Diego, 2011**

**Professor Brian P. Eliceiri, Chair**

**Professor Andrew D. Chisholm, Co-chair**

The blood-brain barrier (BBB) is a multi-cellular vascular structure in the central nervous system that separates the circulating blood from the brain parenchyma. It is composed of microvessels made of endothelial cells with tight intercellular junctions surrounded by a basal lamina, astrocytes, and pericytes. In

many diseased states of the brain, the integrity of the BBB is compromised and the changes in protein levels of key regulators such as transporters leave the brain open for attack. Previous studies have generated detailed databases of the microvessel transcriptome, however, less information is available on the BBB at the protein level. The focus of this study is not to elucidate on proteins that are already widely known and are hallmarks of the BBB but to generate a more extensive protein database of relatively unknown and unique proteins that can be used to gain further insight into the structure and function of the BBB. We specifically focused on the characterization of the membrane fraction of cells within the BBB to generate a more complete understanding of membrane transporters, tight junction proteins, and associated extracellular matrix proteins. Using Multidimensional Protein Identification Technology. 53% out of a total of 1,143 proteins found in mouse brain microvessels were determined to be membrane-associated. Analyses of specific classes of BBB-associated proteins in the context of recent transcriptome and proteome reports provides a unique database to assess the relative contribution of genes at the RNA and protein level in the maintenance of normal BBB integrity.

## INTRODUCTION

### 1.1 Blood-brain barrier – Physiological Function and Morphology

The blood brain barrier (BBB) is a multi-cellular structure that consists primarily of endothelial cells and astrocytes endfeet separated by a basal lamina (Abott et al., 2006). The functional barrier itself between the circulating blood and the brain parenchyma is created by apical junction complexes such as tight junctions between specialized endothelial cells in the brain microvasculature. The different cellular components (i.e. astrocytes, pericytes, and neurons) that cooperatively work to regulate the integrity of the BBB are termed neurovascular unit (NVU). The function of these specialized endothelial-associated cells, their secreted proteins, the tight inter-endothelial barrier itself, the unique transcellular transport, and active efflux pumps combine to yield a unique vascular barrier in the brain (Agarwal et al., 2009; Daneman et al., 2010; Enerson and Drewes, 2006).

### 1.2 Importance of BBB integrity with Focus on Structural Barriers

The brain microvasculature consisting of densely packed microvessels contain the structural basis of the BBB. Due to the high capillary density in the brain, single capillaries are approximately 40  $\mu\text{m}$  apart from each other, a distance short enough for small molecules to diffuse within a second (Rodriguez-Baeza et al., 2003). Generally, tight junction proteins are thought to act as the main physical and functional barrier of the BBB by sealing the space between endothelial cells to create a tight, non-fenestrated capillary that restricts free diffusional exchange of solutes (Nag, 2003), however studies with iron nanoparticles (Neuwelt et al., 1998) demonstrate that additional components of the neurovascular unit need to be considered. The iron nanoparticles used were superparamagnetic dextran-coated

monocrystalline iron oxide particles with sizes of the same order of magnitude as viral particles. Electron microscopy of brain sections from rats that have had their BBB permeability and also have iron nanoparticles intravenously injected, showed that the iron nanoparticles passed the capillary endothelial cells but did not cross beyond the basement membrane, which is evidence for a physiological barrier in the basal lamina between the astrocytes endfeet and the endothelial cells (Neuwelt et al., 1998). With this in mind, the NVU as a whole with its different cell components must be considered when exploring the BBB integrity.

### **1.2 Integrity of the Blood-brain Barrier and Importance in Disease States**

The primary role of the BBB in brain homeostasis is to regulate the transport of various substances between the blood and the brain such as drugs and metabolites using membrane transporters (Banks, 2009). Tight regulation of the BBB is integral in maintaining the metabolic homeostasis in the brain and in preventing toxins or pathogens from entering the brain parenchyma. In brain pathophysiology, the breakdown of the blood-brain barrier and alterations to its transport systems play a significant role in many CNS diseases including glioblastoma multiforme (GBM), traumatic brain injury, edema, multiple sclerosis, Alzheimer's disease, strokes, and HIV-1 encephalitis (Persidsky et al., 2006; Pottiez et al., 2009a). For example, the influence of gliomas on the blood-brain barrier leads to its breakdown and aggressive invasion of the surrounding brain parenchyma through mechanisms that still remains poorly understood (Lee et al., 2009).

### **1.3 Isolation of Mouse Brain Microvessels**

This study focuses on the identification of BBB protein components, especially membrane and extracellular components, from freshly isolated semi-intact microvessel. The microvessels are isolated from brain tissue using a density gradient centrifugation method and are purified using a glass bead column, all within a few hours to yield intact microvessels. These isolated and purified microvessels with a diameter of 1-10um maintain viability and are able to exhibit BBB function (Dallaire et al., 1991). As a form of ion affinity chromatography, the purification process works by extracellular matrix (ECM) glycoproteins, such as fibronectin and laminin (proteins found in our MudPIT results and are classically known ECM proteins surrounding brain capillaries), having strong interactions with the glass beads (Barnes et al., 1984). This fresh isolation procedure eliminates potential fixation, cell culture, and cell suspension artifacts by decreasing the relative time of isolation and purification, thus obtaining the most complete and pure resource of BBB-associated proteins to date.

### **1.4 Novelty of Our Proteomic Study**

Previous studies have examined gene and protein expression profiles of cultured brain endothelial cells which have yielded important expression profile data cataloging gene families using *in vitro* models (Agarwal and Shusta, 2009; Haseloff et al., 2006; Shusta et al., 2002). However, cell culture itself influences the relative abundance of various BBB cell subtypes, gene expression profiles and/or the ECM microenvironment. The integrity or the breakdown of the blood-brain barrier depends on the NVU as a whole. Many other techniques are currently used to isolate microvessels such as mechanical isolation, enzymatic isolation, and laser capture microdissection (Yousif et al., 2006). Microdissection strategies to recover

microvessels from brain tissue have yielded useful, comparative information on the protein expression profiles from an intact BBB. However, this approach requires fixation, laser dissection, immunostaining, and subsequent gel electrophoresis analysis. These conditions may also change the relative protein expression profiles like cell culture. Therefore, we focused on a fresh microvessel isolation to characterize the BBB proteome from mouse brains and then validated the isolation technique using immunoblotting to establish quality control. Based on our interest in developing a resource identifying ECM components and BBB membrane proteins that contribute to ECM interactions, we focused on a membrane-enrichment strategy for mass spectrometry-based protein profiling. Agrin, an ECM protein, and aquaporin-4, an astrocytic endfeet protein, are known to have an important impact on the homeostasis between the brain parenchyma and the circulating blood, but the mechanism cascade from the astrocytes membrane to the endothelial cells to maintain BBB integrity remains completely unknown (Wolburg et al., 2009a).

### **1.5 Proteomics – MudPIT**

Historically, BBB protein analyses depended on various gel electrophoretic-based techniques for protein separation, followed by antibody or mass spectrometry for protein identification. However, protein separation and identification is limited due to the limited insolubility of membrane/ECM proteins and gel electrophoretic resolution. The use of Multidimensional Protein Identification Technology (MudPIT) enables the discrete identification of a large number of proteins from complex protein mixtures obtained from freshly isolated tissues and does not require fixed/frozen samples or single cell suspension-based isolation techniques, i.e. flow cytometry or magnetic beads (Aebersold and Mann, 2003; Pottiez et al., 2009a; Wasburn et al.,



2001; Wu et al., 2003). Moreover, a recent study has used mass spectrometry with the addition of isotope labeled peptide standards to provide quantitation for known proteins (Uchida et al., 2011).

## **1.6 Objectives and Summary**

In our study, freshly isolated normal mouse brain microvessels were isolated, its membranes fractionated, underwent proteolysis, and were followed by protein profiling with MudPIT. We have performed multiple MudPIT runs to maximize coverage of membrane proteins of both high and low abundance and to establish standards for technical replicates within the same samples and between different tissue collections. Identified proteins were assigned to various BBB classes based on classical markers for specific cell types and as well as recent proteomic and transcriptome studies (Daneman et al., 2010; Enerson and Drewes, 2006). Our strategy enabled the identification and classification of BBB membrane transporters, tight junction proteins, integrins, ECM proteins and proteoglycans. We discuss the classes of proteins identified and their relative abundance for the purpose of providing a resource for the membrane protein constituents of the normal BBB.

## **MATERIALS AND METHODS**

### **2.1 Mouse Brain Microvessel Isolation**

Mouse brain microvessels were isolated using a modified protocol based on previously published studies (Hartz et al., 2006; Yousif et al., 2007) based on the IACUC approval of the University of California San Diego. For each preparation, mice (n=6) were over-dosed with 5% isoflourane and the brains were harvested. All subsequent steps were carried out at 4°C or on ice. After a wash in PBS, the brains were homogenized using a dounce (0.25 mm clearance) (whole brain is defined as the S0 fraction, a nomenclature of S0, S1 and S5 to describe the microvessel fractionation steps consistent with the microvessel isolation procedure as described below and previously by Yousif et al., (Yousif et al., 2007)). 15 mL of 30% Ficoll was added to 10 mL of the homogenate and mixed thoroughly. The resulting density gradient was spun at 5800xg for 20 min. (Pellet was defined as the S1 fraction). The pellet was resuspended in 1mL of PBS and passed using PBS with 1% BSA through a glass bead column, with 100 µm nylon mesh filter on the top and a 20 µm nylon mesh filter on the bottom. The glass beads were gently agitated in PBS with 1% BSA to obtain the microvessels. The resulting sample was washed with BSA-free PBS before use in experiments, and defined as the S5 fraction.

### **2.2 Biochemical Analyses in Isolated Endothelial Cells**

Mouse brain microvessels were isolated as described previously. Mice were injected with 1 mg of FITC-lectin (FL 1101-5; Vector Labs) 20 minutes prior to sacrifice to validate the integrity of isolated vessels. The following antibodies were

used to validate microvessel confirmation; anti-Synaptophysin (Abcam; 1:1,000), anti-AQ4 (Millipore; 1:500), anti-P-glycoprotein (Abcam; 1:250).

### **2.3 Endothelial Cell Lysis Using Nitrogen Cavitation**

At 4°C, approximately 40µL of tissue/cells were suspended 0.5mL of relaxation buffer (pH 7.25, 50 mM KCl, 1.5 mM NaCl, 5 mM PIPES, 1.75 mM MgCl<sub>2</sub>, 1 mM ATP, 0.62 mM EGTA). The cell suspension was placed in a 15 mL conical tube, cavitation bomb tubing attached, and nitrogen pressure (500-700 psi) applied for 10 minutes. After placing a clean 15 mL receiving tube at the end of the pressure relief tubing pressure relief valve was gently opened to collect lysed cells. The tubing was then purged to an additional 0.5 mL of relaxation buffer and pooled with the lysate.

### **2.4 Endothelial Cell Lysate Membrane Fractionation and Protein Digest**

At 4°C, the lysate was placed in a microfuge tube and spun at 750xg for 5 minutes. The supernatant was placed in a 5 mL ultracentrifuge tube with relaxation buffer to a total volume of 4.8 mL, and centrifuged at 100,000 xg for 90 min. The supernatant, containing the cytosolic proteins, was discarded, while the pellet, containing the membrane proteins, was resuspended in 15 µL 8M urea followed by 20 µL 0.2% ProteaseMAX acid-labile surfactant (Promega, Madison, WI) in 50 mM ammonium bicarbonate. Aliquots of resuspended membrane proteins were analyzed for total protein content by Bradford protein assay (BioRad Corporation, Hercules, CA). Proteins were then reduced with dithiothreitol and carbamidomethylated with iodoacetamide prior to tryptic digestion according to the ProteaseMAX technical bulletin (ProteaseMAX surfactant, Trypsin Enhancer; #TB373; Promega Corporation,

Madison WI). Digestion was stopped by addition of 90% formic acid to obtain a final in-digest concentration of 5% formic acid.

## **2.5 Multidimensional Liquid Chromatography – Mass Spectrometry (MudPIT)**

Samples in 1.5 mL microfuge tubes were diluted 1:2 with 5% acetonitrile 0.1% formic acid (Buffer A), centrifuged at 13,000 xg for 10 minutes, and supernatant proportionate to 15 µg of protein was transferred to a new microfuge tube for loading to a MudPIT precolumn. The MudPIT precolumn (250 µm internal diameter) with 4 cm of strong cation exchange 5 µm resin (Partisphere SCX, GE Healthcare) was preceded by 3 cm of C18 stationary phase column (5 µm particle size, Jupiter C18, Phenomenex Corporation). The resolving column dimensions were 100 µm (internal diameter) with 10 cm of C18 stationary phase. Operating parameters of the MudPIT experiment are provided in the Supplementary Data Table 2, based on the standards provided by the Minimum Information About a Proteomics Experiment (MIAPE, <http://psidev.sourceforge.net/miape/>)

## **2.6 Bioinformatic Analyses**

Spectra were searched using SEQUEST, and filtered using DTASelect2 [[http://www.scripps.edu/chemphys/cravatt/protomap/dtaselect\\_instructions.html](http://www.scripps.edu/chemphys/cravatt/protomap/dtaselect_instructions.html)]. Spectral count data were then processed using PatternLab for Proteomics (Carvalho et al., 2008). The spectral count information stored in the DTASelect-filter.txt files was extracted by using the Perl script dtarray2.pl (Kaschani et al., 2009). The resulting dtarray2.txt file was loaded into Excel (Microsoft Corp.) and further ordered. This arrangement is the source for the master list of protein identifications in Supplementary Table 1. Information for IPI is located at:

<http://www.ebi.ac.uk/IPI/IPIhelp.html>. Information on Uniprot Knowledgebase is located at <http://www.uniprot.org/help/uniprotkb>. For SEQUEST searches followed by DTASelect2 processing, the following criteria were utilized: (a) a 1% false positive discovery rate; (b) a minimum of two unique tryptic peptides per identification was required; (c) oxidation of methionine (+16Da) was allowed as a differential modification; and (d) cysteines were required to be carbamidomethylated (+57 Da).

Statistical validation was performed for every protein hit using statistical functions and parameters set within the SEQUEST search as we have previously described in detail (Tabb et al.,, 2002). Additional information on the use of DTASelect2 is available at [http://www.scripps.edu/chemphys/cravatt/protomap/dtaselect\\_instructions.html](http://www.scripps.edu/chemphys/cravatt/protomap/dtaselect_instructions.html), along with additional information on data analysis of shotgun proteomics (Yates et al.,, 2009).

## RESULTS

### 3.1 Microvessel Isolation and Validation

The cortices of normal mouse brains were dissected and subjected to a multistep isolation protocol (includes the key steps: homogenization, density centrifugation, and filtration through glass bead column; see overview in Figure 1A) that has been previously described and utilized for transcriptome analysis, BBB transporter, and cell signaling studies (Enerson and Drewes, 2006; Hartz et al., 2006; Yousif et al., 2007). To verify the enrichment of brain microvessel material after density centrifugation in comparison to a step before (after homogenization), we first examined the integrity of the microvessel lumens by staining the vasculature prior to harvest. Mice were subjected to an intravenous injection of fluorescently-labeled endothelial cell-binding lectin from the plant *Griffonia simplicifolia* (Figure 1B). After harvest of the labeled brain and harvest of the microvessels from the Ficoll gradient (termed S1 as described by Yousif et al., and see Materials and Methods for further details), samples were subjected to imaging by laser scanning confocal microscopy to reveal stained microvessels based on the fluorescent lectin binding to the lumen of the endothelia. Further enrichment of microvessel preparations were achieved by filtration through a glass bead column (S5; Yousif et al., 2007), leading to a significant reduction of cellular debris. This reduction was verified by imaging the different sample preparations (S1 vs. S5) by phase microscopy. The staining of the samples with DAPI validates the nuclear integrity and that the endothelial cells in the microvessels are intact (Figure 1C). Consistent with previous studies (Yousif et al., 2007), microvessels were most enriched in the S5 preparation (i.e. after glass bead purification) vs. the S1 preparation (i.e. prior to glass bead purification), as evidenced

by the loss of cellular debris, such as neurons and unnecessary neuroglia. To further confirm the validation of enrichment by biochemical analyses, lysates of whole brain (i.e. S0) were compared to purified microvessels (i.e. S5) by immunoblotting. Antibodies to the endothelial (i.e. P-glycoprotein, Pgp/Abcb1), neuronal (i.e. Synaptophysin, Syn), and astrocyte endfoot (i.e. Aquaporin-4, Aqp4) compartments of the BBB were used. While both Pgp and Aqp4 were increased as predicted in the isolated microvessels (Figure 1D), levels of the neuronal marker (Syn) were decreased. This immunoblot result indicates a relative enrichment of the key components of the BBB, endothelial cells and vessel associated astrocyte endfeet, in the S5 fraction vs. the S0 fraction. These quality control techniques (i.e. lectin staining, microscopy, and immunoblotting) were performed on parallel samples or aliquots of the preparations used for proteomic analysis, to ensure the most rapid processing of fresh unstained microvessels for membrane fractionation and subsequent proteomic analysis.

## **2.2 Proteomic Profile of Mouse Brain Microvessel Membranes**

After the microvessels have been isolated, purified, and enriched (S5), they were subjected to nitrogen cavitation lysis, membrane isolation, tryptic digestion, and prepared for mass spectrometry analysis as described in the Materials and Methods. The gentle lysis by nitrogen cavitation ensured that the proteins stayed intact as cytosolic components were separated from the membrane. The membrane proteins were then subject to tryptic digestion with ProteaseMAX (Promega) for maximal solubilization and peptide recovery. ProteaseMAX, which is a hydrophobic anionic sulfonate surfactant, enhances the solubilization and peptide recovery of hard to digest membrane proteins by allowing the proteins to become more accessible to the

protease trypsin. The surfactant is also classified as a cleavable detergent because acidic conditions would cleave it into products that will not interfere with mass spectrometry results downstream. The resulting peptides were then separated by 2D liquid chromatography composed of a strong cation exchange and reverse phase high performance liquid chromatography using microcapillary columns followed in-line by tandem mass spectrometry as previously described (i.e. MudPIT (Chen et al., 2006)). Utilizing three of the distinct S5 microvessel membrane fractionation samples, independent mass spectrometry runs enabled the identification of a total of 1,645 microvessel proteins, i.e. BBB-associated proteins. The standards and settings used for data acquisition (Taylor et al., 2007) (i.e. MIAPE data) for these studies are found in Supplementary Table 1 along with a complete list of the proteins and their spectral counts from all mass spectrometry runs.

Mass spectrometry is a sampling technique by which probability of detection is a function of protein abundance. By counting the numbers of spectra that map to identified proteins, spectral counting can be used as a surrogate for relative protein abundance (Liu et al., 2004). The criteria used for protein inclusion for the study was that proteins must be identified by MudPIT in 3 technical replicates (i.e. the same sample analyzed at 3 different times; 153-1, 153-2, and 153-3), all of which are considered as a single biological sample, then in an additional 2 independent biological samples (118 and 153). Moreover, the average of the spectral counts from the 3 technical replicates and the 2 biological samples must be  $\geq 5$  (Figure 2 and Supplementary Table 1). However, this should not undervalue the biological relevance of the less abundantly identified proteins (mean spectral counts  $< 5$ ) from our study. Mean spectral counts is just an objective standard to be sure that the proteins are actually present and sufficiently represented. In some cases, proteins



were found in the technical replicates, but not in the biological replicates, as well as the converse (Supplementary Table 1). Each protein sample, 118, 153, and 155, was extracted from isolated, pooled microvessels obtained from different mice.

The first determine the intra-sample variation of proteins identified by MudPIT, three samples from the same S5 microvessel fraction preparation were analyzed (i.e. technical replicates), then the distribution of proteins identified were compared (Figure 2A). For the technical replicates, we found 1,355 proteins (74% of total proteins) common to all three samples, with 247 proteins unique to replicate 153-1, 160 proteins to sample 153-2, and 121 proteins to sample 153-3 (Figure 2A). The distribution of proteins common to any 2 samples was found to be 102 proteins between 153.1 and 153.2, 93 proteins between 153.2 and 153.3, and 164 proteins between 153.1 and 153.3. Thus, a significant degree of protein overlap, 74%, exists between each replicate from the same starting microvessel preparation.

Next, we compared the distribution of proteins identified between the three different S5 samples, where each sample was from pooled tissue of six independent mice (Figure 2B). 1,143 proteins (69% of total proteins) were common to all three samples, whereas 73 proteins were unique to sample 118, 247 proteins to sample 153, and 150 proteins to sample 155. The distribution of proteins common to any two samples was found to be 67 proteins between 118 and 153, 249 proteins between 153 and 155, and 58 proteins between 155 and 118. As assessed with the technical replicates of 153, a significant portion of the proteins identified (69%) was common to all three independent samples and was only slightly less than that identified (74%) from the technical replicates. Moreover, a scatter plot comparison of the proteins identified following the parallel analyses of replicate samples established the degree of overlap in an analysis of intra-sample variation (Figure 2C).

Lastly, Figure 2D, a summary of the bioinformatic analysis of the BBB proteins identified from the 3 samples obtained from different groups of mice (Figure 2B), provided insight as to the number of membrane proteins recovered (53%), as well as mitochondrial (15%), and cell junction proteins (5%) recovered. A complete listing of all proteins identified is found in Supplementary Table 1 and the proteins are grouped in Tables 1-4 based on previously published reports and a classical understanding of their biological function.

### 3.3 Membrane Transporter and Channel Proteins

The BBB is characterized by the expression of a complex range of solute carrier proteins (SLC), ATP-binding cassette proteins (ABC), and voltage gated channel proteins that are essential for normal function of the brain, i.e. import of nutrients and drugs and export of metabolites and toxins across the endothelium. SLC/ABC transporters were ranked based on their number of spectral counts, which can be used for relative quantification between identified proteins. We identified 84 SLC/ABC transmembrane proteins by MudPIT, with proteins having a spectral count of <5 peptides indicated in italics below the dashed line. All of the detailed supporting spectral count data are provided in Supplementary Table 1. Of these microvessel-derived membrane proteins identified, we noted that 21 of these proteins (i.e. Slc25a4, Abcb1a, Slc12a5, Slc3a2, Slc2a1, Slc1a6, Slc12a2, Slc6a1, Slc38a3, Abcc4, Slc9a3r2, Slc16a1, Slc2a3, Slc9a3r1, Abcd3, Slc17a7, Slc30a1, Slc27a1, Asna1, Abcg2, and Slc7a1) were identified in an earlier study based on mRNA expression profiling in rat brain microvessels (Enerson and Drewes, 2006). An additional 13 SLC/ABC proteins (i.e. Slc9a1, Abca7, Abcb8, Abcc9, Slc25a10, Slc5a6, Slc25a20, Slc4a3, Abcb6, Abca2, Slc6a20, Abcc8, and Slc38a2) were detected in mouse

microvessels with a spectral count of <5 that were also present at the mRNA level in rat microvessels (Enerson and Drewes, 2006). Moreover, 5 out of 6 ABC proteins recently quantified in a study of human BBB transporters (Uchida et al., 2011) were also identified in our study, some with spectral counts  $\geq 5$  (i.e. Abcb1a, Abcc4, and Abcg2) and others <5 spectral counts (i.e. Abca2 and Abcc8). In a similar comparison, 7 out of 10 SLC proteins had spectral counts  $\geq 5$  (i.e. Slc1a3, Slc3a2, Slc2a1, Slc2a3, Slc16a1, and Slc7a1), while Slc7a5 had a spectral count <5 (See Table 1 and Supplementary Table 1). An additional 36 transmembrane proteins identified by MudPIT in our study were also detected at the mRNA level from isolated mouse endothelial cells (i.e. Abcb1a, Slc3a2, Slc2a1, Slc38a3, Abcc4, Slc9a3r2, Slc16a1, Slc6a17, Slc25a24, Slc25a25, Slc30a1, Slc1a1, Abcg2, Slc7a1, Slc44a1, Slc25a20, Slc39a10, Slc44a2, Slc12a7, Slc22a8, Slc16a2, Slc7a5, Atp7a, Slc38a2, and Slc7a3; See Table 1, single asterisk), pericytes, (i.e. Atp1a2, Atp1b2, Slc12a2, Atp2a3, Atp13a5, Abcc9, and Abca9; See Table 1, double asterisk), and astrocytes (i.e. Atp1a2, Slc1a2, Slc4a4, Slc25a18, Slc39a12; See Table 1, triple asterisk) (Cahoy et al., 2008; Daneman et al., 2010). Prototypic examples of BBB proteins in the endothelium, such as Abcb1a (i.e. Pgp), insulin receptor (INSR), the transferrin receptor (Tfrc) and  $\alpha_2$  macroglobulin receptor (Lrp1), were detected among the top 50 proteins in our MudPIT studies as well as from mRNA profiling of microvessels (Enerson and Drewes, 2006), isolated endothelium (Daneman et al., 2010), and human microvessel studies (Uchida et al., 2011). Table 1 provides a listing of the 101 transmembrane SLC/ABC and voltage gated channel proteins identified by MudPIT with relevant references to other complementary profiling studies. Additional SLC/ABC and voltage gated channel proteins are listed in Supplementary Data Table

1. About 15% of the genes at the blood-brain barrier encode for transporters and only about 50% of the transporters are known (Pardridge, 2003).

### 3.4 Tight Junction Proteins and Integrins

The physical integrity of the endothelial component of the BBB is generally defined by the expression of tight junction proteins, therefore, we examined their relative abundance following MudPIT of mouse brain microvessels (Table 2). In our analyses, 25 tight junction proteins were identified with a spectral count  $\geq 5$  (Table 2A), including Pecam1 (i.e. CD31), occludin, five catenins (Types  $\alpha 1$ ,  $\alpha 2$ ,  $\beta 1$ ,  $\delta 1$ , and  $\delta 2$ ) and claudin (Type 11). An additional 8 tight junction proteins were identified including Vcam1, Jam 2, and claudin 5, although having limited representation among recovered sequences (i.e.  $< 5$  spectral counts). Cytoplasmic tight junction adapter proteins were also found (9 protein hits with  $\geq 5$  spectral counts; Table 2B) and included cadherins (Types 2 and 13), Esam1, F11r, and Tjp1 and Tjp2 as well as 15 proteins with low representation (Table 2B). In addition to the tight junction proteins, 7 integrins were identified ( $\alpha 1$ ,  $\alpha 2b$ ,  $\alpha 3$ ,  $\alpha 6$ ,  $\alpha 7$ ,  $\alpha v$ , and  $\beta 1$ ), with an additional 7 integrins detected with low recovery ( $\alpha 2$ ,  $\alpha 5$ ,  $\alpha m$ ,  $\beta 2$ ,  $\beta 3$ ,  $\beta 5$ , and  $\beta 8$ ; Supplementary Table 1). These findings provide a resource for the further analysis of cell-cell and cell-ECM mediators expressed in the membrane fraction of brain microvessels.

### 3.5 ECM and Basal Lamina Proteins

To better characterize the ECM microenvironment and basal lamina proteins, we examined the MudPIT proteomic profile from microvessels. We identified 24 ECM proteins including tenascin-R, fibrinogen  $\beta$  chain, 6 laminins (types  $\alpha 1$ ,  $\alpha 2$ ,  $\alpha 4$ ,  $\beta 1$ ,  $\beta 2$ , and  $\gamma 1$ ), and 5 proteoglycans (agrin, brevican, dystroglycan, perlecan, and versican).

In addition, the neuronal associated proteins contactin (Types 1 and 2), and neural cell adhesion molecule L1 were detected. While only collagen IV  $\alpha$ 1 and  $\alpha$ 2, were detected with spectral counts  $\geq 5$ , several additional collagens (types I $\alpha$ 2, IV $\alpha$ 3, VI $\alpha$ 1, XII $\alpha$ 1, XV $\alpha$ 1, and XVIII $\alpha$ 1) were also detected with low spectral abundance. In combination our findings provide a more complete resource of the relevant ECM proteins present in brain microvessel microenvironment than has been previously reported.

### 3.6 BBB-specific Cell Types

The analysis of microvessel proteome included the identification of proteins associated with endothelial cells and their associated astrocytes and pericytes. A total of 24 proteins were associated with endothelial cells based on previously published studies of the RNA expression in isolated cells (Daneman et al., 2010). Of these the ABC transporter Abcb1a was the most abundant, along with several SLC proteins (Slc2a1, Slc1a1; see See Table 1), von Willebrand factor-related proteins (Vwf and Vwa1, See Table 4A), as well as CD31 and Occludin (Pecam1 and Ocln, See Table 2A). A total of 31 endothelial proteins were identified (Table 4A), in addition to the 36 transmembrane endothelial transporter/channel proteins annotated in Table 1. VEGF receptors (Flt1 and Kdr), were also detected, although below the 5 spectral count threshold. Also present were the prototypic astrocyte markers such as GFAP and Aqp4, as well as the more recently characterized astrocyte marker, Aldh1L1. Similarly, recovery of pericyte markers such as Slc1a2, Slc4a4, and laminin  $\alpha$ 2 (Table 3) indicates that the membrane fractionation of microvessels recovers both transmembrane proteins as well as associated ECM proteins.

## DISCUSSION

In this study we validated a combination of brain microvessel isolation, membrane fractionation, proteolytic digestion, liquid chromatography, mass spectrometry and bioinformatic techniques to identify and catalog the protein expression profile in the BBB. The specific strategy enabled the profiling of transmembrane and membrane-associated markers of the BBB at the protein level (Kamiie et al., 2008; Ohtsuki and Terasaki, 2007). We focused on the identification of the canonical classes of membrane-associated BBB proteins that are associated with solute transport functions and normal brain physiology (e.g. SLC/ABC transporters and voltage gated channel proteins), tight junction proteins maintaining integrity (i.e. the transmembrane proteins and associated cytoplasmic adapters) and basal lamina extracellular proteins that constitute the ECM microenvironment and indirect intercellular contacts (i.e. integrin mediated interactions with the basal lamina) (Hawkins and Davis 2005). For the ECM proteins in particular, we were interested in an approach to identify basal lamina proteins from intact microvessels to provide the most physiologically relevant *in vivo* profile, in contrast to *in vitro* assays where cell culture has the potential to introduce artifacts with regard to ECM protein expression.

The intent of our studies was to generate a BBB protein database utilizing MudPIT, which provides a reference for identification of BBB membrane proteins, ECM, and basal lamina. Membrane fractionation of the microvessels followed by a urea-assisted, acid-labile, surfactant-mediated protease digestion was used to identify: 1) the most abundant proteins (i.e. a standard of comparison for proteomic analysis and a measure of contaminants from other cellular compartments); 2) the profile of classical transmembrane transport proteins; and 3) the profile of tight

junction and ECM proteins associated with microvessels (i.e. a measure of proteins that define the physical barrier properties of the NVU (Wolburg et al., 2009b)).

Previous studies by Enerson and Drewes examined the mRNA expression of isolated rat microvessels and validated the techniques to isolate intact fresh microvessels (Enerson and Drewes, 2006). We utilized a similar isolation strategy as described by several laboratories to isolate BBB components, then interrogated which proteins were associated with the BBB membrane fraction utilizing MudPIT. We annotated our proteomic findings for comparison with mRNA expression profiling of SLC/ABC transporters (Table 1), tight junction proteins (Table 2), and endothelial genes (Table 4A) leading to the observation that as expected, many of the genes expressed at the mRNA level in rat brain were also expressed in membrane protein-focused proteomic profiling. For example, 62% of the SLC/ABC transporters expressed at the RNA level by Enerson and Drewes were also detected in our proteomic study. Although a total of 217 SLC/ABC transporters and voltage gated channel proteins were detected by our MudPIT approach, only 58% of these proteins satisfied our selection threshold for identity in 3 different biological samples and with the average spectral count  $\geq 5$ . Although additional membrane transporters were identified, further investigation will be necessary to validate the expression and biological relevance of the less abundant proteins that did not meet the stringent spectral count cutoff criteria. We have compiled a complete listing of all peptides identified with accompanying spectral data (Supplementary Table 1). Similarly, 53% of the tight junction proteins identified by Enerson and Drewes were also detected by MudPIT, while an additional 21 tight junction proteins/adapters (Daneman et al., 2010) were identified along with 7 integrins that have not been previously identified through microvessel proteomics. Tables 1-4 are further annotated with gene expression data

based on protein (Agarwal et al., 2009; Cayrol et al., 2008; Cunnea et al., 2010; Krishna and Redies, 2009; Li et al., 2009; Lu et al., 2008b; Ohtsuki and Terasaki, 2007; Pottiez et al., 2009b; Yousif et al., 2007) and mRNA (Cahoy et al., 2008; Daneman et al., 2010; Enerson and Drewes, 2006) detection strategies. A recent quantitative targeting proteomics strategy, reported as sensitive to the fmol range, for human blood-brain transporters and receptors yielded a similar ABC/SLC transporter profile, with 5 of 6 ABC proteins and 7 of 10 SLC proteins identified by MudPIT present in the reported human study (Uchida et al., 2011). Interestingly, Uchida et al. compared expression levels of ABC transporters between human and mouse microvessel and determined that *Abca8*, which was not detected in our study, and *Abca2*, which had a spectral count <5, were present in their analyses at relatively low concentrations in mouse microvessels. Thus, it seems that our findings compare favorably with findings by Uchida et al. (Uchida et al., 2011) consistent with a high degree of protein detection sensitivity in our studies.

The recent mRNA gene profiling report (Daneman et al., 2010) on isolated endothelial cells, pericytes and astrocytes provided an opportunity to compare and annotate our protein-based findings. We observed that by focusing on the membrane fraction for proteomic analyses we were able to obtain an unexpectedly large number of SLC/ABC transporters present in microvessels, of which the mRNA expression of specific family members of transporters was similar, but not identical to that observed in isolated cells (Daneman et al., 2010). For example, 5 of the 6 transporters *Apccd1*, *Abcb1a*, *Slco1a4*, *Slc22a8*, *Slc7a5*, but not *Slc01c1* were highly expressed at the mRNA level by Daneman et al., (Daneman et al., 2010) and were also identified by proteomics, although with low spectral counts. Similarly, while we obtained a large number of transporters as identified by RNA-based expression studies, the exact



expression profile of transporters varied. We recognized the inherent difficulty of protein isolation from complex tissue sources and identification of low abundance proteins, as compared to RNA transcript analysis of low abundance mRNAs. However, the identification of proteins with low spectral counts provides a starting point for further studies. Most importantly, the further understanding of the interdependent roles of the ~120 transporters recovered with the most abundant spectral counts will provide insight into BBB integrity and function.

Interestingly, among the most abundant transmembrane transporter/channel proteins was *Atp1a2*, whose gene expression was detected in both isolated pericytes (Daneman et al., 2010) and astrocytes (Cahoy et al., 2008). The expression of other pericyte markers such as *PDGFR $\beta$*  was low, or in the case of *CD13* and *NG2*, not represented in the proteomic database generated herein. Expression profiling is another complementary technique, which has been used by Shusta and colleagues to characterize membrane BBB proteins using expression cloning (Agarwal et al., 2009; Agarwal and Shusta 2009). Other *in vitro* proteomic screening studies have focused on triton-soluble proteins (Pottiez et al., 2009b) and proteins lipid membrane rafts in cultured human endothelial cells where *Alcam*, *AHNAK* (desmoyokin), and intracellular trafficking proteins (clathrin light chain and *Cav-1*) were identified (Cayrol et al., 2008). *In vivo* proteomic analysis of integral membrane proteins with colloidal silica recovery technique also identified *Cav-1*, *GPR116*, *G  $\alpha$ q*, *G  $\alpha$ 12*, *Flot-2*, *Fyn*, *ADAM10*, and *contactin-1* (Li et al., 2009). Several of the proteins identified with these complementary approaches have also been identified in our study and are annotated in Table 4A. In combination, these findings underscore that our MudPIT-based approach for microvessel protein identification is useful for high coverage of

membrane proteins and identification of physiologically relevant proteins on specific BBB cell types (Pottiez et al., 2009a).

While transcriptome profiling provides a standard from which to assess mRNA expression levels in microvessels or isolated primary cells, the generation and characterization of a BBB gene expression profile at the protein level is essential for comparative purposes. Therefore, developing techniques to better analyze unknown constituents of the BBB proteome is essential as shown by the laser capture microdissection (LCM) of microvessels from mouse brain slices using cryopreservation, fixation, and immunostaining followed by gel electrophoresis and then mass spectrometry for protein identification (Lu et al., 2008b). In a comparison with our MudPIT strategy of utilizing fresh intact microvessels, 58% of the BBB-enriched/endothelial proteins identified by LCM/gel electrophoresis were also evident in our studies as indicated in Tables 1-4. Quantification of the relative abundance of known membrane transporters from human and mouse microvessels (Uchida et al., 2011), along with our results identifying proteins demonstrates that a MudPIT approach is highly appropriate for proteomic identification from isolated BBB membranes and provides increased protein identification coverage. Each spectral count is a selected peptide precursor ion selected for fragmentation that is subsequently assigned for protein identification. For spectral counts between 5 and 500, abundance is linearly proportionate to spectral count (Collier et al., 2010; Cooper et al., 2010; Liu et al., 2004). Therefore, while there are limitations with relying exclusively on spectral counts for relative quantitation, once protein identifications have been made using a procedure such as the MudPIT, other techniques can be deployed to quantify known proteins (Uchida et al., 2011). The identification of ECM proteins and transmembrane integrins in our study using freshly isolated tissue

provides a profile of the membranes in the contact with the extracellular microenvironment is not influenced by potential tissue culture induced artifacts (McCarty 2009). While many ECM proteins were identified by MudPIT (i.e. six different specific laminin subtypes and five different proteoglycans), the mRNA encoding these and other ECM proteins appeared to be under-represented in transcriptome studies, suggesting the possibility that ECM proteins with a long half-life may help explain the low abundance of their coding mRNAs. Therefore, to better understand the ECM microenvironment in the BBB, a MudPIT strategy provided the most complete protein expression profile to date of ECM proteins.

In contrast to classical 2D isoelectric focusing/SDS-PAGE, where solubilization and resolution of hydrophobic proteins is limited, our solubilization and enzymatic fragmentation yielded experimental material sufficient for a relatively high content analyses of a large number of peptide fragments. This was due, in part, to the use of: 1) a cell membrane lysis approach that minimized nuclear lysis; and 2) a relaxation buffer to better unfold membranes (Borregaard et al., 1983) prior to lysis and centrifugal fractionation; and 3) use of urea as a solubilizing agent and ProteaseMAX acid-labile surfactant to increase the efficiency of tryptic digestion. Similar approaches applied to the analysis of membrane proteins in yeast, heart, and human parotid exosomes have led to the identification of high content membrane protein data sets (Gonzalez-Begne et al., 2009; Lu et al., 2008a; Lu et al., 2009; Washburn et al., 2001). As with most sensitive analytical tools such as MudPIT, an important consideration is the purity of the starting material (i.e. microvessels vs. other cell types) and the quality of the membrane fractionation (i.e. the yield of membrane proteins). Therefore, we have relied on validation techniques consistent with DeClevés and colleagues (Yousif et al., 2007) and our own immunoblotting of

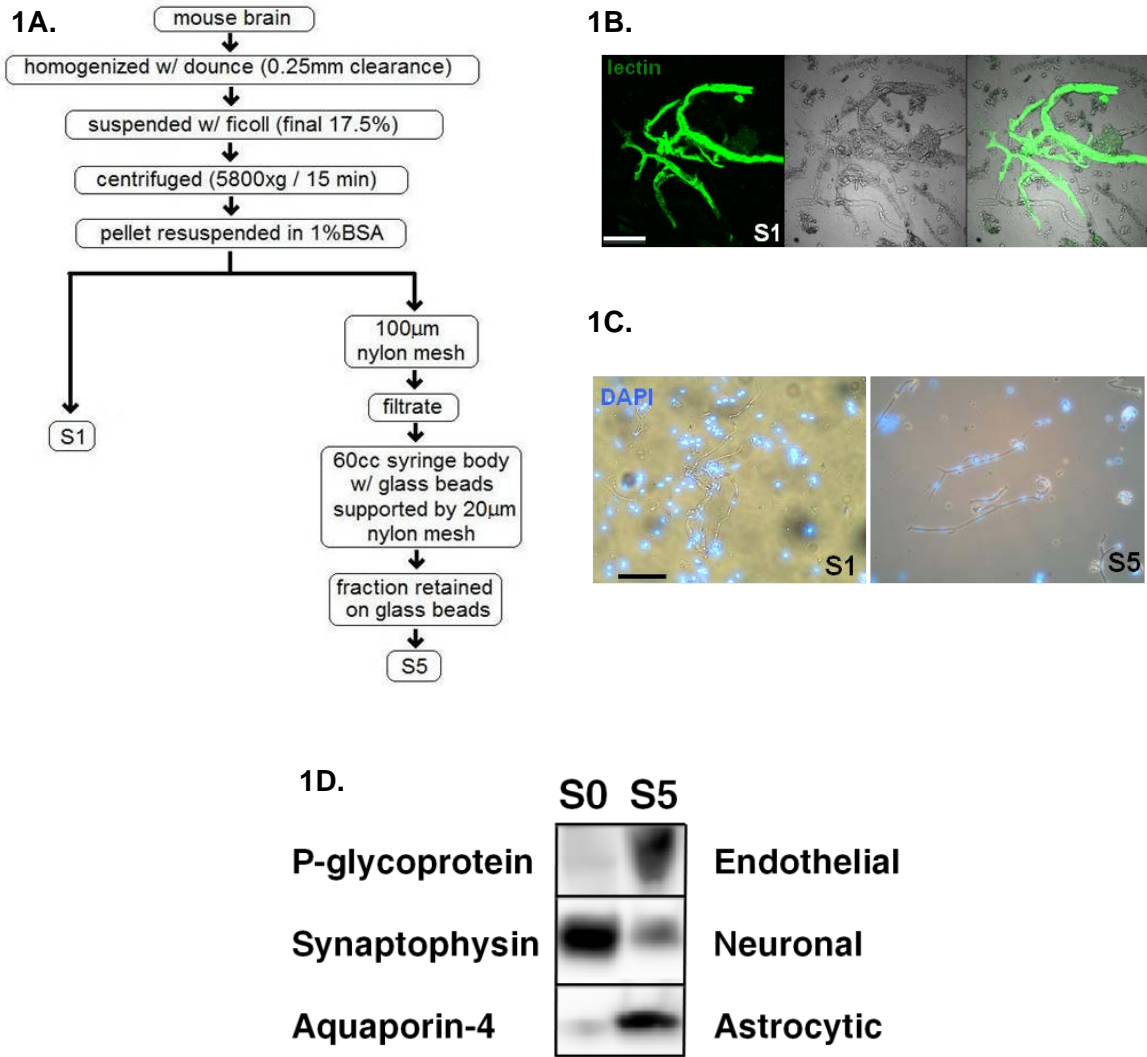
starting microvessel material to validate the expression of membrane proteins, enrichment of endothelium/astrocyte endfeet and a decrease in neuronal proteins.

In a comparison of the proteins identified in this study of mouse brain blood vessels by MudPIT vs. RNA gene profiling and immunohistochemistry, it remains a challenge to determine how source, isolation, and analytical techniques may influence the conclusions (Uchida et al., 2007). For example, in the study of the BBB, expression of claudin 5 and cadherin 11 are widely described (Hawkins and Davis, 2005; Nitta et al., 2003), and yet neither was a major component of the mass spectrometry spectral counts analyzed. In contrast, cadherins 1 and 2 and cadherin 11 predominated in our analysis suggesting that if extraction techniques affect different members of the same family, then such differences in apparent abundance may have other unknown explanations. These differences may reflect the limitations recovery techniques, the source of the lysate (i.e. heterogeneity of brain microvessels), the post-translational status, ex vivo proteolytic activity, and the possibility that other tight junction proteins may deserve closer study in terms of the relative abundance and physiological relevance. For example, a recent *in situ* mRNA study of blood vessel cadherins in developing ferret brains indicated that while many cadherins were expressed in development, surprisingly no mRNA was detected in adult brain capillaries solubilization. This finding further supports the need to characterize BBB components at the protein level (Krishna and Redies, 2009).

It has become increasingly clear that a better understanding of the BBB requires a more comprehensive knowledge of the protein constituents of this important biological barrier (Czeisler and Janigro, 2006; Neuwelt et al., 2008; Pardridge, 2010; Soni et al., 2010). In our study, we have utilized MudPIT on fresh unfixed microvessels to provide a comprehensive database resource for membrane

transporter, tight junction and ECM proteins of the BBB. Our microvessel-based analysis of proteome provides a novel bioinformatic resource to help place into perspective transcriptome and proteomic findings from intact microvessels and isolated cells of the BBB.

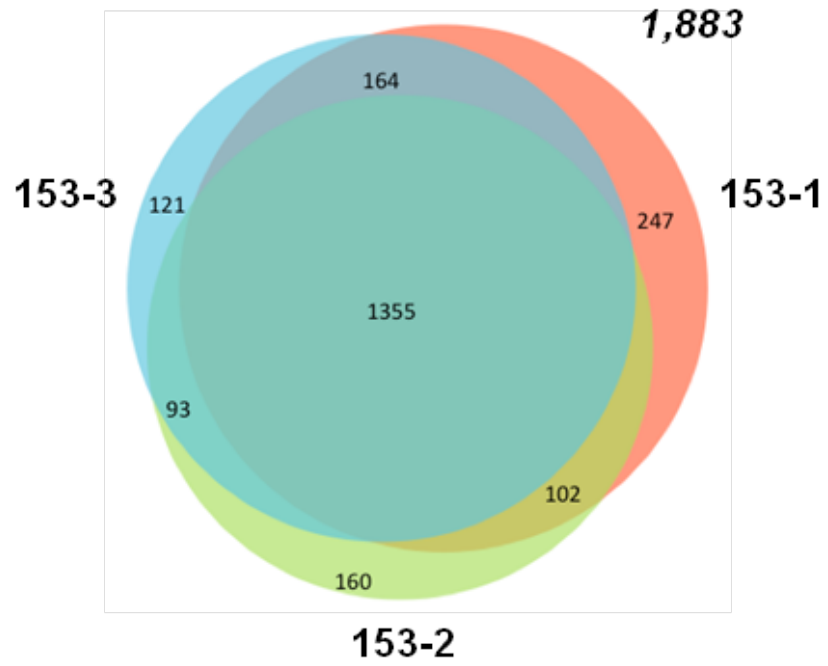
## FIGURES



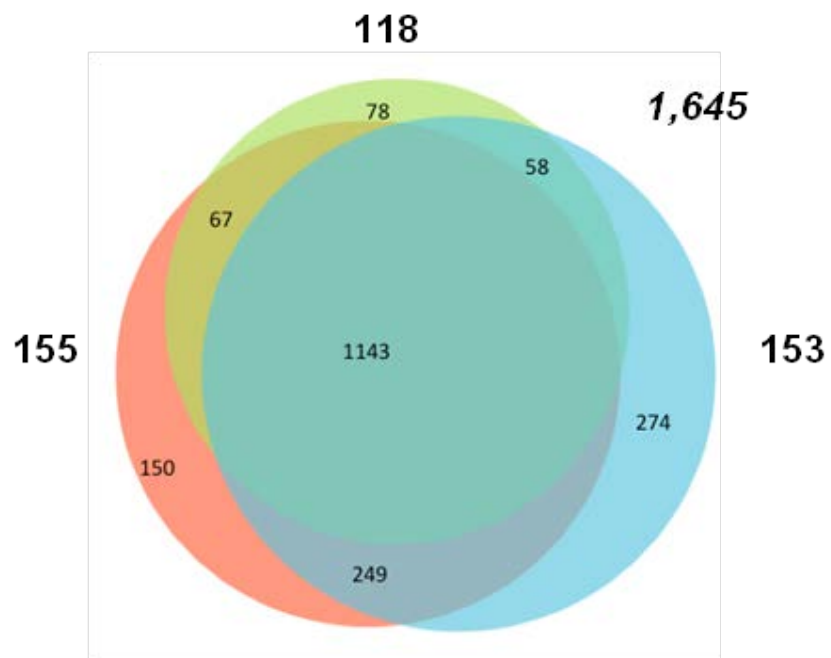
**Figure 1. Scheme of brain microvessel enrichment** (A) Flow chart of microvessel enrichment procedure where  $S_0$  is whole brain lysate,  $S_1$  is brain pellet after ficoll centrifugation and  $S_5$  is obtained from purified microvessels. (B) Confocal imaging of brain microvessels following mouse intravenous injection with FITC-lectin followed by microvessel extraction and isolation. Bar = 50 microns. (C) Fluorescent and phase imaging of DAPI-stained nuclei from  $S_1$  vs.  $S_5$  microvessel fractions. (D) Immunoblotting of  $S_0$  vs.  $S_5$  fractions with anti-P-glycoprotein (Pgp), anti-synaptophysin (Syn), or anti-aquaporin-4 (Aqp4) antibodies.

**Figure 2. Information on proteins identified by MudPIT from technical/biological replicates from the S5 microvessel fraction samples.** (A) Shown in a Venn diagram are the numbers of proteins identified by MudPIT analyses common to or distinct from each other in three technical replicates of the S5 microvessel fraction, 153-1, 153-2, and 153-3. Three single samples were prepared and then stored individually, and then analyzed independently by MudPIT. Inclusion criteria for the 1355 proteins common to all three replicates were  $\geq 5$  spectral counts and common identity to all three replicates, with 1,883 proteins (common + unique proteins) identified from all samples. Individual protein outliers correspond to 247 (153-1), 160 (153-2), and 121 (153-3) proteins from each replicate. Supplementary Table 1 contains the complete listing of all proteins identified and number of spectra from each replicate and all S5 microvessel fraction samples. See Materials and Methods for information on mass spectrometry and bioinformatic analyses. (B) A Venn diagram shows the numbers of proteins common to or distinct from each other in three biological replicates of the S5 microvessel fraction, 118, 154, and 153 (proteins common to 153-1, 153-2, and 153-3). Each S5 microvessel fraction sample was prepared as an individual sample prior to analyses. The criteria used for inclusion of proteins common to, 1149, or distinct from each sample, 78 (118), 150 (154) and 274 (153) were as discussed in A., 1,645 proteins (common + unique proteins) were identified from all samples (C) A comparison of the average of all spectral counts obtained from all mass spectrometry runs was compared to individual spectral counts obtained from each mass spectrometry run. The log scale regression plot presented the average of all spectral counts for all microvessel membrane fraction 5 analyses as compared to the spectral counts for individual analyses. The legend is on the plot for each technological replicate, 1A, 1B, and 1C, and the two biological replicates, 2 and 3.  $R^2 = 0.98624$ . (D) The cellular location of proteins identified from microvessel fractions, with numbers of proteins given after the category in the bar chart, for a given subcellular location relative to the total number of location-annotated Uniprot IDs for all five MudPIT analyses of the S5 membrane fractions. For each identified protein the International Protein Index (IPI) (Kersey *et al.*, 2004) identifiers were first mapped to Uniprot identifiers (IDs). The Uniprot Knowledgebase was then used to correlate IDs to subcellular location. Membrane refers to all proteins identified that can be designated as a membrane protein. The percentages do not add to 100%, as many proteins have more than one subcellular location. For additional information please see Material and Methods.

2A.

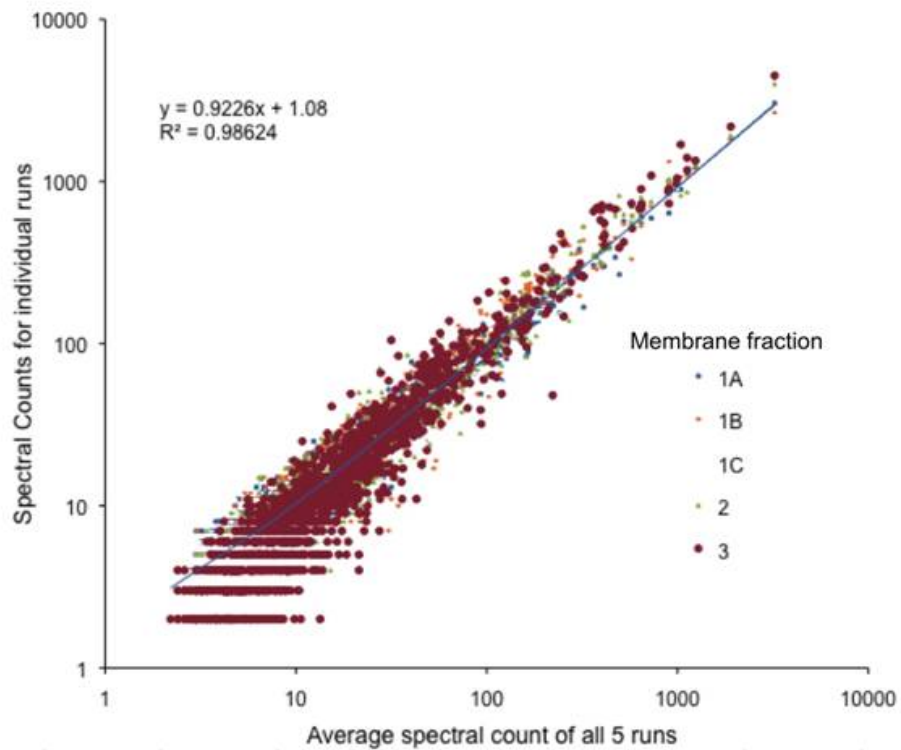


2B.





2C.



2D.

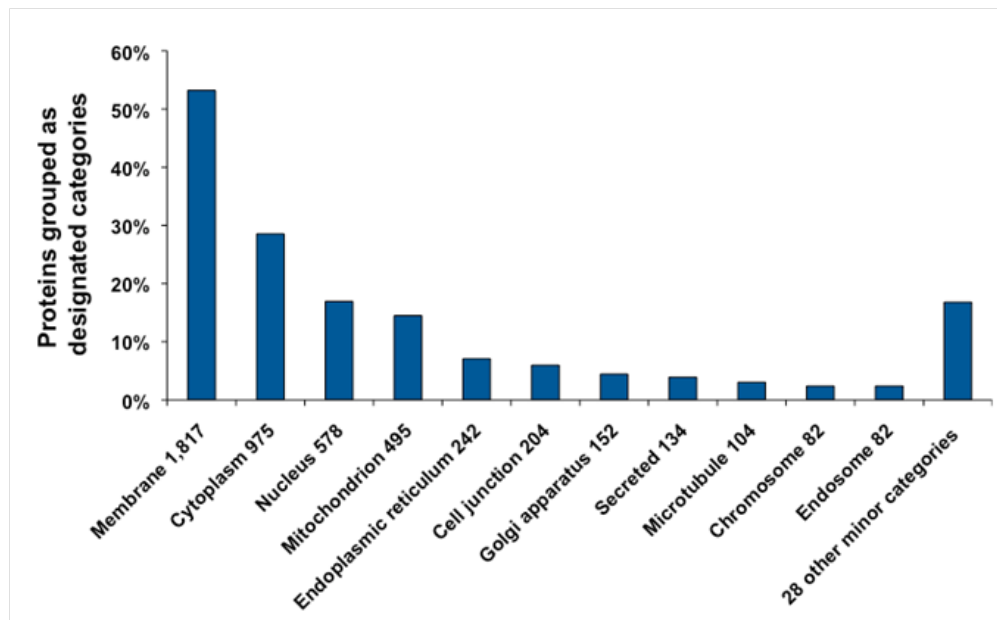


Figure 2. continued.

## TABLES

**Table 1.** Transporters and Channel Proteins

Protein	Gene	Mean	S.D.	Ref
IPI00752412	Atp1a3	1954.7	211.5	3
IPI00420569	Atp1a2	1249.7	99.1	2**, 5***
IPI00311682	Atp1a1	1135.0	79.9	
IPI00468481	Atp5b	1033.0	318.0	
IPI00130280	Atp5a1	849.0	158.4	
IPI00378485	Atp1a4	639.7	48.0	
IPI00115564	Slc25a4	568.0	50.5	1
IPI00875419	Slc1a2	500.0	110.0	5***
IPI00127841	Slc25a5	397.3	11.9	
IPI00857092	Slc4a4	211.0	13.1	5***
IPI00308162	Slc25a12	209.0	53.7	
IPI00114279	Slc1a3	208.3	35.3	7
IPI00121550	Atp1b1	184.3	46.2	
IPI00119113	Atp6v1b2	178.7	35.0	
IPI00831180	Atp2b2	175.0	16.1	
IPI00313475	Atp5c1	166.3	53.3	
IPI00553834	Abcb1a	150.0	46.7	1,2*,4,6,7
IPI00556827	Atp2b1	140.7	19.8	
IPI00407692	Atp6v1a	130.7	29.7	
IPI00341282	Atp5f1	121.3	35.0	
IPI00407499	Abat	118.0	41.6	
IPI00338964	Atp2a2	118.0	18.7	3
IPI00828950	Atp6v0a1	113.0	19.3	
IPI00751474	Atp2b4	107.7	28.9	
IPI00265299	Atp2b3	107.0	18.0	
IPI00877254	Slc12a5	91.0	6.1	1
IPI00124771	Slc25a3	88.3	8.7	3
IPI00109275	Slc25a22	79.7	16.3	
IPI00114641	Slc3a2	75.3	15.0	1,2*,6,7
IPI00313841	Atp6v0d1	69.7	23.7	
IPI00230507	Atp5h	58.7	5.0	
IPI00230680	Atp4a	56.3	8.1	6
IPI00118986	Atp5o	56.0	9.2	
IPI00315999	Atp6v1f	52.7	7.4	
IPI00123704	Atp1b2	51.0	14.9	2**
IPI00230754	Slc25a11	49.3	10.1	
IPI00119115	Atp6v1e1	49.3	5.5	
IPI00752501	Slc4a10	42.0	5.0	
IPI00120761	Slc4a1	39.0	9.2	3
IPI00311654	Atp2a1	32.0	27.9	
IPI00308691	Slc2a1	29.7	4.5	1,2*,4,6,7
IPI00626793	Cacna2d1	26.3	5.9	
IPI00131666	Slc1a6	25.7	3.1	1
IPI00111770	Atp5k	25.0	7.8	

1-Enerson and Drewes, 2006  
2-Daneman et al., 2010  
3-Lu et al., 2008  
4-Yousif et al., 2007  
5-Cahoy et al., 2008  
6-Kamiie et al., 2008  
7-Uchida et al., 2011  
8-Armulik et al., 2010

\*endothelial  
\*\*pericyte  
\*\*\*astrocyte

Protein	Gene	Mean	S.D.	Ref
IPI00135324	Slc12a2	21.7	4.0	1,2**
IPI00875016	Slc25a18	18.7	2.3	5***
IPI00227928	Slc6a1	17.7	2.3	1
IPI00322156	Slc38a3	17.0	7.5	1,2*
IPI00111686	Abcc4	17.0	3.6	1, 2*,6,7
IPI00308063	Slc9a3r2	16.7	4.7	1,2*
IPI00828762	Abca1	15.7	5.1	7
IPI00311200	Atp2a3	12.7	2.5	2**
IPI00137194	Slc16a1	11.7	4.7	1,2*,6,7
IPI00134191	Slc2a3	11.3	2.1	1,7
IPI00109311	Slc9a3r1	11.3	1.5	1
IPI00553576	Abcd3	11.0	3.0	1
IPI00109153	Slc17a7	10.0	4.6	1
IPI00229554	Slc6a17	10.0	2.6	2*
IPI00468924	Slc25a24	9.3	5.5	2*
IPI00221921	CD9	8.0	6.6	7
IPI00265568	Slc25a25	7.3	1.5	2*
IPI00120166	Slc30a1	7.3	5.9	1,2*
IPI00420244	Slc1a1	7.0	3.6	2*
IPI00468691	Abcg2	6.7	3.2	1,2*,4,6,7
IPI00120097	Slc27a1	6.3	1.5	1
IPI00624501	Asna1	6.0	5.6	1
IPI00121634	Slc7a1	6.0	5.3	1,2*,7
IPI00122522	Ggt1	5.3	3.1	6*

1-Enerson and Drewes, 2006

2-Daneman et al., 2010

3-Lu et al., 2008

4-Yousif et al., 2007

5-Cahoy et al., 2008

6-Kamiie et al., 2008

7-Uchida et al., 2011

8-Armulik et al., 2010

\*endothelial

\*\*pericyte

\*\*\*astrocyte

Table 1. Continued

Protein	Gene	Mean	S.D.	Ref
<i>IPI00225534</i>	<i>Atp13a5</i>	4.7	1.5	2**
<i>IPI00622815</i>	<i>Slc9a1</i>	4.3	1.5	1
<i>IPI00125970</i>	<i>Abca7</i>	4.0	4.0	1,7
<i>IPI00459383</i>	<i>Abcb8</i>	4.0	2.0	1
<i>IPI00136334</i>	<i>Abcc9</i>	3.7	3.2	1,2**,8
<i>IPI00317074</i>	<i>Slc25a10</i>	3.7	2.1	1
<i>IPI00828826</i>	<i>Slc44a1</i>	3.7	1.5	2*
<i>IPI00553414</i>	<i>Slc5a6</i>	3.3	2.9	1
<i>IPI00131584</i>	<i>Slc25a20</i>	3.0	1.7	1,2*
<i>IPI00128358</i>	<i>Insr</i>	3.0	3.6	7
<i>IPI00858037</i>	<i>Abca9</i>	2.7	3.1	2**
<i>IPI00273801</i>	<i>Slc39a10</i>	2.7	0.6	2*
<i>IPI00750694</i>	<i>Slc39a12</i>	2.7	3.1	5***
<i>IPI00169896</i>	<i>Slc44a2</i>	2.7	2.3	2*
<i>IPI00470963</i>	<i>Slc4a3</i>	2.3	2.1	1
<i>IPI00120094</i>	<i>Abcb6</i>	2.0	1.7	1
<i>IPI00112616</i>	<i>Abca2</i>	1.7	1.5	1,7
<i>IPI00331175</i>	<i>Slc12a7</i>	1.7	1.5	2*
<i>IPI00387413</i>	<i>Slc22a8</i>	1.7	1.5	2*,6
<i>IPI00742415</i>	<i>Abca3</i>	1.3	2.3	7
<i>IPI00353962</i>	<i>Slc16a2</i>	1.3	1.2	2*
<i>IPI00331577</i>	<i>Slc7a5</i>	1.3	2.3	2*,6
<i>IPI00314205</i>	<i>Slc6a20</i>	1.0	1.7	1
<i>IPI00624726</i>	<i>Abcc8</i>	0.7	1.2	1,2,7
<i>IPI00830169</i>	<i>Atp7a</i>	0.7	1.2	2*
<i>IPI00453817</i>	<i>Slc38a2</i>	0.7	1.2	1,2*,6
<i>IPI00109242</i>	<i>Slc7a3</i>	0.7	1.2	2*

1-Enerson and Drewes, 2006

2-Daneman et al., 2010

3-Lu et al., 2008

4-Yousif et al., 2007

5-Cahoy et al., 2008

6-Kamiie et al., 2008

7-Uchida et al., 2011

8-Armulik et al., 2010

\*endothelial

\*\*pericyte

\*\*\*astrocyte

*Italics: Mean spectral count < 5*

Table 1. Continued

**Table 2A.** Tight Junction Transmembrane Proteins

<b>Protein</b>	<b>Gene</b>	<b>Mean</b>	<b>S.D.</b>	<b>Ref</b>
IPI00122971	Ncam1	205	51	3
IPI00850457	Cadm2	42.7	9.1	
IPI00380273	Gja1	35.3	7	1
IPI00121378	Alcam	32.7	7.4	1
IPI00115762	L1cam	31.7	2.5	3
IPI00136135	Ctnnd2	22.3	3.8	
IPI00395042	Nrcam	21.7	4.2	1
IPI00127556	Ncam2	21.7	2.5	
IPI00471176	Hepacam	21.3	8.5	
IPI00856723	Cadm1	19.3	7.1	
IPI00857780	Cadm3	19	6.1	
IPI00877299	Icam5	18	3.6	1
IPI00229475	Jup	17	1	1,3
IPI00473693	Pkp4	16.7	5	1
IPI00749860	Cdh2	14.3	4	2
IPI00875877	CD47	10.7	1.2	
IPI00153840	Cadm4	9.7	4.9	
IPI00406901	Pecam1	8	3.5	
IPI00120295	Cldn11	7.7	1.2	1,3
IPI00154057	Pcdh1	6.7	1.5	5
<i>IPI00117828</i>	<i>Ocln</i>	<i>4.3</i>	<i>1.5</i>	<i>2,6</i>
<i>IPI00775975</i>	<i>Cdh13</i>	<i>4.3</i>	<i>1.2</i>	<i>1,2</i>
<i>IPI00117424</i>	<i>Icam2</i>	<i>3.3</i>	<i>1.5</i>	<i>1,2,6</i>
<i>IPI00410802</i>	<i>CD44</i>	<i>3</i>	<i>1</i>	<i>4,5</i>
<i>IPI00111902</i>	<i>Gjb6</i>	<i>2.3</i>	<i>2.1</i>	<i>4</i>
<i>IPI00138180</i>	<i>Cdh5</i>	<i>2.3</i>	<i>2.1</i>	<i>1,2,5,6</i>
<i>IPI00138190</i>	<i>Cdh11</i>	<i>2</i>	<i>2</i>	<i>1,2,5</i>
<i>IPI00755700</i>	<i>Pcdh9</i>	<i>2</i>	<i>2</i>	<i>5</i>
<i>IPI00469123</i>	<i>Mcam</i>	<i>1.7</i>	<i>1.5</i>	<i>1,4</i>
<i>IPI00848632</i>	<i>Pcdh10</i>	<i>1.3</i>	<i>2.3</i>	<i>5</i>
<i>IPI00122973</i>	<i>Icam1</i>	<i>1.3</i>	<i>1.2</i>	<i>6</i>
<i>IPI00356667</i>	<i>Pcdh17</i>	<i>1.3</i>	<i>1.2</i>	<i>5</i>
<i>IPI00119893</i>	<i>Cldn5</i>	<i>1</i>	<i>1.7</i>	<i>1,2</i>
<i>IPI00346482</i>	<i>Cdh10</i>	<i>1</i>	<i>1.7</i>	<i>2</i>
<i>IPI00775835</i>	<i>Cdh6</i>	<i>1</i>	<i>1.7</i>	<i>2,5</i>
<i>IPI00109330</i>	<i>Cdh15</i>	<i>0.7</i>	<i>1.2</i>	<i>2</i>
<i>IPI00622727</i>	<i>Jam2</i>	<i>0.7</i>	<i>1.2</i>	<i>2</i>
<i>IPI00653571</i>	<i>Vcam1</i>	<i>0.7</i>	<i>1.2</i>	<i>4,6</i>

1-Enerson and Drewes, 2006

2-Daneman et al., 2010

3-Lu et al., 2008

4-Yousif et al., 2007

5-Cahoy et al., 2008

6-Kamiie et al., 2008

7-Uchida et al., 2011

8-Armulik et al., 2010

*Italics: Mean spectral count < 5*

**Table 2B.** Tight Junction Cytoplasmic Adapters

<b>Protein</b>	<b>Gene</b>	<b>Mean</b>	<b>S.D.</b>	<b>Ref</b>
IPI00125899	Ctnnb1	55	4.4	2,4
IPI00119870	Ctnna2	49	3	
IPI00112963	Ctnna1	43.3	7.6	
IPI00874522	Tjp1	24.3	10	1,2,3
IPI00125861	Dlg1	20	3	2
IPI00752108	Ctnnd1	16.7	4.2	
IPI00126827	Esam1	9.7	5.5	1,2
IPI00136498	Lin7c	9	1.7	2
IPI00137706	Mpp1	8.7	3.1	2
IPI00124051	Mpp5	6	2	2
IPI00323349	Tjp2	6	2.6	1,2
IPI00330186	Dlgap4	5	1	2
IPI00121362	F11r	5	3	1,2
<i>IPI00877180</i>	<i>Cgnl1</i>	<i>3.3</i>	<i>1.5</i>	<i>2</i>
<i>IPI00761693</i>	<i>Mpp7</i>	<i>2</i>	<i>2</i>	<i>2</i>
<i>IPI00129388</i>	<i>Scrib</i>	<i>2</i>	<i>2</i>	<i>2</i>
<i>IPI00399953</i>	<i>Wnk1</i>	<i>1</i>	<i>1.7</i>	<i>2</i>
<i>IPI00761515</i>	<i>Magi3</i>	<i>0.7</i>	<i>1.2</i>	<i>2</i>
<i>IPI00762889</i>	<i>Tjp3</i>	<i>0.7</i>	<i>1.2</i>	<i>2</i>

1-Enerson and Drewes, 2006

2-Daneman et al., 2010

3-Lu et al., 2008

4-Yousif et al., 2007

5-Cahoy et al., 2008

6-Kamiie et al., 2008

7-Uchida et al., 2011

8-Armulik et al., 2010

*Italics: Mean spectral count < 5***Table 2C.** Integrins

<b>Protein</b>	<b>Gene</b>	<b>Mean</b>	<b>S.D.</b>	<b>Ref</b>
IPI00132474	Itgb1	44.7	9.6	4
IPI00620873	Itga6	25.3	4.2	
IPI00466371	Itga1	22.3	6.4	4
IPI00877232	Itga2b	11.7	5.8	
IPI00466276	Itga7	11.3	1.5	
IPI00857195	Itgav	8.7	1.5	6
IPI00468674	Itga3	5.3	0.7	

1-Enerson and Drewes, 2006

2-Daneman et al., 2010

3-Lu et al., 2008

4-Yousif et al., 2007

5-Cahoy et al., 2008

6-Kamiie et al., 2008

7-Uchida et al., 2011

8-Armulik et al., 2010

**Table 3.** Extracellular Matrix Proteins

<b>Protein</b>	<b>Gene</b>	<b>Mean</b>	<b>S.D.</b>	<b>Ref</b>
IPI00123058	Cntn1	125.0	16.5	8*
IPI00515360	Hspg2	60.3	44.0	
IPI00227126	Tnr	53.7	4.2	
IPI00854028	Cntnap1	46.0	14.8	
IPI00400016	Lamc1	43.0	36.4	
IPI00109612	Lamb2	33.0	24.6	
IPI00756745	Lama2	30.3	34.5	2**
IPI00648938	Agrn	26.7	20.4	
IPI00111793	Nid1	24.7	17.5	
IPI00676957	Lama5	24.0	23.6	
IPI00420554	Cntnap2	18.7	3.8	
IPI00620601	Astn1	16.7	0.6	
IPI00122273	Dag1	16.7	9.0	
IPI00113726	Lama1	13.3	7.4	
IPI00279079	Fgb	13.0	6.1	
IPI00119970	Cntn2	12.7	3.8	
IPI00338785	Lamb1	12.3	6.0	
IPI00223446	Lama4	12.0	5.3	
IPI00109588	Col4a1	11.7	15.3	
IPI00121038	Vcan	11.0	3.5	
IPI00338452	Col4a2	10.7	11.0	
IPI00719919	Nid2	10.0	9.2	
IPI00270915	Cntn3	8.3	3.5	
IPI00869394	Bcan	6.3	2.1	
IPI00115522	Fga	6.3	1.2	
IPI00113539	Fn1	6.3	5.5	
IPI00830749	Col6a3	5.0	5.6	
<i>IPI00469123</i>	<i>Mcam</i>	<i>1.7</i>	<i>1.5</i>	<i>5*</i>
<i>IPI00480306</i>	<i>Cntn4</i>	<i>0.7</i>	<i>1.2</i>	<i>5*</i>
<i>IPI00129240</i>	<i>Vtn</i>	<i>0.7</i>	<i>1.2</i>	<i>2**</i>

1-Enerson and Drewes, 2006

2-Daneman et al., 2010

3-Lu et al., 2008

4-Yousif et al., 2007

5-Cahoy et al., 2008

6-Kamiie et al., 2008

7-Uchida et al., 2011

8-Armulik et al., 2010

\*endothelial

\*\*pericyte

\*\*\*astrocyte

*Italics: Mean spectral count < 5*

**Table 4A.** Endothelial Proteins

<b>Protein</b>	<b>Gene</b>	<b>Mean</b>	<b>S.D.</b>	<b>Ref</b>
IPI00227299	Vim	191.3	63.0	3,5,8
IPI00115240	Mbp	75.3	28.4	1,3
IPI00119063	Lrp1	57.0	12.2	11
IPI00228618	Gnaq	37.3	0.6	7
IPI00553798	Ahnak	37.3	22.9	5
IPI00121378	Alcam	32.7	7.4	5
IPI00224776	Hspa12b	31.7	14.6	2
IPI00620256	Lmna	24.7	6.7	4
IPI00131138	Flna	24.0	6.9	8
IPI00313884	Podxl	23.0	1.0	4
IPI00124700	Tfrc	22.3	12.9	1,11
IPI00408495	Bsg	20.0	3.5	2,5
IPI00653398	Flot2	18.0	4.0	7
IPI00118569	Gna13	16.0	3.5	5
IPI00648927	Cita	14.3	2.5	5
IPI00311671	Nos3	13.0	2.6	6
IPI00130589	Sod1	12.7	4.7	6
IPI00134131	Scp2	12.3	1.5	4
IPI00114594	Rras	11.7	3.8	5
IPI00127942	Dstn	10.7	1.5	4
IPI00331609	Vwa1	10.7	6.0	2
IPI00762435	Fyn	10.3	2.3	7
IPI00117829	Cav1	9.7	5.0	5,7
IPI00129253	Ly75	8.0	3.0	2
IPI00131881	Adam10	7.7	4.9	7
IPI00278804	Scamp1	7.7	1.5	4
IPI00315280	Sema7a	6.7	0.6	2
IPI00120066	Prom1	6.3	1.5	2
IPI00133608	Ociad1	5.7	0.6	4
IPI00849927	Rpl10	5.7	4.0	4
IPI00113377	Rplp1	5.0	4.4	4

1-Enerson and Drewes, 2006

2-Daneman et al., 2010

3-Lu et al., 2008

4-Agarwal et al. 2010

5-Cayrol et al., 2007

6-Cunnea et al., 2010

7-Li et al., 2009

8-Pottiez et al., 2009

9-Cahoy et al., 2008

10-Armulik et al., 2010

11-Uchida et al., 2011

12-Yousif et al., 2007



<b>Protein</b>	<b>Gene</b>	<b>Mean</b>	<b>S.D.</b>	<b>Ref</b>	
<i>IPI00338094</i>	<i>Bmpr2</i>	4.0	2.0	3	
<i>IPI00762331</i>	<i>Cobll1</i>	4.0	3.6	2	1-Enerson and Drewes, 2006
<i>IPI00828826</i>	<i>Slc44a1</i>	3.7	1.5	2	2-Daneman et al., 2010
<i>IPI00322245</i>	<i>Fmo2</i>	3.3	2.9	2	3-Lu et al., 2008
<i>IPI00222369</i>	<i>Itih5</i>	3.3	3.5	2	4-Agarwal et al. 2010
<i>IPI00111600</i>	<i>Spock2</i>	3.3	0.6	2	5-Cayrol et al., 2007
<i>IPI00321477</i>	<i>Eng</i>	3.0	2.6	2,3	6-Cunnea et al., 2010
<i>IPI00128358</i>	<i>Insr</i>	3.0	3.6	11	7-Li et al., 2009
<i>IPI00129423</i>	<i>Thbd</i>	3.0	3.6	6	8-Pottiez et al., 2009
<i>IPI00798576</i>	<i>Vwf</i>	2.7	0.6	1	9-Cahoy et al., 2008
<i>IPI00876541</i>	<i>Gpr116</i>	2.3	2.5	2,7	10-Armulik et al., 2010
<i>IPI00221547</i>	<i>Tek</i>	2.3	2.1	1	11-Uchida et al., 2011
<i>IPI00458230</i>	<i>Mfsd2</i>	2.0	2.0	2	12-Yousif et al., 2007
<i>IPI00131498</i>	<i>Cd93</i>	1.7	1.5	2	
<i>IPI00467127</i>	<i>Chid1</i>	1.7	2.9	4	
<i>IPI00625441</i>	<i>Tpd52</i>	1.7	1.5	2	
<i>IPI00133257</i>	<i>CD34</i>	1.3	2.3	1	<i>Italics: Mean spectral count &lt; 5</i>
<i>IPI00124497</i>	<i>Flt1</i>	1.3	1.2	1,2,6	
<i>IPI00230191</i>	<i>Gna12</i>	1.3	2.3	7	
<i>IPI00123996</i>	<i>Nrp1</i>	1.3	2.3	7	
<i>IPI00137172</i>	<i>Ppapdc2</i>	1.3	2.3	2	
<i>IPI00226730</i>	<i>Clic5</i>	1.0	1.7	2	

Table 4A. Continued

**Table 4B.** Astrocyte Proteins

Protein	Gene	Mean	S.D.	Ref
IPI00117042	Gfap	282.0	5.2	1,3,9,12
IPI00119458	Aldoc	66.3	5.7	9
IPI00230738	Aqp4	51.3	12.1	9
IPI00153317	Aldh111	17.0	6.9	9
IPI00453834	Acsbg1	16.0	4.0	9
IPI00131995	Hapln1	11.0	7.2	9
IPI00322760	Prodh	7.7	2.1	9
IPI00119130	1500005I02Rik	6.7	2.5	9
<i>IPI00830222</i>	<i>Mlc1</i>	4.7	2.5	9
<i>IPI00118069</i>	<i>F3</i>	1.3	1.2	9
<i>IPI00329896</i>	<i>Pla2g7</i>	1.3	2.3	9

1-Enerson and Drewes, 2006  
2-Daneman et al., 2010  
3-Lu et al., 2008  
4-Agarwal et al. 2010  
5-Cayrol et al., 2007  
6-Cunnea et al., 2010  
7-Li et al., 2009  
8-Pottiez et al., 2009  
9-Cahoy et al., 2008  
10-Armulik et al., 2010  
11-Uchida et al., 2011  
12-Yousif et al., 2007

**Table 4C.** Pericyte Proteins

Protein	Gene	Mean	S.D.	Ref
IPI00117043	Acta2	133.3	116.4	1
IPI00416906	Gnas	33.3	2.5	10
IPI00134585	Enpep	31.0	12.8	2,10
IPI00756745	Lama2	30.3	34.5	2
IPI00113517	Ctsb	26.0	4.6	10
IPI00153202	Ace2	21.7	6.4	2
IPI00136716	Grm3	14.3	0.6	2
IPI00126186	Mrc1	12.0	9.2	10
IPI00133208	Hspa1a	6.0	10.4	10
IPI00344142	Lin7a	5.3	5.0	2
IPI00118880	Gucy1b3	5.0	1.0	2
<i>IPI00128915</i>	<i>Cspg4</i>	4.3	1.5	2
<i>IPI00229516</i>	<i>Itgb5</i>	4.3	2.3	2
<i>IPI00121827</i>	<i>Pdgfrb</i>	2.7	1.2	2
<i>IPI00329896</i>	<i>Pla2g7</i>	1.3	2.3	2
<i>IPI00123125</i>	<i>Xtrp3s1</i>	1.0	1.7	2
<i>IPI00221980</i>	<i>Pde8b</i>	0.7	1.2	2,10

1-Enerson and Drewes, 2006  
2-Daneman et al., 2010  
3-Lu et al., 2008  
4-Agarwal et al. 2010  
5-Cayrol et al., 2007  
6-Cunnea et al., 2010  
7-Li et al., 2009  
8-Pottiez et al., 2009  
9-Cahoy et al., 2008  
10-Armulik et al., 2010  
11-Uchida et al., 2011  
12-Yousif et al., 2007

**SUPPLEMENTARY TABLES**

**Supplementary Table 1.** Complete listing of all proteins identified from the microvessel S5 fraction

Gene Name	Biol Reps RSD	Biol Reps SD	Tech Reps RSD	Tech Reps Standard Deviation (SD)	Avg of all 5 SC	Median of All 5 Samples	Total Spectral Counts (SC)	Spectral Counts for 118-1	Spectral Counts for 117-155-1	Spectral Counts for 117-153-3	Spectral Counts for 117-153-2	Spectral Counts for 117-153-1	IPI Identifier
RP23-403O15.4	38.88%	1352.0	21.79%	554.6	3225.0	3035	16125	4494	3995	1943	2658	3035	IP 00273646.9
Atp1a3	10.82%	211.5	2.35%	42.3	1899.6	1845	9498	2181	1921	1762	1845	1789	IP 00752412.1
Atp1a2	7.93%	99.1	3.69%	44.2	1239.6	1227	6198	1346	1255	1148	1227	1222	IP 00420569.2
Atp1a1	7.04%	79.9	4.37%	47.5	1124.0	1137	5620	1175	1187	1043	1078	1137	IP 00468481.2
Atp5b	30.79%	318.0	23.65%	263.8	1121.4	1145	5607	1400	861	838	1363	1145	IP 00648173.1
Cltc	49.24%	531.3	19.00%	170.0	1039.0	892	5195	1690	821	726	1066	892	IP 00403810.2
Tuba1a	8.68%	87.4	5.08%	48.5	995.8	1004	4979	1045	1069	907	1004	954	IP 00311682.5
Tuba1b	8.79%	86.6	9.97%	95.0	982.0	974	4910	974	1078	906	1062	890	IP 00110753.1
Atp5a1	18.65%	158.4	39.47%	362.0	902.2	787	4511	731	1029	787	1326	638	IP 00462072.3
Tuba4a	9.69%	86.3	14.30%	125.7	902.0	900	4510	900	973	801	1024	812	IP 00117348.4
Tuba1c	5.26%	46.7	11.21%	99.3	893.0	869	4465	869	940	852	997	807	IP 00117350.1
Actb	37.57%	293.5	16.63%	99.7	727.8	702	3639	1088	753	503	702	593	IP 00798592.1
Spna2	11.88%	79.1	6.56%	39.9	648.8	653	3244	729	691	577	653	594	IP 00378485.2
LOC100044223	9.73%	63.5	12.09%	78.9	644.2	683	3221	683	579	694	703	562	IP 00874964.1
ENSMUSG0000057666	39.01%	271.0	21.90%	111.2	644.0	606	3220	898	799	387	530	606	IP 00130280.1
Atp1a4	7.51%	48.0	7.09%	43.2	631.4	651	3157	681	651	587	658	580	IP 00850779.1
Slc25a4	8.89%	50.5	4.54%	26.7	578.6	576	2893	514	614	576	619	570	IP 00875419.1
Vdac1	17.43%	106.3	35.67%	187.3	573.6	561	2868	732	561	537	332	706	IP 00114375.2
LOC100040109	39.01%	271.0	92.71%	306.9	538.0	606	2690	898	799	387	0	606	IP 00110850.1
Slc1a2	22.00%	110.0	11.86%	61.8	522.4	544	2612	423	626	451	544	568	IP 00775863.1
Hist1h2bc	52.60%	268.8	52.07%	219.2	494.4	390	2472	390	819	324	672	267	IP 00169463.1
Gnb1	41.04%	214.2	24.75%	88.1	471.4	451	2357	674	615	277	451	340	IP 00122549.1
Tubb4	44.44%	205.9	17.31%	64.0	440.2	398	2201	694	398	298	421	390	IP 00461514.4

IP100115564.5	0	648	1517	0	2165	0	433.0	374.1	173.21%	761.2	105.48%	Hist1h2ao
IP00127841.3	434	362	412	468	2081	412	416.2	36.2	9.05%	53.0	12.81%	Spnb2
IP00109073.5	382	407	384	401	2076	401	415.2	63.3	14.71%	11.9	3.00%	Slc25a5
IP00120716.3	300	241	533	553	2067	440	413.4	102.2	31.26%	174.6	39.48%	Gnb2
IP00319830.7	299	339	486	451	2033	451	406.6	82.7	22.64%	76.8	18.05%	Dpysl2
IP00117352.1	368	305	387	714	2005	368	401.0	68.6	22.76%	246.5	55.52%	Tubb2c
IP00415403.1	363	354	371	666	1992	363	398.4	69.7	21.90%	219.1	51.54%	Tubb5
IP00331704.7	314	353	318	578	1972	353	394.4	47.8	13.31%	141.1	33.89%	Stxbp1
IP00109061.1	344	288	355	689	1888	344	377.6	66.3	23.55%	244.8	58.47%	Tubb2b
IP00338039.1	304	292	352	654	1814	304	362.8	50.0	18.57%	225.9	55.64%	Tubb2a
IP00387413.2	0	674	0	1053	1727	0	345.4	389.1	173.21%	607.9	173.21%	Actg1
IP00319602.3	168	434	530	260	1608	260	321.6	141.8	51.99%	170.0	50.70%	Hist2h2bb
IP00187543.3	263	522	250	259	1602	263	320.4	138.4	37.98%	31.2	11.46%	Hspd1
IP00117042.3	262	366	282	312	1540	312	308.0	52.1	16.51%	19.3	6.34%	Eho2
IP00308885.6	382	197	429	262	1530	262	306.0	94.1	33.63%	97.0	30.60%	Cnp
IP00136703.1	272	358	276	285	1476	285	295.2	46.4	15.20%	5.2	1.84%	Gfap
IP00308162.3	270	285	360	239	1385	270	277.0	27.9	10.64%	72.3	26.13%	Ckb
IP00467841.6	257	403	343	208	1350	257	270.0	73.4	21.95%	103.8	45.11%	Nefl
IP00857092.1	204	252	185	415	1271	215	254.2	34.5	16.16%	125.0	46.02%	Actc1
IP00230192.5	196	445	242	147	1268	238	253.6	132.5	45.01%	53.7	25.71%	Slc25a12
IP00114279.1	187	203	155	476	1220	199	244.0	24.4	13.45%	174.0	62.90%	Tubb6
IP00119113.3	155	424	186	190	1204	190	240.8	136.5	49.46%	35.3	16.93%	Slc1a3
IP00121550.3	235	250	221	254	1171	235	234.2	19.7	8.48%	22.5	9.84%	Uqrcr1
IP00330000.4	173	0	516	233	1147	225	229.4	117.8	88.79%	165.7	51.05%	Hist2h2aa2
IP00114593.1	172	195	137	381	1117	195	223.4	29.2	17.38%	123.0	49.20%	Calm2
IP00323592.2	172	195	137	381	1117	195	223.4	29.2	17.38%	123.0	49.20%	Calm2
IP00228548.6	268	343	243	48	1108	243	221.6	52.0	18.28%	103.6	62.52%	Hbb-b1
IP00122971.1	187	284	226	202	1104	205	220.8	48.8	21.01%	13.1	6.20%	Slc4a4
IP00117043.1	178	257	193	244	1084	212	216.8	42.0	20.04%	25.8	11.91%	Mdh2
IP00653598.2	243	185	130	236	1080	236	216.0	56.5	30.38%	79.7	36.65%	Gnao1
IP00135965.2	174	264	190	163	1053	190	210.6	48.0	22.94%	51.2	24.96%	Ncam1
IP00761696.2	186	183	149	207	1020	186	204.0	20.6	11.90%	73.5	33.88%	Tubb3
IP00341282.2	180	263	157	153	1017	180	203.4	48.2	20.46%	63.0	32.91%	Vim
IP00313475.1	189	201	149	291	985	189	197.0	23.9	13.13%	80.3	40.49%	Aco2

IP00762198.2	154	174	146	242	246	962	174	192.4	14.4	9.13%	56.6	26.79%	ENSMUSG00000060832
IP00122928.1	171	228	131	212	210	952	210	190.4	48.7	27.59%	46.2	25.06%	Alp1b1
IP00407130.4	296	132	166	349	0	943	166	188.6	86.6	43.72%	174.6	101.69%	LOC640374
IP00831180.1	136	255	143	180	213	927	180	185.4	66.8	37.51%	35.0	19.60%	Atp6v1b2
IP00406447.3	168	265	180	112	183	908	180	181.6	52.9	25.88%	40.2	25.36%	Ina
IP00621789.1	133	221	190	158	177	879	177	175.8	44.6	24.62%	16.1	9.20%	Atp2b2
IP00227299.6	204	252	185	215	0	856	204	171.2	34.5	16.16%	116.4	87.33%	Acta2
IP00283611.1	160	236	194	101	155	846	160	169.2	38.1	19.36%	46.7	31.13%	Abcb1a
IP00378063.1	142	212	154	178	153	839	154	167.8	37.4	22.11%	14.2	8.75%	Hk1
IP00116074.1	188	228	220	194	0	830	194	166.0	21.2	9.98%	120.2	87.11%	Eno3
IP00751677.3	127	204	134	174	190	829	174	165.8	42.6	27.47%	28.8	17.38%	Dpysl3
IP00128296.1	89	224	108	273	130	824	130	164.8	73.1	52.07%	89.6	52.60%	Hist1h2bf
IP00876255.1	172	273	161	108	95	809	161	161.8	61.7	30.56%	35.0	28.81%	Atp5f1
IP00776087.1	84	217	108	265	130	804	130	160.8	70.9	51.99%	85.0	50.70%	Hist3h2ba
IP00626385.4	170	231	174	83	146	804	170	160.8	34.1	17.80%	46.6	34.70%	Nefm
IP00323357.3	199	180	148	149	121	797	149	159.4	25.8	14.67%	15.9	11.40%	Dync1h1
IP00554928.3	117	65	57	248	299	786	117	157.2	32.6	40.89%	127.6	63.36%	4732495G21Rik
IP00273801.3	0	0	0	785	0	785	0	157.0	0.0	#DIV/0!	453.2	173.21%	LOC100041204
IP00323800.6	131	164	131	176	178	780	164	156.0	19.1	13.42%	26.6	16.44%	Pkm2
IP00459725.2	100	167	118	269	121	775	121	155.0	34.7	27.02%	86.3	50.98%	Hist2h2ab
IP00119876.1	158	149	123	153	183	766	153	153.2	18.2	12.68%	30.0	19.61%	Ap2b1
IP00112251.1	112	155	193	201	105	766	155	153.2	40.5	26.43%	53.3	32.02%	Atp5c1
IP00122547.1	180	118	162	162	139	761	162	152.2	31.9	20.80%	13.3	8.60%	Vdac2
IP00131695.3	180	174	135	140	126	755	140	151.0	24.4	14.99%	7.1	5.31%	Gnai2
IP00272878.6	107	209	143	173	117	749	143	149.8	51.7	33.81%	28.0	19.42%	Dnm1
IP00331182.4	144	160	167	108	129	708	144	141.6	11.8	7.51%	29.9	22.21%	Plec1
IP00558827.1	128	145	173	129	130	705	130	141.0	22.7	15.28%	25.1	17.44%	Alb
IP00553774.3	149	124	132	126	168	699	132	139.8	12.8	9.46%	22.7	16.00%	Nduif1
IP00553834.3	144	107	97	157	188	693	144	138.6	24.8	21.34%	46.3	31.40%	Hspa8
IP00652902.1	128	128	106	139	168	669	128	133.8	12.7	10.53%	31.0	22.53%	Ckmt1
IP00407692.3	114	159	113	114	165	665	114	133.0	26.3	20.42%	29.7	22.76%	Atp6v1a
IP00221723.1	656	0	0	0	0	656	0	131.2	378.7	173.21%	0.0	#DIV/0!	Tuba8
IP00230351.1	123	184	161	88	89	645	123	129.0	30.8	19.75%	41.9	37.15%	Nefh
IP00338854.3	101	188	95	93	166	643	101	128.6	52.0	40.66%	41.6	35.24%	Abat

IP00123058.1	110	108	162	137	123	640	123	128.0	30.6	24.17%	19.8	14.05%	Alp2b1
IP00114642.4	117	145	133	136	106	637	133	127.4	14.0	10.67%	16.5	13.22%	Cntn1
IP00131376.5	81	195	126	142	91	635	126	127.0	57.4	42.85%	26.1	21.80%	Cox5a
IP00338964.3	113	107	90	121	204	635	113	127.0	11.9	11.55%	58.9	42.61%	Glud1
IP00114241.2	109	83	63	128	244	627	109	125.4	23.1	27.14%	91.7	63.23%	Nsf
IP00828950.1	104	176	99	105	135	619	105	123.8	43.1	34.10%	19.3	17.07%	Alp6v0a1
IP00623951.3	100	0	119	269	123	611	119	122.2	63.9	87.57%	85.5	50.18%	H2afx
IP00221528.1	119	136	115	101	138	609	119	121.8	11.2	9.04%	18.7	15.83%	Alp2a2
IP00132653.1	59	248	170	73	49	599	73	119.8	95.0	59.74%	64.1	65.82%	Syp
IP00117312.1	96	186	70	121	121	594	121	118.8	60.9	51.88%	29.4	28.31%	ldh3a
IP00116498.1	85	103	58	207	138	591	103	118.2	22.6	27.62%	74.6	55.51%	H2afy
IP00308882.4	106	66	146	136	137	591	136	118.2	40.0	37.74%	5.5	3.94%	Ywhaz
IP00129618.1	76	135	110	107	150	578	110	115.6	29.6	27.68%	24.0	19.62%	Syt1
IP00134344.6	123	124	106	95	124	572	123	114.4	10.1	8.60%	14.6	13.51%	Gpd2
IP00845652.1	138	118	105	91	118	570	118	114.0	16.6	13.81%	13.5	12.90%	Got2
IP00467152.4	85	108	164	106	106	569	106	113.8	40.6	34.14%	33.5	26.72%	Camk2a
IP00788324.1	113	131	114	144	62	564	114	112.8	10.1	8.48%	41.5	38.90%	Myh9
IP00464317.3	101	145	83	82	147	558	101	111.6	31.9	29.08%	37.2	35.81%	Cs
IP00751474.1	105	146	97	86	112	546	105	109.2	26.3	22.66%	13.1	13.27%	Ogdh
IP00230730.4	148	116	75	108	98	545	108	109.0	36.6	32.38%	16.9	18.07%	Gnai1
IP00319509.5	108	113	111	103	102	537	108	107.4	2.5	2.27%	4.9	4.68%	Spnb3
IP00877254.1	123	124	122	91	76	536	122	107.2	1.0	0.81%	23.5	24.35%	Spnb1
IP00120193.1	62	206	123	78	64	533	78	106.6	72.3	55.46%	30.8	34.90%	Cdc42
IP00319992.1	88	114	122	112	87	523	112	104.6	17.8	16.46%	18.0	16.85%	Alp2b3
IP00420725.4	94	47	60	105	207	513	94	102.6	24.3	36.22%	75.3	60.74%	Hist2h3c2
IP00227235.3	102	151	105	64	79	501	102	100.2	27.5	23.02%	20.7	25.09%	Anep
IP00119138.1	78	95	141	89	93	496	93	99.2	32.6	31.14%	28.9	26.88%	Alp2b4
IP00469221.5	112	117	118	75	60	482	112	96.4	3.2	2.78%	30.1	35.70%	Hspa5
IP00230264.5	77	87	73	121	123	481	87	96.2	7.2	9.13%	28.3	26.79%	Hist1h4a
IP00648683.3	132	104	64	93	86	479	93	95.8	34.2	34.18%	15.1	18.68%	Gnai3
IP00123644.1	0	252	0	215	0	467	0	93.4	145.5	173.21%	124.1	173.21%	Acta1
IP00649886.1	102	153	85	95	32	467	95	93.4	35.4	31.22%	33.9	47.91%	Hba-a2
IP00227432.6	49	160	102	114	39	464	102	92.8	55.5	53.56%	40.3	47.40%	Dnm3
IP00108780.7	87	104	88	87	98	464	88	92.8	9.5	10.26%	6.1	6.68%	Sic12a5

IP00323230.5	76	113	81	86	98	454	86	90.8	20.1	22.31%	8.7	9.89%	Slc25a3
IP00407339.7	102	90	71	91	99	453	91	90.6	15.6	17.83%	14.4	16.58%	Uqrcr2
IP00753468.2	70	49	53	92	184	448	70	89.6	11.2	19.45%	67.3	61.33%	Basp1
IP00124771.1	90	89	82	93	89	443	89	88.6	4.4	5.01%	5.6	6.33%	Ap2a2
IP00407499.1	76	69	78	100	117	440	78	88.0	4.7	6.36%	19.6	19.88%	Aldoa
IP00263013.4	81	110	86	71	86	434	86	86.8	15.5	16.79%	8.7	10.69%	Sdha
IP00856379.1	66	117	66	81	102	432	81	86.4	29.4	35.48%	18.1	21.79%	Syn1
IP00115240.1	85	92	79	82	83	421	83	84.2	6.5	7.62%	2.1	2.56%	Ap1b1
IP00395193.1	78	90	85	86	81	420	85	84.0	6.0	7.15%	2.6	3.15%	Ap2a1
IP00123505.2	94	98	77	67	74	410	77	82.0	11.2	12.44%	5.1	7.06%	Hadha
IP00230108.6	69	77	134	71	57	408	71	81.6	35.4	37.98%	41.0	46.97%	Camk2b
IP00113141.1	62	65	57	100	117	401	65	80.2	4.0	6.59%	30.9	33.86%	Ywhae
IP00129519.3	90	82	89	70	69	400	82	80.0	4.4	5.01%	11.3	14.83%	Rtn1
IP00310091.8	83	106	50	118	40	397	83	79.4	28.1	35.33%	42.4	61.21%	Plp1
IP00109275.1	68	85	87	61	91	392	85	78.4	10.4	13.05%	16.3	20.45%	Slc25a22
IP00230707.6	78	59	76	86	85	384	78	76.8	10.4	14.70%	5.5	6.69%	Ank2
IP00312527.4	79	79	61	108	57	384	79	76.8	10.4	14.24%	28.4	37.65%	Mbp
IP00347110.2	103	71	49	63	97	383	71	76.6	27.2	36.53%	24.7	35.43%	Pdhb
IP00110658.1	82	50	32	115	99	378	82	75.6	25.3	46.33%	44.0	53.70%	Ywhag
IP00342603.7	84	71	60	38	118	371	71	74.2	12.0	16.76%	41.3	57.40%	Oxct1
IP00265299.3	0	0	112	258	0	370	0	74.0	64.7	173.21%	129.4	104.90%	H2afj
IP00114209.1	0	0	113	256	0	369	0	73.8	65.2	173.21%	128.3	104.30%	Hist1h2af
IP00622235.5	80	65	49	68	107	369	68	73.8	15.5	23.97%	29.6	39.60%	Stx1b2
IP00114641.2	59	83	76	60	90	368	76	73.6	12.3	16.98%	15.0	19.93%	Slc3a2
IP00554845.2	54	100	70	80	64	368	70	73.6	23.4	31.28%	8.1	11.33%	Vcp
IP00118384.1	89	81	63	54	80	367	80	73.4	13.3	17.15%	13.2	20.11%	Gls
IP00515360.8	99	84	106	28	46	363	84	72.6	11.2	11.67%	40.8	68.07%	Spna1
IP00623553.1	65	140	75	62	18	360	65	72.0	40.7	43.63%	29.9	57.82%	Krt73
IP00663736.2	68	129	71	43	45	356	68	71.2	34.4	38.49%	15.6	29.47%	Maoa
IP00828664.1	82	70	68	70	61	351	70	70.2	7.6	10.33%	4.7	7.12%	Immt
IP00130460.1	67	68	55	57	97	344	67	68.8	7.2	11.42%	23.7	34.01%	Atp6v0d1
IP00229542.1	39	90	77	112	25	343	77	68.6	26.5	38.59%	43.8	61.37%	Hist1h2aa
IP00132042.1	75	77	72	42	77	343	75	68.6	2.5	3.37%	18.9	29.73%	ltp1
IP00122548.3	64	99	64	58	54	339	64	67.8	20.2	26.71%	5.0	8.58%	Atp5h



IP100626790.3	62	90	54	68	65	339	65	67.8	18.9	27.53%	7.4	11.83%	Glul
IP00125899.1	64	99	64	58	54	339	64	67.8	20.2	26.71%	5.0	8.58%	LOC100039281
IP00126635.1	76	90	68	56	48	338	68	67.6	11.1	14.28%	10.1	17.56%	Ndufs2
IP00378480.1	63	0	65	105	104	337	65	67.4	37.0	86.63%	22.8	24.98%	Hist1h4a
IP00337893.2	63	85	71	60	57	336	63	67.2	11.1	15.25%	7.4	11.76%	Ogdhl
IP00229080.7	52	66	63	54	97	332	63	66.4	7.4	12.22%	22.7	31.79%	Pdha1
IP00230507.5	50	91	62	60	63	326	62	65.2	21.1	31.15%	1.5	2.48%	Gpi1
IP00123390.8	71	72	77	48	57	325	71	65.0	3.2	4.38%	14.8	24.47%	Pdia3
IP00153660.4	78	48	63	68	65	322	65	64.4	15.0	23.81%	2.5	3.85%	Hsp90ab1
IP00113149.1	87	37	68	67	62	321	67	64.2	25.2	39.44%	3.2	4.90%	Hsp90aa1
IP0011218.1	67	66	53	56	76	318	66	63.6	7.8	12.60%	12.5	20.28%	Dlat
IP00785343.2	62	35	39	44	138	318	44	63.6	14.6	32.14%	55.8	75.71%	Sv2a
IP00131176.1	49	70	49	77	70	315	70	63.0	12.1	21.65%	14.6	22.30%	Pgk1
IP00128023.3	64	51	68	71	60	314	64	62.8	8.9	14.57%	5.7	8.57%	Aldoc
IP00229796.3	55	75	47	50	86	313	55	62.6	14.4	24.44%	21.7	35.58%	ldh3b
IP00118986.1	82	73	55	57	45	312	57	62.4	13.7	19.64%	6.4	12.28%	Acs16
IP00555069.3	44	80	39	78	70	311	70	62.2	22.4	41.17%	20.6	33.05%	mt-Co2
IP00162781.2	76	68	58	57	50	309	58	61.8	9.0	13.39%	4.4	7.93%	Ctnnb1
IP00330804.4	63	47	55	66	72	303	63	60.6	8.0	14.55%	8.6	13.40%	Crmp1
IP00828253.1	70	54	46	54	77	301	54	60.2	12.2	21.57%	16.1	27.28%	Canx
IP00129622.3	60	65	52	64	57	298	60	59.6	6.6	11.11%	6.0	10.45%	Tln1
IP00227126.1	71	59	44	49	75	298	59	59.6	13.5	23.32%	16.6	29.72%	Vdac3
IP00420570.1	55	73	57	63	49	297	57	59.4	9.9	16.00%	7.0	12.47%	Syt2
IP00119063.2	58	55	53	40	87	293	55	58.6	2.5	4.55%	24.3	40.45%	Opa1
IP00663180.4	59	94	36	40	63	292	59	58.4	29.2	46.36%	14.6	31.45%	Cyc1
IP00120761.3	49	87	62	44	48	290	49	58.0	19.3	29.26%	9.5	18.41%	Sept7
IP00129526.1	71	50	54	66	48	289	54	57.8	11.2	19.12%	9.2	16.37%	Atp5o
IP00113223.2	61	47	38	111	32	289	47	57.8	11.6	23.82%	44.0	72.90%	Hspg2
IP00553798.2	49	85	57	49	48	288	49	57.6	18.9	29.69%	4.9	9.61%	Calr
IP00649778.1	56	37	45	74	76	288	56	57.6	9.5	20.74%	17.3	26.69%	Nfasc
IP00757312.1	79	52	50	40	64	285	52	57.0	16.2	26.85%	12.1	23.48%	Aqp4
IP00230682.6	63	30	29	47	116	285	47	57.0	19.3	47.58%	45.9	71.76%	Fh1
IP00465810.3	64	69	72	35	43	283	64	56.6	4.0	5.91%	19.5	38.94%	Ank1
IP00854028.2	53	73	70	34	53	283	53	56.6	10.8	16.51%	18.0	34.41%	Ly6c1



IP100122565.1	36	55	38	79	33	241	38	48.2	10.4	24.28%	25.2	50.48%	Gpm6a
IP100132728.2	46	25	60	47	62	240	47	48.0	17.6	40.34%	8.1	14.46%	Atp4a
IP100464296.5	53	42	52	47	46	240	47	48.0	6.1	12.41%	3.2	6.65%	Rtn4
IP100553538.3	46	55	38	47	51	237	47	47.4	8.5	18.36%	6.7	14.69%	Epb4.113
IP100230754.5	32	46	47	50	61	236	47	47.2	8.4	20.13%	7.4	14.00%	Atp6v1f
IP100133240.1	61	19	25	45	85	235	45	47.0	22.7	64.90%	30.6	59.13%	H3f3a
IP100315999.4	38	38	29	47	83	235	38	47.0	5.2	14.85%	27.5	51.88%	Snap91
IP100119115.2	59	47	47	43	38	234	47	46.8	6.9	13.58%	4.5	10.57%	Myo5a
IP100222833.1	54	51	37	56	36	234	51	46.8	9.1	19.17%	11.3	26.21%	Pygb
IP100467833.5	39	39	32	50	73	233	39	46.6	4.0	11.02%	20.6	39.78%	Syn2
IP100880852.1	43	42	37	63	48	233	43	46.6	3.2	7.90%	13.1	26.46%	Ywhaq
IP100129134.3	38	56	53	56	29	232	53	46.4	9.6	19.68%	14.8	32.17%	Cntnap1
IP100119458.7	33	35	26	63	75	232	35	46.4	4.7	15.08%	25.5	46.72%	Rab3a
IP100123704.1	50	45	45	53	38	231	45	46.2	2.9	6.19%	7.5	16.56%	Ank3
IP100132474.3	42	45	42	33	68	230	42	46.0	1.7	4.03%	18.2	38.13%	Phb
IP100308976.1	38	52	38	39	62	229	39	45.8	8.1	18.94%	13.6	29.30%	Actn4
IP100114560.5	40	44	92	34	18	228	40	45.6	28.9	49.32%	38.9	81.12%	Rps27a
IP100128450.1	33	64	34	55	42	228	42	45.6	17.6	40.34%	10.6	24.27%	Sirt2
IP100407954.2	40	44	92	34	18	228	40	45.6	28.9	49.32%	38.9	81.12%	Uba52
IP100556723.1	27	45	92	32	29	225	32	45.0	33.6	61.39%	35.5	69.68%	Camk2g
IP100353420.7	41	41	33	44	64	223	41	44.6	4.6	12.05%	15.7	33.44%	Ppia
IP100468121.1	47	30	30	50	64	221	47	44.2	9.8	27.52%	17.1	35.60%	Hist1h3g
IP100330754.1	35	46	29	36	74	220	36	44.0	8.6	23.51%	24.2	52.26%	Rab1
IP100132799.4	37	55	33	41	53	219	41	43.8	11.7	28.13%	10.1	23.78%	Mdh1
IP100554989.3	50	39	36	40	54	219	40	43.8	7.4	17.69%	9.5	21.81%	Pcx
IP100229510.5	31	81	25	40	42	219	40	43.8	30.7	67.33%	9.3	26.05%	Rala
IP100124700.1	45	72	37	29	36	219	37	43.8	18.3	35.73%	4.4	12.82%	Stxn3
IP100307837.6	35	38	82	32	31	218	35	43.6	26.3	50.93%	29.2	60.33%	Dlst
IP100875717.2	44	40	33	38	63	218	40	43.6	5.6	14.28%	16.1	35.98%	Sucl2
IP100224626.3	66	36	52	49	11	214	49	42.8	15.0	29.24%	22.9	61.22%	Ahnak
IP100115875.7	35	48	40	38	52	213	40	42.6	6.6	15.99%	7.6	17.47%	Ctnna1
IP100113517.1	57	49	30	30	47	213	47	42.6	13.9	30.59%	9.8	27.52%	Syngap1
IP100850200.1	44	51	47	37	34	213	44	42.6	3.5	7.42%	6.8	17.31%	Tln2
IP100133440.1	42	37	41	41	51	212	41	42.4	2.6	6.61%	5.8	13.02%	Dpp6

IP100121309.2	34	50	36	46	44	210	44	42.0	8.7	21.79%	5.3	12.60%	Ndufs3
IP100853924.1	43	45	44	42	35	209	43	41.8	1.0	2.27%	4.7	11.72%	Anxa6
IP100317309.5	41	73	48	20	27	209	41	41.8	16.8	31.15%	14.6	46.02%	Hspa12b
IP100380273.1	48	52	43	31	35	209	43	41.8	4.5	9.46%	6.1	16.82%	Mbc2
IP100311515.2	44	40	37	36	52	209	40	41.8	3.5	8.71%	9.0	21.51%	Me3
IP100380436.1	33	27	43	53	52	208	43	41.6	8.1	23.54%	5.5	11.16%	Atp6v1e1
IP100118120.1	36	42	46	36	48	208	42	41.6	5.0	12.18%	6.4	14.84%	Dlg4
IP100228807.3	89	0	0	0	119	208	0	41.6	51.4	173.21%	68.7	173.21%	Hist2h2ac
IP100336324.11	49	28	29	62	40	208	40	41.6	11.8	33.53%	16.8	38.48%	Ldhb
IP100880742.1	47	34	41	45	40	207	41	41.4	6.5	16.00%	2.6	6.30%	Pl4ka
IP100406790.6	46	64	51	45	0	206	46	41.2	9.3	17.31%	27.9	87.11%	Alp2a1
IP100466699.3	53	38	41	47	27	206	41	41.2	7.9	18.04%	10.3	26.77%	Macf1
IP100850457.1	55	40	41	39	30	205	40	41.0	8.4	18.50%	5.9	15.98%	Dmxl2
IP100755916.3	47	32	35	52	39	205	39	41.0	7.9	20.89%	8.9	21.16%	Fasn
IP100118899.1	47	44	39	41	34	205	41	41.0	4.0	9.33%	3.6	9.49%	Gria2
IP100124282.1	40	36	36	53	39	204	39	40.8	2.3	6.19%	9.1	21.27%	Cadm2
IP100663627.1	43	43	47	36	34	203	43	40.6	2.3	5.21%	7.0	17.95%	Cadps
IP100261627.1	39	41	38	39	45	202	39	40.4	1.5	3.88%	3.8	9.31%	Gdi2
IP100133706.1	54	40	37	36	35	202	37	40.4	9.1	20.78%	1.0	2.78%	Rap1a
IP100137227.1	37	39	47	42	37	202	39	40.4	5.3	12.91%	5.0	11.90%	Slc4a10
IP100405986.2	77	31	27	27	40	202	31	40.4	27.8	61.74%	7.5	23.95%	Sod2
IP100310220.3	33	67	23	40	38	201	38	40.2	23.1	56.26%	9.3	27.60%	Ralb
IP100110265.1	43	23	38	53	41	198	41	39.6	10.4	30.02%	7.9	18.04%	C1qbp
IP100228618.5	50	35	36	42	34	197	36	39.4	8.4	20.79%	4.2	11.15%	Ndufa10
IP100121378.2	37	42	25	30	62	196	37	39.2	8.7	25.20%	20.1	51.47%	Arf1
IP100403993.5	56	37	36	36	31	196	36	39.2	11.3	26.21%	2.9	8.41%	Rap1b
IP100396743.1	51	42	33	31	38	195	38	39.0	9.0	21.43%	3.6	10.60%	Gnb4
IP100136134.1	47	50	30	32	36	195	36	39.0	10.8	25.48%	3.1	9.35%	Tom70a
IP100169788.1	39	24	23	85	21	192	24	38.4	9.0	31.27%	36.4	84.62%	Lamc1
IP100402778.1	47	55	34	28	28	192	34	38.4	10.6	23.38%	3.5	11.55%	Rpn2
IP100469548.2	39	39	38	39	37	192	39	38.4	0.6	1.49%	1.0	2.63%	Ube1y1
IP100649135.2	46	36	50	35	24	191	36	38.2	7.2	16.39%	13.1	35.92%	Acadvl
IP100400180.1	35	44	41	39	31	190	39	38.0	4.6	11.46%	5.3	14.30%	Sept8
IP100138892.2	27	47	39	46	30	189	39	37.8	10.1	26.73%	8.0	20.92%	Amph

IP100330163.4	32	38	29	47	43	189	38	37.8	4.6	13.89%	9.5	23.83%	Napb
IP100331332.6	27	65	36	29	32	189	32	37.8	19.9	46.54%	3.5	10.86%	Ndufv2
IP100625588.1	45	50	37	29	27	188	37	37.6	6.6	14.90%	5.3	17.07%	Bdh1
IP100762897.2	49	47	37	30	25	188	37	37.6	6.4	14.50%	6.0	19.66%	Flnb
IP100405227.3	53	33	31	36	34	187	34	37.4	12.2	31.19%	2.5	7.48%	Gna11
IP100122409.1	37	36	37	37	38	185	37	37.0	0.6	1.57%	0.6	1.55%	Gnaq
IP100109169.1	35	48	35	36	30	184	35	36.8	7.5	19.08%	3.2	9.55%	Cand1
IP100227898.3	34	57	33	28	32	184	33	36.8	13.6	32.85%	2.6	8.53%	Prkcc
IP100112963.1	30	34	40	33	46	183	34	36.6	5.0	14.52%	6.5	16.40%	ENSMUSG0000074081
IP100123313.1	27	54	38	39	24	182	38	36.4	13.6	34.23%	8.4	24.91%	Lsmp
IP100676243.2	37	37	33	30	45	182	37	36.4	2.3	6.47%	7.9	22.05%	Rab1b
IP100126042.3	50	47	45	23	17	182	45	36.4	2.5	5.32%	14.7	52.03%	Utrn
IP100752501.2	33	32	21	38	56	180	33	36.0	6.7	23.23%	17.5	45.66%	Ndrp2
IP100122974.1	50	38	38	43	11	180	38	36.0	6.9	16.50%	17.2	56.13%	Pripf8
IP100331541.5	47	32	31	33	36	179	33	35.8	9.0	24.44%	2.5	7.55%	Gnas
IP100605677.5	34	37	40	43	25	179	37	35.8	3.0	8.11%	9.6	26.79%	H2afz
IP100469392.2	37	35	47	46	14	179	37	35.8	6.4	16.21%	18.8	52.63%	Syne1
IP100331734.5	44	24	30	36	44	178	36	35.6	10.3	31.42%	7.0	19.16%	Idh3g
IP100122965.1	30	46	34	39	28	177	34	35.4	8.3	22.71%	5.5	16.36%	Cox4i1
IP100113480.1	45	43	26	19	44	177	43	35.4	10.4	27.47%	12.9	43.47%	Rab2
IP100138406.1	24	46	36	42	28	176	36	35.2	11.0	31.17%	7.0	19.88%	Gja1
IP100120984.5	36	34	33	33	39	175	34	35.0	1.5	4.45%	3.5	9.90%	Prdx6
IP100462803.5	81	20	0	46	28	175	28	35.0	42.2	125.33%	23.2	93.97%	Prss3
IP100119667.1	31	45	41	30	27	174	31	34.8	7.2	18.49%	7.4	22.56%	Alcam
IP100134809.2	30	42	37	34	31	174	34	34.8	6.0	16.59%	3.0	8.82%	Ndufa8
IP100117978.1	52	24	34	29	35	174	34	34.8	14.2	38.70%	3.2	9.84%	Rpn1
IP100126115.1	34	34	38	31	36	173	34	34.6	2.3	6.54%	3.6	10.30%	Pfkfb
IP100416906.2	30	10	16	32	84	172	30	34.4	10.3	54.98%	35.6	80.80%	Hist2h3c2
IP100751137.1	35	32	31	34	40	172	34	34.4	2.1	6.37%	4.6	13.09%	Stxn5
IP100420562.5	31	33	30	41	34	169	33	33.8	1.5	4.88%	5.6	15.91%	Prkcb1
IP100263863.8	36	43	0	27	62	168	36	33.6	23.1	87.62%	31.1	104.78%	Arf3
IP100620873.3	30	36	29	28	45	168	30	33.6	3.8	11.96%	9.5	28.06%	Rhob
IP100470981.2	28	37	25	25	52	167	28	33.4	6.2	20.82%	15.6	45.85%	Acaa2
IP100405625.9	36	36	41	31	23	167	36	33.4	2.9	7.66%	9.0	28.48%	Car2

IP100309035.2	29	42	27	33	35	166	33	33.2	8.1	24.93%	4.2	13.15%	Eef1a1
IP00121387.1	33	40	42	34	17	166	34	33.2	4.7	12.33%	12.8	41.18%	Enpep
IP00111770.7	34	40	33	29	30	166	33	33.2	3.8	10.61%	2.1	6.79%	Ndufb10
IP00758024.1	41	25	27	33	39	165	33	33.0	8.7	28.12%	6.0	18.18%	Pgam1
IP00330862.5	28	40	29	30	37	164	30	32.8	6.7	20.59%	4.4	13.62%	Ncdn
IP00221614.3	33	38	36	41	15	163	36	32.6	2.5	7.06%	13.8	44.99%	Gstm1
IP00308691.1	26	42	34	30	30	162	30	32.4	8.0	23.53%	2.3	7.37%	Cap1
IP00124792.1	38	34	27	36	26	161	34	32.2	5.6	16.87%	5.5	18.56%	Mtap1a
IP00221613.5	26	29	29	40	37	161	29	32.2	1.7	6.19%	5.7	16.09%	Ppp3ca
IP00109109.1	27	40	34	30	30	161	30	32.2	6.5	19.33%	2.3	7.37%	Tufm
IP00115627.4	22	37	27	27	47	160	27	32.0	7.6	26.64%	11.5	34.30%	Dnajc5
IP00474883.2	31	40	30	25	34	160	31	32.0	5.5	16.36%	4.5	15.20%	Slc2a1
IP00115679.1	36	48	30	29	16	159	30	31.8	9.2	24.12%	7.8	31.24%	Atp5k
IP00828762.1	13	11	10	19	105	158	13	31.6	1.5	13.48%	52.4	117.41%	Gap43
IP00331299.9	35	35	29	37	22	158	35	31.6	3.5	10.50%	7.5	25.59%	Vcl
IP00457898.3	27	36	26	32	36	157	32	31.4	5.5	18.56%	5.0	16.06%	Rhoa
IP00122815.3	24	48	30	25	29	156	29	31.2	12.5	36.74%	2.6	9.45%	Acly
IP00133006.1	35	37	30	27	27	156	30	31.2	3.6	10.60%	1.7	6.19%	Hpcal4
IP00877205.1	33	26	23	28	46	156	28	31.2	5.1	18.77%	12.1	37.41%	Prdx3
IP00135869.3	32	35	34	24	31	156	32	31.2	1.5	4.54%	5.1	17.30%	Rab11b
IP00121288.5	26	33	29	32	34	154	32	30.8	3.5	11.97%	2.5	7.95%	L1cam
IP00131138.10	31	29	48	27	18	153	29	30.6	10.4	29.00%	15.4	49.66%	Hnt
IP00170310.1	52	7	10	25	59	153	25	30.6	25.2	109.39%	25.1	80.13%	mt-Co1
IP00116748.1	30	27	39	35	22	153	30	30.6	6.2	19.52%	8.9	27.78%	Mtap1b
IP00134585.1	26	29	20	44	34	153	29	30.6	4.6	18.33%	12.1	36.90%	Pacsin1
IP00115762.1	30	27	31	33	30	151	30	30.2	2.1	7.10%	1.5	4.88%	Eef2
IP00648938.1	26	26	27	60	12	151	26	30.2	0.6	2.19%	24.6	74.41%	Lamb2
IP00869430.1	24	35	30	36	25	150	30	30.0	5.5	18.56%	5.5	18.16%	Cox6b1
IP00221602.1	23	38	23	33	33	150	33	30.0	8.7	30.93%	5.8	19.46%	Dnm1l
IP00348094.4	35	32	33	24	25	149	32	29.8	1.5	4.58%	4.9	18.05%	Cyfp2
IP00116753.4	28	36	30	24	31	149	30	29.8	4.2	13.29%	3.8	13.36%	Etfa
IP00118825.2	29	20	19	39	42	149	29	29.8	5.5	24.30%	12.5	37.51%	Lrnmb1
IP00475154.1	35	19	24	33	38	149	33	29.8	8.2	31.48%	7.1	22.40%	Ndufa5
IP00130920.1	27	28	28	35	29	147	28	29.4	0.6	2.09%	3.8	12.35%	Actr3

IP100133342.1	34	34	33	24	22	147	33	29.4	0.6	1.71%	5.9	22.25%	Cacna2d1
IP100831373.1	31	38	31	25	22	147	31	29.4	4.0	12.12%	4.6	17.63%	Ctsb
IP100225275.5	40	40	27	15	25	147	27	29.4	7.5	21.04%	6.4	28.79%	Iiga1
IP100407543.2	32	39	30	22	24	147	30	29.4	4.7	14.04%	4.2	16.43%	Iiga6
IP100110588.4	35	34	36	26	16	147	34	29.4	1.0	2.86%	10.0	38.46%	P140
IP100124221.1	25	43	18	25	36	147	25	29.4	12.9	44.99%	9.1	34.46%	Rab14
IP100466371.2	26	25	24	45	26	146	26	29.2	1.0	4.00%	11.6	36.60%	Kpnb1
IP100466069.3	38	22	23	30	33	146	30	29.2	9.0	32.40%	5.1	17.90%	P4hb
IP100338309.4	37	30	29	28	21	145	29	29.0	4.4	13.62%	4.4	16.76%	Ryr2
IP100135677.1	39	38	37	17	13	144	37	28.8	1.0	2.63%	12.9	57.57%	Tfrc
IP100121545.1	15	46	28	31	23	143	28	28.6	15.6	52.47%	4.0	14.79%	Hspe1
IP100775779.11	33	29	26	27	28	143	28	28.6	3.5	11.97%	1.0	3.70%	Sirpa
IP100420329.1	28	36	28	23	27	142	28	28.4	4.6	15.06%	2.6	10.18%	Anxa5
IP100116192.1	25	33	27	30	27	142	27	28.4	4.2	14.69%	1.7	6.19%	Bin1
IP100124695.1	21	20	41	29	31	142	29	28.4	11.8	43.34%	6.4	19.10%	Got1
IP100274407.1	27	31	29	30	25	142	29	28.4	2.0	6.90%	2.6	9.45%	Mtap2
IP100453798.3	28	33	18	50	12	141	28	28.2	7.6	29.00%	20.4	76.61%	Agrn
IP100119203.4	20	19	36	41	25	141	25	28.2	9.5	38.16%	8.2	24.07%	Citb
IP100224768	39	0	0	41	61	141	39	28.2	22.5	173.21%	31.1	91.46%	Csl
IP100221608.3	21	36	37	18	29	141	29	28.2	9.0	28.60%	9.5	34.07%	Letm1
IP100133215.3	24	38	28	24	26	140	26	28.0	7.2	24.04%	2.0	7.69%	Atp1b3
IP100653158.1	36	32	28	28	16	140	28	28.0	4.0	12.50%	6.9	28.87%	Flna
IP100874522.1	26	29	21	39	25	140	26	28.0	4.0	15.95%	9.5	33.36%	Mtap6
IP100315100.1	24	27	22	36	31	140	27	28.0	2.5	10.34%	7.1	23.91%	Nptn
IP100313884.1	30	27	22	21	40	140	27	28.0	4.0	15.35%	10.7	38.65%	Prkar2b
IP100453834.2	40	27	36	19	18	140	27	28.0	6.7	19.39%	10.1	41.57%	Tjp1
IP100667086.3	0	139	0	0	0	139	0	27.8	80.3	173.21%	0.0	#DIV/0!	4732456N10Rik
IP100131177.1	22	31	20	28	38	139	28	27.8	5.9	24.08%	9.0	31.46%	Epb4.111
IP100230394.5	15	40	30	28	26	139	28	27.8	12.6	44.41%	2.0	7.14%	Ndufab1
IP100111686.5	15	24	15	19	66	139	19	27.8	5.2	28.87%	28.4	85.08%	Ywhah
IP100404182.2	31	23	25	30	29	138	29	27.6	4.2	15.81%	2.6	9.45%	Adam22
IP100225390.5	17	33	24	30	34	138	30	27.6	8.0	32.52%	5.0	17.16%	Capzb
IP100880252.1	23	32	31	41	11	138	31	27.6	4.9	17.21%	15.3	55.21%	Ezr
IP100474959.2	23	33	37	22	23	138	23	27.6	7.2	23.26%	8.4	30.68%	Piprs

IP00608063.3	32	31	37	22	16	138	31	27.6	3.2	9.64%	10.8	43.27%	Ugcg1
IP00134093.4	19	24	36	34	24	137	24	27.4	8.7	33.18%	6.4	20.52%	Hspa2
IP00323122.1	26	20	14	70	7	137	20	27.4	6.0	30.00%	34.5	113.84%	Lama2
IP00387494.1	26	38	27	23	23	137	26	27.4	6.7	21.95%	2.3	9.49%	Sucg1
IP00453499.3	29	38	24	22	23	136	24	27.2	7.1	23.39%	1.0	4.35%	Aldh6a1
IP00122069.1	31	15	19	39	32	136	31	27.2	8.3	38.43%	10.1	33.83%	Eef1a2
IP00877320.1	25	29	33	24	25	136	25	27.2	4.0	13.79%	4.9	18.05%	Lonp1
IP00762713.1	27	39	28	24	18	136	27	27.2	6.7	21.25%	5.0	21.57%	Uqcrb
IP00169925.2	20	25	33	26	31	135	26	27.0	6.6	25.22%	3.6	12.02%	Igsf8
IP00355248.5	27	52	24	32	0	135	27	27.0	15.4	44.78%	16.7	89.21%	Mpo
IP00114352.1	32	27	27	37	12	135	27	27.0	2.9	10.07%	12.6	49.67%	Msn
IP00308063.3	30	34	31	20	20	135	30	27.0	2.1	6.57%	6.4	26.83%	Samm50
IP00132347.1	34	18	25	20	37	134	25	26.8	8.0	31.25%	8.7	31.96%	Cfl1
IP00742334.1	21	36	29	25	23	134	25	26.8	7.5	26.18%	3.1	11.90%	Slc1a6
IP00395042.1	29	31	31	21	21	133	29	26.6	1.2	3.81%	5.8	23.73%	Ganab
IP00761641.1	28	23	30	27	25	133	27	26.6	3.6	13.35%	2.5	9.21%	Rtn3
IP00845544.1	30	30	27	28	18	133	28	26.6	1.7	5.97%	5.5	22.63%	Scn2a1
IP00153202.1	18	31	23	32	29	133	29	26.6	6.6	27.32%	4.6	16.37%	Sept6
IP00649556.1	33	32	35	16	17	133	32	26.6	1.5	4.58%	10.7	47.17%	Serpinh1
IP00109612.2	17	40	28	22	25	132	25	26.4	11.5	40.60%	3.0	12.00%	Rab35
IP00125266.1	28	38	32	15	19	132	28	26.4	5.0	15.41%	8.9	40.40%	Scppdh
IP00229008.2	28	54	22	27	0	131	27	26.2	17.0	49.07%	14.4	87.94%	Epx
IP00230444.1	30	32	28	25	16	131	28	26.2	2.0	6.67%	6.2	27.15%	Kcnj10
IP00139301.3	20	24	17	28	42	131	24	26.2	3.5	17.27%	12.5	43.21%	Kras
IP00221788.3	27	40	34	30	0	131	30	26.2	6.5	19.33%	18.6	87.11%	LOC100041265
IP00626793.3	21	22	22	33	33	131	22	26.2	0.6	2.66%	6.4	21.65%	Sept11
IP00757771.1	19	33	28	26	24	130	26	26.0	7.1	26.60%	2.0	7.69%	Alp8a1
IP00648567.1	27	22	21	23	36	129	23	25.8	3.2	13.78%	8.1	30.54%	Pde2a
IP00130391.1	18	27	27	22	34	128	27	25.6	5.2	21.65%	6.0	21.79%	4631427C17Rik
IP00620256.3	27	34	24	18	25	128	25	25.6	5.1	18.11%	3.8	16.95%	Ctnnd2
IP00461964.2	26	28	36	15	23	128	26	25.6	5.3	17.64%	10.6	42.97%	Por
IP00118569.1	27	44	26	14	17	128	26	25.6	10.1	31.29%	6.2	32.87%	Trf
IP00331628.5	28	30	25	24	20	127	25	25.4	2.5	9.10%	2.6	11.50%	Ptkar2a
IP00153266.1	20	23	16	27	40	126	23	25.2	3.5	17.86%	12.0	43.42%	Hras1



IP100131693.2	31	29	29	17	19	125	29	25.0	1.2	3.89%	6.4	29.67%	Ace2
IP100311461.1	25	23	23	32	19	125	25	25.0	1.5	6.19%	6.7	26.99%	Lmna
IP100130185.1	37	22	21	14	31	125	22	25.0	9.0	33.61%	8.5	38.84%	Rab2b
IP100857911.1	24	28	21	21	30	124	24	24.8	3.5	14.43%	5.2	21.65%	Atp6v1h
IP100113248.1	27	28	23	22	24	124	24	24.8	2.6	10.18%	1.0	4.35%	Podxl
IP100124444.1	23	39	23	15	24	124	23	24.8	9.2	32.60%	4.9	23.87%	Ppp1ca
IP100118075.1	21	18	27	29	28	123	27	24.6	4.6	20.83%	1.0	3.57%	Pclo
IP100126248.3	19	25	31	29	18	122	25	24.4	6.0	24.00%	7.0	26.92%	Dnajc6
IP100221456.1	19	28	17	35	23	122	23	24.4	5.9	27.47%	9.2	36.66%	Tmod2
IP100279443.1	34	33	26	11	17	121	26	24.2	4.4	14.06%	7.5	41.94%	Alpl
IP100321348.3	21	22	25	15	38	121	22	24.2	2.1	9.18%	11.5	44.36%	Atp6v1c1
IP100130489.1	23	14	10	51	23	121	23	24.2	6.7	42.50%	21.0	74.83%	H2afy2
IP100387312.2	23	31	20	21	26	121	23	24.2	5.7	23.05%	3.2	14.39%	Napa
IP100136000.1	22	31	19	34	15	121	22	24.2	6.2	26.02%	10.0	44.19%	Slc6a11
IP100515604.2	17	27	19	29	29	121	27	24.2	5.3	25.20%	5.8	22.49%	Slc8a2
IP100330523.1	20	30	18	21	31	120	21	24.0	6.4	28.36%	6.8	29.17%	Negr1
IP100308446.2	17	22	31	23	27	120	23	24.0	7.1	30.41%	4.0	14.81%	Opcml
IP100118930.1	26	36	36	22	0	120	26	24.0	5.8	17.67%	18.1	93.87%	Prph
IP100123613.1	21	15	28	31	25	120	25	24.0	6.5	30.50%	3.0	10.71%	Sh3gl2
IP100323897.3	15	9	56	17	23	120	17	24.0	25.6	95.92%	21.0	65.63%	Vamp3
IP100857780.1	27	27	19	24	22	119	24	23.8	4.6	18.98%	2.5	11.62%	Ncam2
IP100226215.1	23	22	20	44	10	119	22	23.8	1.5	7.05%	17.5	70.84%	Nid1
IP100230715.5	25	36	22	23	12	118	23	23.6	7.4	26.64%	6.1	32.01%	Cadm3
IP100282848.1	19	21	15	33	30	118	21	23.6	3.1	16.66%	9.6	37.09%	Cd81
IP100308077.6	29	24	17	31	17	118	24	23.6	6.0	25.83%	8.1	37.31%	Lap3
IP100139788.2	24	14	24	36	20	118	24	23.6	5.8	27.94%	8.3	31.22%	Ndufb5
IP100849670.1	24	30	28	22	14	118	24	23.6	3.1	11.18%	7.0	32.92%	Ptprd
IP100131666.1	17	23	23	22	32	117	23	23.4	3.5	16.50%	5.5	21.46%	Ap2m1
IP100453826.2	32	25	30	22	8	117	25	23.4	3.6	12.43%	11.1	55.68%	Hsd17b4
IP100379694.4	29	34	33	12	9	117	29	23.4	2.6	8.27%	13.1	72.65%	Krt5
IP100420706.4	23	28	27	23	16	117	23	23.4	2.6	10.18%	5.6	25.31%	Ndufa13
IP100115833.2	18	26	27	25	20	116	25	23.2	4.9	20.84%	3.6	15.02%	Sv2b
IP100111168.1	115	0	0	0	0	115	0	23.0	66.4	173.21%	0.0	#DIV/0!	Abcb1b
IP100228978.2	22	17	24	13	39	115	22	23.0	3.6	17.17%	13.1	51.52%	Echs1

IP100320217.9	36	18	19	22	20	115	20	23.0	10.1	41.57%	1.5	7.51%	Hadhb
IP100752192.1	21	24	26	16	28	115	24	23.0	2.5	10.63%	6.4	27.55%	Iars2
IP100400016.1	31	16	18	23	27	115	23	23.0	8.1	37.59%	4.5	19.89%	Ndufs4
IP100127556.1	28	22	25	17	23	115	23	23.0	3.0	12.00%	4.2	19.22%	Nrcam
IP100403929.3	25	28	25	15	22	115	25	23.0	1.7	6.66%	5.1	24.83%	Rab11a
IP100330887.1	27	32	28	20	7	114	27	22.8	2.6	9.12%	10.6	57.81%	Ascc311
IP100136135.1	23	19	24	29	19	114	23	22.8	2.6	12.03%	5.0	20.83%	Mpp2
IP100652934.3	32	19	22	22	19	114	22	22.8	6.8	27.97%	1.7	8.25%	Pcyox1
IP100856723.1	20	22	46	17	9	114	20	22.8	14.5	49.32%	19.5	81.12%	Rps27a
IP100119152.1	30	23	18	20	22	113	22	22.6	6.0	25.47%	2.0	10.00%	Gna14
IP100621548.2	21	22	28	25	17	113	22	22.6	3.8	16.00%	5.7	24.37%	Ndufa12
IP100121949.1	27	32	28	8	18	113	27	22.6	2.6	9.12%	10.0	55.56%	Ndufb7
IP100339468.4	19	29	19	28	18	113	19	22.6	5.8	25.85%	5.5	25.42%	Sept3
IP100844763.1	14	36	20	10	31	111	20	22.2	11.4	48.74%	10.5	51.66%	Cntrf
IP100626132.3	23	28	17	22	21	111	22	22.2	5.5	24.30%	2.6	13.23%	Gdap111
IP100123342.4	24	19	20	24	24	111	24	22.2	2.6	12.60%	2.3	10.19%	Pfkip
IP100752108.1	21	25	27	19	19	111	21	22.2	3.1	12.56%	4.6	21.32%	Rab7
IP100403586.3	32	32	27	0	20	111	27	22.2	2.9	9.52%	14.0	89.44%	Tubb1
IP100387557.3	25	17	18	20	30	110	20	22.0	4.4	21.79%	6.4	28.36%	Acad9
IP100125861.2	14	32	19	11	34	110	19	22.0	9.3	42.88%	11.7	54.73%	Arf5
IP100153317.3	23	22	20	29	16	110	22	22.0	1.5	7.05%	6.7	30.73%	Gda
IP100402982.2	25	22	20	22	21	110	22	22.0	2.5	11.27%	1.0	4.76%	Maob
IP100224570.3	31	14	26	27	12	110	26	22.0	8.7	36.92%	8.4	38.71%	Pygm
IP100875866.1	14	21	20	27	27	109	21	21.8	3.8	20.65%	4.0	16.38%	Camkv
IP100132705.1	34	30	16	14	15	109	16	21.8	9.5	35.44%	1.0	6.67%	Cyb5f3
IP100221615.5	28	17	15	18	31	109	18	21.8	7.0	35.00%	8.5	39.87%	Hepacam
IP100138378.4	27	29	25	13	15	109	25	21.8	2.0	7.41%	6.4	36.39%	Hyou1
IP100454142.5	23	41	24	21	0	109	23	21.8	10.1	34.49%	13.1	87.18%	Matr3
IP100624192.3	24	22	17	19	27	109	22	21.8	3.6	17.17%	5.3	25.20%	Prdx5
IP100221754.3	16	42	24	18	9	109	18	21.8	13.3	48.72%	7.5	44.41%	Sic38a3
IP100756238.2	30	18	22	26	12	108	22	21.6	6.1	26.19%	7.2	36.06%	Abhd12
IP100475031.1	27	25	26	21	9	108	25	21.6	1.0	3.85%	8.7	46.80%	Atp5l
IP100754853.1	35	8	21	26	18	108	21	21.6	13.5	63.30%	4.0	18.65%	Sic12a2
IP100109213.1	16	31	24	18	18	107	18	21.4	7.5	31.71%	3.5	17.32%	Bsg

PI00129577.1	20	15	18	50	4	107	18	21.4	2.5	14.24%	23.6	98.25%	Lama5
PI00319518.4	21	23	24	24	15	107	23	21.4	1.5	6.74%	5.2	24.74%	Pfkl
PI00875110.1	20	30	20	16	21	107	20	21.4	5.8	24.74%	2.6	13.93%	Ppp1cb
PI00126191.5	25	39	26	12	5	107	25	21.4	7.8	26.03%	10.7	74.60%	Rcn2
PI00845581.1	29	15	17	24	21	106	21	21.2	7.6	37.24%	3.5	16.99%	Cpe
PI00116356.5	24	16	24	20	22	106	22	21.2	4.6	21.65%	2.0	9.09%	Dpysl4
PI00310533.2	16	29	18	22	21	106	21	21.2	7.0	33.33%	2.1	10.24%	ENSMUSG00000048362
PI00468203.3	20	22	18	35	11	106	20	21.2	2.0	10.00%	12.3	57.85%	Mag
PI00380799.5	19	18	12	21	36	106	19	21.2	3.8	23.18%	12.1	52.71%	Nras
PI00605187.4	18	27	18	20	23	106	20	21.2	5.2	24.74%	2.5	12.38%	Pebp1
PI00226140.5	24	22	24	21	15	106	22	21.2	1.2	4.95%	4.6	22.91%	Rap2b
PI00406442.2	21	21	25	30	9	106	21	21.2	2.3	10.34%	11.0	51.42%	Rdx
PI00408892.2	21	29	20	10	25	105	21	21.0	4.9	21.14%	7.6	41.66%	Actn2
PI00187430.3	26	23	21	22	13	105	22	21.0	2.5	10.79%	4.9	26.43%	Rap2a
PI00266942.6	20	26	15	18	25	104	20	20.8	5.5	27.09%	5.1	26.54%	Add1
PI00122273.1	23	21	20	22	18	104	21	20.8	1.5	7.16%	2.0	10.00%	Hpca
PI00222496.3	22	26	23	14	19	104	22	20.8	2.1	8.80%	4.5	24.16%	Hsd17b10
PI00808203.2	15	29	20	20	20	104	20	20.8	7.1	33.26%	0.0	0.00%	Ndufc2
PI00344004.2	23	23	26	21	11	104	23	20.8	1.7	7.22%	7.6	39.50%	Pcca
PI00315334.5	26	24	25	10	18	103	24	20.6	1.0	4.00%	7.5	42.48%	Lrpprc
PI00330289.3	25	19	17	25	17	103	19	20.6	4.2	20.48%	4.6	23.49%	Pccb
PI00122486.3	22	19	19	23	20	103	20	20.6	1.7	8.66%	2.1	10.07%	Spnb4
PI00761331.2	22	29	18	20	13	102	20	20.4	5.6	24.21%	3.6	21.21%	Abcc4
PI00123756.1	23	18	28	21	12	102	21	20.4	5.0	21.74%	8.0	39.45%	Cct2
PI00115663.2	23	19	23	17	20	102	20	20.4	2.3	10.66%	3.0	15.00%	Dlg1
PI00387430.1	27	24	18	17	16	102	18	20.4	4.6	19.92%	1.0	5.88%	Jup
PI00405736.3	18	23	15	20	26	102	20	20.4	4.0	21.65%	5.5	27.09%	Mif
PI00116279.3	23	20	24	17	18	102	20	20.4	2.1	9.32%	3.8	19.25%	Pdia6
PI00111265.3	25	27	22	13	15	102	22	20.4	2.5	10.20%	4.7	28.35%	Slc9a32
PI00378438.6	26	28	20	17	10	101	20	20.2	4.2	16.88%	5.1	32.75%	Abca1
PI00653398.1	26	16	14	17	28	101	17	20.2	6.4	34.44%	7.4	37.48%	EG620772
PI00119114.2	27	20	11	15	28	101	20	20.2	8.0	41.49%	8.9	49.38%	Kcnab2
PI00130186.4	19	19	21	22	20	101	20	20.2	1.2	5.87%	1.0	4.76%	Nme2
PI00230193.7	22	22	17	23	17	101	22	20.2	2.9	14.20%	3.5	18.23%	Rap1gds1

IP100473693.3	15	29	13	21	23	101	21	20.2	8.7	45.88%	5.3	27.85%	Tkt
IP100230427.5	16	27	19	19	20	101	19	20.2	5.7	27.51%	0.6	2.99%	Vapa
IP100131459.1	25	27	20	16	12	100	20	20.0	3.6	15.02%	4.0	25.00%	Acsbg1
IP100313998.1	29	21	20	12	18	100	20	20.0	4.9	21.14%	4.2	24.98%	Ctnnd1
IP100119219.2	20	25	25	13	17	100	20	20.0	2.9	12.37%	6.1	33.33%	Hibch
IP100222419.5	22	28	22	16	12	100	22	20.0	3.5	14.43%	5.0	30.20%	Pkpa4
IP100132002.1	24	21	23	18	13	99	21	19.8	1.5	6.74%	5.0	27.78%	Dctn1
IP100133284.1	18	26	16	15	24	99	18	19.8	5.3	26.46%	4.9	26.91%	Hibadh
IP100453996.1	22	18	22	21	16	99	21	19.8	2.3	11.17%	3.2	16.35%	ldh2
IP100124900.1	18	15	19	14	33	99	18	19.8	2.1	12.01%	9.8	44.77%	Thy1
IP100131896.1	16	22	15	18	28	99	18	19.8	3.8	21.43%	6.8	33.48%	Uchl1
IP100880610.1	23	27	18	18	12	98	18	19.6	4.5	19.89%	3.5	21.65%	Gna13
IP100115607.3	18	18	13	30	19	98	18	19.6	2.9	17.67%	8.6	41.72%	Hist1h1d
IP100119970.4	26	19	13	13	27	98	19	19.6	6.5	33.65%	8.1	45.75%	Mgst3
IP100132531.1	17	18	27	23	13	98	18	19.6	5.5	26.65%	7.2	34.34%	Ppp3cb
IP100420554.2	23	13	15	31	16	98	16	19.6	5.3	31.13%	9.0	43.37%	Prdx2
IP100117181.1	16	25	23	16	18	98	18	19.6	4.7	22.15%	3.6	18.98%	Sdhb
IP100751369.1	15	24	19	23	16	97	19	19.4	4.5	23.32%	3.5	18.16%	Brp44
IP100653686.1	25	25	17	17	13	97	17	19.4	4.6	20.68%	2.3	14.74%	Cyfp1
IP100132443.3	26	21	27	12	11	97	21	19.4	3.2	13.03%	9.0	53.78%	Dag1
IP100759881.1	12	8	9	19	49	97	12	19.4	2.1	21.53%	20.8	81.10%	Dynll2
IP100454049.4	18	17	15	26	21	97	18	19.4	1.5	9.17%	5.5	26.65%	Epb4.1l2
IP100753326.1	29	23	24	9	12	97	23	19.4	3.2	12.69%	7.9	52.92%	Fer1l3
IP100405742.6	23	19	19	24	12	97	19	19.4	2.3	11.36%	6.0	32.88%	mt-Nd5
IP100311873.5	19	23	13	22	19	96	19	19.2	5.0	27.45%	4.6	25.46%	Dpp10
IP100126083.3	20	19	18	23	16	96	19	19.2	1.0	5.26%	3.6	18.98%	Ehd1
IP100353563.4	24	17	18	21	16	96	18	19.2	3.8	19.25%	2.5	13.73%	Ehd3
IP100165731.3	18	15	20	21	22	96	20	19.2	2.5	14.24%	1.0	4.76%	Ldha
IP100154047.1	17	17	19	11	32	96	17	19.2	1.2	6.54%	10.6	51.28%	LOC672195
IP100342766.8	28	22	18	15	13	96	18	19.2	5.0	22.21%	2.5	16.41%	Nomo1
IP100480233.1	24	18	19	17	18	96	18	19.2	3.2	15.81%	1.0	5.56%	Pixnb1
IP100135324.2	20	23	20	20	13	96	20	19.2	1.7	8.25%	4.0	22.88%	Pixnb2
IP100229475.1	21	22	16	17	19	95	19	19.0	3.2	16.35%	1.5	8.81%	C130026L21Rik
IP100460063.2	23	15	14	21	22	95	21	19.0	4.9	28.46%	4.4	22.94%	Cask

IP100315359.1	19	16	24	19	17	95	19	19.0	4.0	20.55%	3.6	18.03%	Cct5
IP100330476.3	19	21	15	16	24	95	19	19.0	3.1	16.66%	4.9	26.91%	Fscn1
IP100111793.1	21	16	21	21	16	95	21	19.0	2.9	14.93%	2.9	14.93%	Mpp6
IP100170093.3	23	21	16	17	18	95	18	19.0	3.6	18.03%	1.0	5.88%	Ndufs8
IP100276515.4	20	17	18	21	19	95	19	19.0	1.5	8.33%	1.5	7.90%	Rab10
IP100133980.1	16	18	12	36	12	94	16	18.8	3.1	19.92%	13.9	69.28%	Ahsg
IP100468539.3	24	20	17	19	14	94	19	18.8	3.5	17.27%	2.5	15.10%	Anxa2
IP100750256.2	9	15	39	18	13	94	15	18.8	15.9	75.59%	13.8	59.13%	Atp5d
IP100116222.1	22	16	16	23	17	94	17	18.8	3.5	19.25%	3.8	20.28%	Cntnap2
IP100319973.3	22	22	17	19	14	94	19	18.8	2.9	14.20%	2.5	15.10%	Flot1
IP100756745.5	20	14	19	23	18	94	19	18.8	3.2	18.20%	2.6	13.23%	Gprin1
IP100128249.1	19	21	15	33	6	94	19	18.8	3.1	16.66%	13.7	76.38%	Mog
IP100453777.2	14	34	13	14	19	94	14	18.8	11.8	58.26%	3.2	20.96%	Uqcrh
IP100109221.3	20	20	19	21	13	93	20	18.6	0.6	2.94%	4.2	23.57%	Auh
IP100396687.3	25	10	13	18	27	93	18	18.6	7.9	49.61%	7.1	36.70%	Cadm1
IP100133387.1	30	21	27	13	2	93	21	18.6	4.6	17.63%	12.5	89.50%	Dysf
IP100227928.3	20	36	0	15	22	93	20	18.6	18.0	96.63%	11.2	91.13%	Ppp1cc
IP100340165.5	20	13	9	20	31	93	20	18.6	5.6	39.77%	11.0	55.00%	Snca
IP100131614.1	26	19	17	14	17	93	17	18.6	4.7	22.87%	1.7	10.83%	Sntb1
IP100785314.1	16	22	18	22	14	92	18	18.4	3.1	16.37%	4.0	22.22%	Flot2
IP100470152.2	14	24	14	21	19	92	19	18.4	5.8	33.31%	3.6	20.03%	Icam5
IP100113726.3	22	25	25	15	5	92	22	18.4	1.7	7.22%	10.0	66.67%	Myh14
IP100858116.1	20	16	18	19	19	92	19	18.4	2.0	11.11%	0.6	3.09%	Nrxn3
IP100137409.3	17	24	21	20	9	91	20	18.2	3.5	16.99%	6.7	39.95%	Aadacl1
IP100346834.1	19	27	22	12	11	91	19	18.2	4.0	17.83%	6.1	40.55%	Aifm1
IP100227773.1	18	19	23	19	12	91	19	18.2	2.6	13.23%	5.6	30.93%	Gstp1
IP100471176.2	21	13	17	23	17	91	17	18.2	4.0	23.53%	3.5	18.23%	Marcks
IP100761443.1	10	30	16	20	15	91	16	18.2	10.3	54.98%	2.6	15.56%	Scamp5
IP100170357.5	21	24	17	19	10	91	19	18.2	3.5	16.99%	4.7	30.82%	Spg3a
IP100469987.3	16	13	23	18	21	91	18	18.2	5.1	29.61%	2.5	12.18%	Tppp
IP100653794.2	21	16	21	10	22	90	21	18.0	2.9	14.93%	6.7	37.69%	Cp
IP100317710.1	20	19	19	16	16	90	19	18.0	0.6	2.99%	1.7	10.19%	Lanc12
IP100396701.3	22	16	21	22	9	90	21	18.0	3.2	16.35%	7.2	41.74%	Myh11
IP100555023.2	21	14	21	18	16	90	18	18.0	4.0	21.65%	2.5	13.73%	Pgrmc1

IP00331597.6	10	24	14	17	25	17	18.0	7.2	45.07%	5.7	30.46%	Rab3c
IP00110825.1	14	18	19	25	13	18	17.8	2.6	15.56%	6.0	31.58%	Baiap2
IP00653921.2	22	14	21	20	12	20	17.8	4.4	22.94%	4.9	27.92%	Dhx9
IP00112129.1	23	20	15	8	23	20	17.8	4.0	20.90%	7.5	48.95%	Dpysl5
IP00331076.4	0	0	0	0	89	0	17.8	0.0	#DIV/0!	51.4	173.21%	EG433923
IP00408207.2	18	20	20	17	14	18	17.8	1.2	5.97%	3.0	17.65%	Gnaz
IP00120374.1	25	17	8	16	23	17	17.8	8.5	51.03%	7.5	47.91%	Nckap1
IP00128155.2	21	18	20	15	15	18	17.8	1.5	7.77%	2.9	17.32%	Nrxn1
IP00125267.4	17	22	11	21	18	18	17.8	5.5	33.05%	5.1	30.79%	Prnd
IP00856469.1	15	26	19	19	10	19	17.8	5.6	27.84%	5.2	32.48%	Rac1
IP00346073.1	17	29	18	13	11	17	17.6	6.7	31.21%	3.6	25.75%	Nnt
IP00223714.5	20	22	20	15	11	20	17.6	1.2	5.59%	4.5	29.41%	Sacm1l
IP00876408.1	21	11	20	20	16	20	17.6	5.5	31.77%	2.3	12.37%	Slc25a18
IP00228680.2	19	16	18	20	15	18	17.6	1.5	8.65%	2.5	14.24%	Tns1
IP00230310.5	14	15	20	21	17	17	17.4	3.2	19.68%	2.1	10.77%	Aak1
IP00816946.1	22	24	20	14	7	20	17.4	2.0	9.09%	6.5	47.61%	Asah1
IP00409405.2	19	15	13	24	16	16	17.4	3.1	19.50%	5.7	32.19%	Hist1h1e
IP00408215.1	17	21	16	21	12	17	17.4	2.6	14.70%	4.5	27.61%	Hspa4l
IP00331692.1	13	31	18	14	11	14	17.4	9.3	44.96%	3.5	24.50%	Ndufb4
IP00515151.2	14	22	14	17	20	17	17.4	4.6	27.71%	3.0	17.65%	Psap
IP00130118.1	19	16	16	18	18	18	17.4	1.7	10.19%	1.2	6.66%	Rab5c
IP00755241.1	15	23	21	16	12	16	17.4	4.2	21.17%	4.5	27.61%	Slc1a4
IP00123276.1	24	17	17	15	13	17	17.2	4.0	20.90%	2.0	13.33%	Afg3l2
IP00877299.1	15	16	20	17	18	17	17.2	2.6	15.56%	1.5	8.33%	Nme1
IP00124046.1	19	10	13	24	20	19	17.2	4.6	32.73%	5.6	29.30%	Plxna4
IP00676957.3	12	22	17	16	19	17	17.2	5.0	29.41%	1.5	8.81%	Ppp2ca
IP00222731.2	14	24	10	20	18	18	17.2	7.2	45.07%	5.3	33.07%	Shcb
IP00113895.1	19	20	19	17	10	19	17.0	0.6	2.99%	4.7	30.82%	Ak1
IP00138716.1	12	22	21	21	9	21	17.0	5.5	30.04%	6.9	40.75%	Aldh1l1
IP00408495.1	17	21	10	19	18	18	17.0	5.6	34.80%	4.9	31.49%	Dlg2
IP00338536.1	19	18	19	17	12	18	17.0	0.6	3.09%	3.6	22.53%	Hspa12a
IP00458393.3	23	0	15	47	0	15	17.0	11.7	92.18%	24.0	116.16%	Rab3d
IP00115157.2	18	17	20	19	10	18	16.8	1.5	8.33%	5.5	33.72%	5730469M1ORik
IP00127408.3	16	16	14	20	18	16	16.8	1.2	7.53%	3.1	17.63%	Capza2

IP100322156.1	17	19	11	20	17	84	17	16.8	4.2	26.57%	4.6	28.64%	D10Jhu81e
IP100316495.3	18	22	16	23	5	84	18	16.8	3.1	16.37%	9.1	61.87%	Enpp6
IP100127417.1	15	12	10	19	28	84	15	16.8	2.5	20.40%	9.0	47.37%	Gbas
IP100187545.4	13	19	13	21	18	84	18	16.8	3.5	23.09%	4.0	23.32%	MaPt
IP100227392.5	22	13	18	22	9	84	18	16.8	4.5	25.52%	6.7	40.77%	Myo1d
IP100132720.2	22	18	10	16	18	84	18	16.8	6.1	36.66%	4.2	28.39%	Ndufb8
IP100875016.1	16	15	15	19	19	84	16	16.8	0.6	3.77%	2.3	13.07%	Sloc6a1
IP100850737.1	21	25	20	9	9	84	20	16.8	2.6	12.03%	6.4	50.14%	Txndc10
IP100620601.1	22	16	17	18	11	84	17	16.8	3.2	17.53%	3.8	24.69%	Usp5
IP100123802.5	12	25	17	14	15	83	15	16.6	6.6	36.43%	1.5	9.96%	Alg2
IP100129319.3	20	25	10	16	12	83	16	16.6	7.6	41.66%	3.1	24.12%	Alp6v0c
IP100130144.2	15	21	21	11	15	83	15	16.6	3.5	18.23%	5.0	32.13%	Cat
IP100468653.3	16	14	12	23	18	83	16	16.6	2.0	14.29%	5.5	31.17%	Ctbp1
IP100314950.2	22	12	17	17	15	83	17	16.6	5.0	29.41%	1.2	7.07%	Lrrc7
IP100229534.5	20	13	16	19	15	83	16	16.6	3.5	21.50%	2.1	12.49%	Pip5k1c
IP100126072.2	25	23	12	14	8	82	14	16.4	7.0	35.00%	3.1	26.96%	2410015M20Rik
IP100649104.1	16	14	17	12	23	82	16	16.4	1.5	9.75%	5.5	31.77%	Acadl
IP100123604.4	20	19	22	9	12	82	19	16.4	1.5	7.51%	6.8	47.49%	D12Erttd551e
IP100123619.6	16	23	14	14	15	82	15	16.4	4.7	26.75%	0.6	4.03%	Grm3
IP100131369.1	22	28	13	13	6	82	13	16.4	7.5	35.95%	4.0	37.89%	Hnrpm
IP100331556.5	21	15	19	15	12	82	15	16.4	3.1	16.66%	3.5	22.90%	Myo1c
IP100608097.1	12	25	18	12	15	82	15	16.4	6.5	35.49%	3.0	20.00%	Pcsk1n
IP100229992.1	17	14	16	17	17	81	17	16.2	1.5	9.75%	0.6	3.46%	Astn1
IP100131407.1	26	17	16	12	10	81	16	16.2	5.5	28.00%	3.1	24.12%	Ephx1
IP100138232.4	20	6	11	18	26	81	18	16.2	7.1	57.52%	7.5	40.94%	Nipsnap1
IP100749860.1	20	18	21	14	8	81	18	16.2	1.5	7.77%	6.5	45.39%	Rplp0
IP100875107.1	12	19	17	17	16	81	17	16.2	3.6	22.53%	0.6	3.46%	Wdr7
IP100112312.3	12	18	17	14	19	80	17	16.0	3.2	20.52%	2.5	15.10%	Cisd1
IP100136716.1	13	26	16	11	14	80	14	16.0	6.8	37.13%	2.5	18.41%	Hsbp1
IP100467447.3	13	13	11	26	17	80	13	16.0	1.2	9.36%	7.5	41.94%	Lrmb2
IP100875695.1	17	13	15	16	19	80	16	16.0	2.0	13.33%	2.1	12.49%	Myc2b
IP100649184.2	16	24	14	18	8	80	16	16.0	5.3	29.40%	5.0	37.75%	Plxna1
IP100128346.1	20	18	17	17	8	80	17	16.0	1.5	8.33%	5.2	37.12%	Sic32a1
IP100119067.3	17	15	17	17	13	79	17	15.8	1.2	7.07%	2.3	14.74%	Acsi3

IP00120302.1	18	20	11	13	17	79	17	15.8	4.7	28.93%	3.1	22.35%	Actr1a
IP00123494.3	11	25	18	15	10	79	15	15.8	7.0	38.89%	4.0	28.20%	Cdh2
IP00109727.1	16	15	17	17	14	79	16	15.8	1.0	6.25%	1.7	10.83%	Hsp110
IP00655177.1	16	19	12	15	17	79	16	15.8	3.5	22.42%	2.5	17.16%	Hspa4
IP00116283.1	23	12	13	14	17	79	14	15.8	6.1	38.02%	2.1	14.19%	Ngef
IP00875011.1	13	18	14	16	18	79	16	15.8	2.6	17.64%	2.0	12.50%	Picalm
IP00404579.1	16	13	12	17	21	79	16	15.8	2.1	15.23%	4.5	27.06%	Scg2
IP00649950.3	18	16	13	13	19	79	16	15.8	2.5	16.06%	3.5	23.09%	Stxn1
IP00321190.1	15	17	17	16	14	79	16	15.8	1.2	7.07%	1.5	9.75%	Sgjp1
IP00136965.1	16	19	20	13	10	78	16	15.6	2.1	11.35%	5.1	35.80%	5730596K20Rik
IP00222429.6	16	11	21	17	13	78	16	15.6	5.0	31.25%	4.0	23.53%	Dst
IP00110885.1	18	11	10	24	15	78	15	15.6	4.4	33.53%	7.1	43.44%	Gphn
IP00117910.3	18	17	11	15	17	78	17	15.6	3.8	24.69%	3.1	21.31%	Igsec1
IP00121575.1	17	11	14	22	14	78	14	15.6	3.0	21.43%	4.6	27.71%	Myo18a
IP00108410.1	11	17	14	14	21	77	14	15.4	3.0	21.43%	4.0	24.74%	Acst2
IP00123870.1	26	0	0	0	51	77	0	15.4	15.0	173.21%	29.4	173.21%	Aldoa
IP00658313.2	9	6	8	13	41	77	9	15.4	1.5	19.92%	17.8	86.06%	Dynll1
IP00119500.1	17	13	22	13	12	77	13	15.4	4.5	26.01%	5.5	35.15%	Etfldh
IP0022325.5	13	19	13	12	20	77	13	15.4	3.5	23.09%	4.4	29.06%	Hadh
IP00123281.1	12	22	22	12	9	77	12	15.4	5.8	30.93%	6.8	47.49%	Hmgcs2
IP00310518.3	15	18	14	11	19	77	15	15.4	2.1	13.29%	4.0	27.56%	Mtch2
IP00311200.1	11	17	14	28	7	77	14	15.4	3.0	21.43%	10.7	65.47%	Ndufa4
IP00137311.1	13	20	16	13	15	77	15	15.4	3.5	21.50%	1.5	10.41%	Ndufb9
IP00515349.1	18	17	16	11	15	77	16	15.4	1.0	5.88%	2.6	18.90%	Npepps
IP00222430.5	16	18	14	14	15	77	15	15.4	2.0	12.50%	0.6	4.03%	Oat
IP00131472.4	18	17	12	14	15	76	15	15.2	3.2	20.52%	1.5	11.18%	Actr1b
IP00121280.5	27	13	10	4	22	76	13	15.2	9.1	54.44%	9.2	76.38%	Mrc1
IP00226891.1	18	18	17	12	11	76	17	15.2	0.6	3.27%	3.2	24.11%	Myo1b
IP00116414.2	16	17	16	17	10	76	16	15.2	0.6	3.53%	3.8	26.41%	Rhot1
IP00113214.1	21	13	5	16	21	76	16	15.2	8.0	61.54%	8.2	58.47%	S100b
IP00651782.2	7	26	17	14	11	75	14	15.0	9.5	57.03%	3.0	21.43%	150003119Rik
IP00272681.2	18	18	18	11	10	75	18	15.0	0.0	0.00%	4.4	33.53%	Hsd17b12
IP00132623.3	21	14	16	14	10	75	14	15.0	3.6	21.21%	3.1	22.91%	Rap2c
IP00225254.6	19	21	19	8	8	75	19	15.0	1.2	5.87%	6.4	54.44%	Sqrdl



IP100229725.2	19	17	13	15	10	74	15	14.8	3.1	18.70%	2.5	19.87%	Alp2a3
IP00133418.1	14	20	16	9	15	74	15	14.8	3.1	18.33%	3.8	28.39%	Decr1
IP00128345.2	14	27	19	9	5	74	14	14.8	6.6	32.79%	7.2	65.56%	Gabbr1
IP00459493.5	19	20	16	7	12	74	16	14.8	2.1	11.35%	4.5	38.65%	Gatm
IP00405699.2	13	8	13	14	25	73	13	14.6	2.9	25.47%	6.7	38.41%	Actr2
IP00128360.1	12	22	14	14	11	73	14	14.6	5.3	33.07%	1.7	13.32%	Adcy5
IP00380717.1	8	18	9	10	28	73	10	14.6	5.5	47.21%	10.7	68.25%	Arf4
IP00311671.4	15	19	11	12	16	73	15	14.6	4.0	26.67%	2.6	20.35%	Cyp2d22
IP00230692.1	16	9	13	20	15	73	15	14.6	3.5	27.73%	3.6	22.53%	Dbi
IP00750595.2	20	18	21	5	9	73	18	14.6	1.5	7.77%	8.3	71.37%	lggap1
IP00624363.4	0	0	73	0	0	73	0	14.6	42.1	173.21%	42.1	173.21%	Krt7
IP00114380.1	15	15	11	13	19	73	15	14.6	2.3	16.90%	4.2	29.05%	Palm
IP00380997.3	14	19	18	16	6	73	16	14.6	2.6	15.56%	6.4	48.22%	Psmd2
IP00466588.3	17	18	13	12	13	73	13	14.6	2.6	16.54%	0.6	4.56%	Vat1
IP00649326.2	18	15	17	13	9	72	15	14.4	1.5	9.17%	4.0	30.77%	Add2
IP00331543.1	15	22	17	14	4	72	15	14.4	3.6	20.03%	6.8	58.34%	Ccdc47
IP00337980.5	10	24	15	13	10	72	13	14.4	7.1	43.44%	2.5	19.87%	Glod4
IP00125929.2	15	18	15	11	13	72	15	14.4	1.7	10.83%	2.0	15.38%	Gria3
IP00118380.2	18	16	8	19	11	72	16	14.4	5.3	37.80%	5.7	44.89%	Hp1bp3
IP00135651.1	14	23	16	14	5	72	14	14.4	4.7	26.75%	5.9	50.22%	Iga2b
IP00623241.3	14	19	15	9	15	72	15	14.4	2.6	16.54%	3.5	26.65%	Ntrk2
IP00828920.1	13	29	16	6	7	71	13	14.2	8.5	43.99%	5.5	56.97%	Esam1
IP00226932.2	10	16	11	14	20	71	14	14.2	3.2	26.06%	4.6	30.55%	Hspa1b
IP00271951.5	15	18	10	15	13	71	15	14.2	4.0	28.20%	2.5	19.87%	Ncald
IP00222767.1	19	13	16	11	12	71	13	14.2	3.0	18.75%	2.6	20.35%	Nos3
IP00647986.1	16	19	18	9	8	70	16	14.0	1.5	8.65%	5.5	47.21%	Mosc2
IP00875068.1	15	17	19	9	10	70	15	14.0	2.0	11.76%	5.5	43.48%	Ptrf
IP00458612.5	12	11	11	17	19	70	12	14.0	0.6	5.09%	4.2	26.57%	Ttyh1
IP00121514.3	21	12	12	14	10	69	12	13.8	5.2	34.64%	2.0	16.67%	Agk
IP00222557.5	8	20	16	10	15	69	15	13.8	6.1	41.66%	3.2	23.52%	Dci
IP00263638.8	14	15	14	21	5	69	14	13.8	0.6	4.03%	8.0	60.16%	Gstm5
IP00132276.1	18	11	16	19	5	69	16	13.8	3.6	24.04%	7.4	55.28%	Lama1
IP00468605.4	14	15	10	12	18	69	14	13.8	2.6	20.35%	4.2	31.22%	Lgi1
IP00126186.1	21	7	12	17	12	69	12	13.8	7.1	53.21%	2.9	21.12%	Lphn1

IP100869393.1	9	15	14	18	13	69	14	13.8	3.2	25.38%	2.6	17.64%	Mapk1
IP100471246.2	19	14	17	11	8	69	14	13.8	2.5	15.10%	4.6	38.19%	Pdia4
IP100264501.8	9	15	18	12	15	69	15	13.8	4.6	32.73%	3.0	20.00%	Rasa1
IP100651798.1	11	11	21	17	9	69	11	13.8	5.8	40.28%	6.1	39.00%	RP23-298F22.4
IP100130589.8	17	22	16	10	4	69	16	13.8	3.2	17.53%	6.0	60.00%	Sfpq
IP100556840.1	18	17	13	12	9	69	13	13.8	2.6	16.54%	2.1	18.37%	Slc25a23
IP100474756.1	16	13	10	19	11	69	13	13.8	3.0	23.08%	4.9	37.00%	Slp1
IP100118787.1	15	24	12	11	7	69	12	13.8	6.2	36.74%	2.6	26.46%	Stoml2
IP100876412.1	16	15	15	8	14	68	15	13.6	0.6	3.77%	3.8	30.70%	C230096C10Rik
IP100466276.2	15	17	11	15	10	68	15	13.6	3.1	21.31%	2.6	22.05%	Cct3
IP100420718.4	14	16	10	17	11	68	14	13.6	3.1	22.91%	3.8	29.89%	Cntn2
IP100135231.2	15	7	8	26	12	68	12	13.6	4.4	43.59%	9.5	61.64%	EG667952
IP100337844.4	12	18	16	13	9	68	13	13.6	3.1	19.92%	3.5	27.73%	Rgs7
IP100828488.1	20	13	16	10	9	68	13	13.6	3.5	21.50%	3.8	32.45%	Rras
IP100113606.2	12	22	14	8	12	68	12	13.6	5.3	33.07%	3.1	26.96%	Yes1
IP100331436.4	0	0	23	44	0	67	0	13.4	13.3	173.21%	22.0	98.54%	Alp12a
IP100129178.1	14	15	16	14	8	67	14	13.4	1.0	6.67%	4.2	32.87%	Eftud2
IP100127415.1	15	13	11	16	12	67	13	13.4	2.0	15.38%	2.6	20.35%	Hcn2
IP100117689.1	19	12	5	14	17	67	14	13.4	7.0	58.33%	6.2	52.04%	Nduif7
IP100128973.1	13	23	16	13	2	67	13	13.4	5.1	29.61%	7.4	71.33%	Npm1
IP100132390.5	14	14	15	13	11	67	14	13.4	0.6	4.03%	2.0	15.38%	Rab21
IP100223446.5	10	13	10	12	22	67	12	13.4	1.7	15.75%	6.4	43.83%	Stx12
IP100121276.4	11	19	16	9	11	66	11	13.2	4.0	26.36%	3.6	30.05%	1200015F23Rik
IP100229554.5	14	19	11	14	8	66	14	13.2	4.0	27.56%	3.0	27.27%	Abcd3
IP100122633.3	20	13	14	14	5	66	14	13.2	3.8	24.17%	5.2	47.24%	Usp9x
IP100120342.4	19	15	16	10	5	65	15	13.0	2.1	12.49%	5.5	53.30%	Atad1
IP100131471.1	13	13	11	11	17	65	13	13.0	1.2	9.36%	3.5	26.65%	Dnm2
IP100661414.2	13	16	19	13	4	65	13	13.0	3.0	18.75%	7.5	62.92%	Gm996
IP100271986.6	16	19	12	11	7	65	12	13.0	3.5	22.42%	2.6	26.46%	Igsec2
IP100137313.2	23	11	15	9	7	65	11	13.0	6.1	37.41%	4.2	40.29%	Krt10
IP100123004.3	12	14	10	17	12	65	12	13.0	2.0	16.67%	3.6	27.74%	Napg
IP100116228.1	18	19	7	7	14	65	14	13.0	6.7	45.40%	4.0	43.30%	Scarb2
IP100153792.1	16	15	14	4	15	64	15	12.8	1.0	6.67%	6.1	55.30%	Aldh4a1
IP100387379.1	11	10	14	12	17	64	12	12.8	2.1	17.84%	2.5	17.56%	Ctla

IP100135686.2	14	14	13	12	11	64	13	12.8	0.6	4.22%	1.0	8.33%	Gabrg2
IP100798492.2	13	17	15	9	10	64	13	12.8	2.0	13.33%	3.2	28.36%	Hmgcl
IP100331174.5	11	18	14	13	8	64	13	12.8	3.5	24.50%	3.2	27.55%	Pdk3
IP100135604.5	11	16	9	14	14	64	14	12.8	3.6	30.05%	2.9	23.41%	Prkacb
IP100473293.1	12	20	14	6	12	64	12	12.8	4.2	27.15%	4.2	39.03%	Shank3
IP100110721.5	16	17	13	13	5	64	13	12.8	2.1	13.58%	4.6	44.70%	Stom
IP100313390.3	19	15	11	11	7	63	11	12.6	4.0	26.67%	2.3	23.89%	Anks1b
IP100816879.1	13	14	19	6	11	63	13	12.6	3.2	20.96%	6.6	54.65%	Atad3a
IP100321446.4	11	18	10	11	13	63	11	12.6	4.4	33.53%	1.5	13.48%	Atp5j2
IP100117829.1	17	11	13	15	7	63	13	12.6	3.1	22.35%	4.2	35.69%	Cct7
IP100742329.1	18	7	9	13	16	63	13	12.6	5.9	51.70%	3.5	27.73%	Ctsd
IP100877232.1	9	12	9	13	20	63	12	12.6	1.7	17.32%	5.6	39.77%	Gnal
IP100135130.3	10	14	16	15	8	63	14	12.6	3.1	22.91%	4.4	33.53%	Gria1
IP100125960.1	14	15	15	8	11	63	14	12.6	0.6	3.94%	3.5	30.99%	Mthfd1l
IP100126913.5	10	17	15	13	8	63	13	12.6	3.6	25.75%	3.6	30.05%	Ppp1r7
IP100128989.1	13	22	10	14	4	63	13	12.6	6.2	41.63%	5.0	53.93%	Psmab5
IP100122865.1	14	8	15	19	7	63	14	12.6	3.8	30.70%	6.1	44.71%	Ranbp2
IP100153990.1	16	13	15	19	0	63	15	12.6	1.5	10.41%	10.0	88.38%	Sic25a13
IP100126827.1	16	14	13	13	7	63	13	12.6	1.5	10.66%	3.5	31.49%	Tcp1
IP100474660.1	11	14	14	11	12	62	12	12.4	1.7	13.32%	1.5	12.39%	Cpne6
IP100128454.1	18	19	12	8	5	62	12	12.4	3.8	23.18%	3.5	42.14%	Rpl4
IP100114594.1	14	11	14	12	11	62	12	12.4	1.7	13.32%	1.5	12.39%	Sep2
IP100134191.3	25	6	10	10	11	62	10	12.4	10.0	73.29%	0.6	5.59%	Sic24a2
IP100115458.1	10	16	19	12	4	61	12	12.2	4.6	30.55%	7.5	64.33%	Cend1
IP100408176.2	10	20	12	6	13	61	12	12.2	5.3	37.80%	3.8	36.64%	Crym
IP100129516.1	12	12	15	13	9	61	12	12.2	1.7	13.32%	3.1	24.77%	Csnk2a1
IP100465871.1	13	9	10	20	9	61	10	12.2	2.1	19.52%	6.1	46.79%	Fgb
IP100466708.3	21	9	7	15	9	61	9	12.2	7.6	61.39%	4.2	40.29%	Gng2
IP100117300.3	14	8	14	12	13	61	13	12.2	3.5	28.87%	1.0	7.69%	Grin2b
IP100318645.6	15	12	13	10	11	61	12	12.2	1.5	11.46%	1.5	13.48%	Iitga7
IP100221998.4	12	17	13	11	8	61	12	12.2	2.6	18.90%	2.5	23.59%	Prkaca
IP100120232.1	8	14	12	13	14	61	13	12.2	3.1	26.96%	1.0	7.69%	Rab18
IP100131445.2	9	18	16	13	5	61	13	12.2	4.7	32.97%	5.7	50.17%	Snx22
IP100758248.3	14	9	11	9	18	61	11	12.2	2.5	22.21%	4.7	37.31%	Sod1

IP00129430.1	8	12	15	15	10	60	12	12.0	3.5	30.10%	2.9	21.65%	Actr3b
IP00223987.1	14	12	14	10	10	60	12	12.0	1.2	8.66%	2.3	20.38%	Atp5j
IP00320016.7	18	14	15	5	8	60	14	12.0	2.1	13.29%	5.1	54.98%	C33000219Rik
IP00881096.1	11	14	14	10	11	60	11	12.0	1.7	13.32%	2.1	17.84%	Dlg3
IP00453821.2	17	10	9	9	15	60	10	12.0	4.4	36.32%	3.5	31.49%	Fahd2a
IP00462752.4	12	16	12	10	10	60	12	12.0	2.3	17.32%	1.2	10.83%	Gria4
IP00118769.2	14	9	12	12	13	60	12	12.0	2.5	21.57%	0.6	4.68%	Grin2a
IP00469204.2	11	14	20	9	6	60	11	12.0	4.6	30.55%	7.4	63.18%	Hmox2
IP00116558.1	16	18	13	8	5	60	13	12.0	2.5	16.06%	4.0	46.63%	Pdhx
IP00120225.1	15	10	10	17	8	60	10	12.0	2.9	24.74%	4.7	40.51%	Sic16a1
IP00875833.1	9	11	17	13	10	60	11	12.0	4.2	33.76%	3.5	26.34%	Tom1l2
IP00807902.1	11	7	19	8	14	59	11	11.8	6.1	49.54%	5.5	40.30%	Acot9
IP00224069.2	19	13	11	10	6	59	11	11.8	4.2	29.05%	2.6	29.40%	Angptl7
IP00667471.1	17	7	6	29	0	59	7	11.8	6.1	60.83%	15.3	131.21%	Col4a1
IP00230449.7	16	8	16	10	9	59	10	11.8	4.6	34.64%	3.8	32.45%	ltp2
IP00875877.1	13	14	14	12	6	59	13	11.8	0.6	4.22%	4.2	39.03%	Lrrc59
IP00404687.5	10	13	7	19	10	59	10	11.8	3.0	30.00%	6.2	52.04%	Ndufs6
IP00420221.3	11	20	13	15	0	59	13	11.8	4.7	32.22%	8.1	87.26%	Rac3
IP00404551.1	9	15	13	8	13	58	13	11.6	3.1	24.77%	2.9	25.47%	Arpc2
IP00109153.6	14	13	12	10	9	58	12	11.6	1.0	7.69%	1.5	14.78%	Bes1
IP00775867.1	13	16	15	9	5	58	13	11.6	1.5	10.41%	5.0	52.07%	Cav1
IP00341550.1	13	13	12	10	10	58	12	11.6	0.6	4.56%	1.2	10.83%	Cd47
IP00623284.4	17	11	13	9	8	58	11	11.6	3.1	22.35%	2.6	26.46%	Ddost
IP00322033.7	11	17	9	11	10	58	11	11.6	4.2	33.76%	1.0	10.00%	Dhrs1
IP00620582.1	14	10	12	13	9	58	12	11.6	2.0	16.67%	2.1	18.37%	Dnajc11
IP00125681.4	17	9	9	14	9	58	9	11.6	4.6	39.59%	2.9	27.06%	Grin1
IP00468924.5	12	15	10	11	10	58	11	11.6	2.5	20.40%	0.6	5.59%	Igf1r
IP00131771.3	21	8	12	6	11	58	11	11.6	6.7	48.72%	3.2	33.25%	LOC433157
IP00112986.1	8	17	12	8	13	58	12	11.6	4.5	36.56%	2.6	24.05%	Mecr
IP00754876.2	11	14	13	11	9	58	11	11.6	1.5	12.06%	2.0	18.18%	Piprf
IP00776169.1	14	13	5	24	2	58	13	11.6	4.9	46.25%	11.9	115.46%	Tinagl
IP00268673.5	11	15	14	11	6	57	11	11.4	2.1	15.61%	4.0	39.11%	Dync1l1i
IP00122399.1	10	3	9	13	22	57	10	11.4	3.8	51.63%	6.7	45.40%	ENSMUSG00000075245
IP00406901.2	14	16	15	4	8	57	14	11.4	1.0	6.67%	5.6	61.86%	Glg1

IP00119663.3	6	11	11	13	16	57	11	11.4	2.9	30.93%	2.5	18.87%	Gpr158
IP00134476.2	11	20	9	11	6	57	11	11.4	5.9	43.95%	2.5	29.04%	Gpsn2
IP00115117.1	10	14	11	8	14	57	11	11.4	2.1	17.84%	3.0	27.27%	Gyk
IP00118895.1	11	17	14	6	9	57	11	11.4	3.0	21.43%	4.0	41.81%	mt-Nd1
IP00420870.4	14	10	7	9	17	57	10	11.4	3.5	33.99%	5.3	48.10%	Myl9
IP00875100.1	13	8	12	8	16	57	12	11.4	2.6	24.05%	4.0	33.33%	Ndufa11
IP00553576.3	15	9	4	15	14	57	14	11.4	5.5	59.01%	6.1	55.30%	Ndufs5
IP00463468.3	7	21	14	8	7	57	8	11.4	7.0	50.00%	3.8	39.16%	Pfn1
IP00555114.1	14	13	12	8	10	57	12	11.4	1.0	7.69%	2.0	20.00%	Piges2
IP00134131.2	14	9	13	11	10	57	11	11.4	2.6	22.05%	1.5	13.48%	Sic9a3r1
IP00331663.6	14	14	8	10	11	57	11	11.4	3.5	28.87%	1.5	15.80%	Syt7
IP00116770.1	9	10	14	12	12	57	12	11.4	2.6	24.05%	1.2	9.12%	Ttc7b
IP00230185.5	6	11	10	20	10	57	10	11.4	2.6	29.40%	5.8	43.30%	Uqcrq
IP00874996.1	18	11	11	10	6	56	11	11.2	4.0	30.31%	2.6	29.40%	Apoo
IP00458583.3	16	5	7	10	18	56	10	11.2	5.9	62.78%	5.7	48.74%	Atf8b
IP00115454.3	11	5	9	16	15	56	11	11.2	3.1	36.66%	3.8	28.39%	Arpc1a
IP00761241.2	11	12	10	13	10	56	11	11.2	1.0	9.09%	1.7	15.75%	Alp6v1d
IP00129924.6	15	7	10	14	10	56	10	11.2	4.0	37.89%	2.3	20.38%	Cacna2d2
IP00119575.3	17	15	13	6	5	56	13	11.2	2.0	13.33%	4.4	54.49%	Dock9
IP00177038.1	14	6	13	14	9	56	13	11.2	4.4	39.63%	2.6	22.05%	Etfb
IP00114671.1	8	17	13	9	9	56	9	11.2	4.5	35.60%	2.3	22.35%	Fyn
IP00127472.2	20	9	5	9	13	56	9	11.2	7.8	68.54%	4.0	44.44%	Gad2
IP00323748.2	8	12	12	6	18	56	12	11.2	2.3	21.65%	6.0	50.00%	lvd
IP00312174.6	14	8	14	11	9	56	11	11.2	3.5	28.87%	2.5	22.21%	Neo1
IP00127987.1	13	13	11	13	6	56	13	11.2	1.2	9.36%	3.6	36.06%	Nlgn2
IP0011412.3	11	11	12	11	11	56	11	11.2	0.6	5.09%	0.6	5.09%	Shank1
IP00457611.2	12	14	9	13	8	56	12	11.2	2.5	21.57%	2.6	26.46%	Sic6a17
IP00116154.2	14	10	10	12	10	56	10	11.2	2.3	20.38%	1.2	10.83%	Sorbs1
IP00281761.3	13	8	15	14	6	56	13	11.2	3.6	30.05%	4.9	42.28%	Stxbp5
IP00125460.1	14	8	13	12	8	55	12	11.0	3.2	27.55%	2.6	24.05%	Acs11
IP00323442.8	11	15	13	9	7	55	11	11.0	2.0	15.38%	3.1	31.60%	Aldh1a1
IP00128986.1	17	9	9	9	11	55	9	11.0	4.6	39.59%	1.2	11.95%	Cadps2
IP00229256.5	14	11	15	10	5	55	11	11.0	2.1	15.61%	5.0	50.00%	Cct4
IP00651878.1	12	14	12	7	10	55	12	11.0	1.2	9.12%	2.5	26.03%	Dbn1

IP00314069.1	11	12	12	11	9	55	11	11.0	0.6	4.95%	1.5	14.32%	Dstn
IP00165688.2	13	11	10	12	9	55	11	11.0	1.5	13.48%	1.5	14.78%	Gg1a1
IP00154004.1	9	13	9	19	5	55	9	11.0	2.3	22.35%	7.2	65.56%	Hap1n1
IP00474705.2	13	9	9	14	10	55	10	11.0	2.3	22.35%	2.6	24.05%	Hcn1
IP00337050.1	13	9	13	15	5	55	13	11.0	2.3	19.79%	5.3	48.10%	ldh1
IP00672663.1	13	13	10	15	4	55	13	11.0	1.7	14.43%	5.5	56.97%	Kif5c
IP00129159.1	11	12	8	14	10	55	11	11.0	2.1	20.15%	3.1	28.64%	Qdpr
IP00853905.2	11	16	9	11	8	55	11	11.0	3.6	30.05%	1.5	16.37%	Rph3a
IP00111158.2	13	13	10	10	8	54	10	10.8	1.7	14.43%	1.2	12.37%	Abcb7
IP00122826.1	10	12	12	12	8	54	12	10.8	1.2	10.19%	2.3	21.65%	Aldh1b1
IP00108783.1	9	7	6	7	25	54	7	10.8	1.5	20.83%	10.7	84.42%	Ap2s1
IP00283600.2	9	8	18	12	7	54	9	10.8	5.5	47.21%	5.5	44.66%	Arfgap1
IP00331361.2	7	11	18	7	11	54	11	10.8	5.6	46.40%	5.6	46.40%	Cds2
IP00116277.3	7	17	11	14	5	54	11	10.8	5.0	43.14%	4.6	45.83%	Cse1l
IP00118654.2	14	14	17	5	4	54	14	10.8	1.7	11.55%	7.2	83.47%	Dtnb
IP00116688.1	5	13	16	14	6	54	13	10.8	5.7	50.17%	5.3	44.10%	Lama4
IP00117811.3	12	13	8	11	10	54	11	10.8	2.6	24.05%	1.5	15.80%	Mgll
IP00127447.3	6	10	10	13	15	54	10	10.8	2.3	26.65%	2.5	19.87%	Nup155
IP00121440.4	12	9	13	15	5	54	12	10.8	2.1	18.37%	5.3	48.10%	Ptpra
IP00127492.1	0	0	54	0	0	54	0	10.8	31.2	173.21%	31.2	173.21%	Slc25a31
IP00387580.1	14	17	10	5	8	54	10	10.8	3.5	25.70%	2.5	32.83%	Tmem24
IP00381088.5	16	9	9	6	14	54	9	10.8	4.0	35.66%	4.0	41.81%	Vps35
IP00230005.3	11	8	11	14	9	53	11	10.6	1.7	17.32%	2.5	22.21%	2310001A20Rik
IP00653841.1	9	14	14	11	5	53	11	10.6	2.9	23.41%	4.6	45.83%	Ckap4
IP00313539.4	8	13	9	12	11	53	11	10.6	2.6	26.46%	1.5	14.32%	Cox6c
IP00127050.1	12	14	13	8	6	53	12	10.6	1.0	7.69%	3.6	40.06%	Cyp46a1
IP00626662.3	8	13	9	12	11	53	11	10.6	2.6	26.46%	1.5	14.32%	EG621837
IP00127942.4	12	14	11	11	5	53	11	10.6	1.5	12.39%	3.5	38.49%	Gpdl
IP00108388.1	8	8	13	18	6	53	8	10.6	2.9	29.86%	6.0	48.87%	Lamb1-1
IP00320462.4	12	13	10	16	2	53	12	10.6	1.5	13.09%	7.0	75.25%	Mybbp1a
IP00468999.4	17	6	5	14	11	53	11	10.6	6.7	71.34%	4.6	45.83%	Slc17a7
IP00331609.7	11	14	15	9	4	53	11	10.6	2.1	15.61%	5.5	59.01%	Slc25a24
IP00113073.3	8	11	13	9	12	53	11	10.6	2.5	23.59%	2.1	18.37%	Slc2a3
IP00623995.3	8	12	12	10	11	53	11	10.6	2.3	21.65%	1.0	9.09%	Trio

IP	11	12	7	11	52	11	10.4	0.6	5.09%	2.6	26.46%	A930041I02Rik
IP100153840.2	11	11	7	11	52	11	10.4	0.6	5.09%	2.6	26.46%	Acadsb
IP100719919.2	13	7	7	17	52	8	10.4	3.2	34.44%	5.8	55.87%	Arl6ip5
IP100135443.2	13	9	10	11	52	10	10.4	2.3	22.35%	1.0	10.00%	Cplx2
IP100309322.3	8	10	8	18	52	8	10.4	1.2	13.32%	5.3	44.10%	Ddx1
IP100223047.2	13	12	11	6	52	11	10.4	1.5	13.09%	3.2	33.25%	Endogl1
IP100459279.2	13	7	14	10	52	10	10.4	3.2	34.44%	3.1	28.64%	Gad1
IP100118692.6	13	14	9	8	52	9	10.4	3.2	27.55%	3.2	31.11%	Gpm6b
IP100116697.1	6	9	10	13	52	10	10.4	4.0	41.81%	2.1	19.52%	Hpcal1
IP100121634.1	11	12	7	9	52	11	10.4	1.0	8.33%	3.1	31.60%	Me2
IP100469194.2	9	13	10	8	52	10	10.4	2.1	18.37%	2.0	20.00%	Oxr1
IP100828306.2	8	14	10	8	52	10	10.4	3.1	26.96%	2.0	20.00%	Rims1
IP100112032.1	10	14	9	7	52	10	10.4	2.0	16.67%	2.5	26.96%	Sic27a4
IP100131995.1	12	19	9	3	52	9	10.4	5.1	38.49%	3.5	49.49%	Sic4a8
IP100265386.5	10	12	11	9	52	10	10.4	1.2	10.83%	1.0	10.00%	Top2b
IP100122011.2	7	11	19	4	52	11	10.4	2.3	23.89%	7.5	66.23%	Vamp1
IP100678532.3	9	11	10	12	52	10	10.4	1.0	10.00%	1.0	9.09%	Cyb5b
IP100123199.1	13	8	8	14	51	8	10.2	2.9	29.86%	3.5	34.64%	Dnaja1
IP100110264.2	6	13	9	10	51	10	10.2	4.0	37.89%	2.1	19.52%	Farp1
IP100468140.2	10	12	6	14	51	10	10.2	1.5	14.78%	4.0	41.81%	Itgav
IP100121443.1	9	16	9	7	51	9	10.2	3.8	32.45%	1.5	17.63%	Lnpep
IP100880817.1	11	15	6	6	51	11	10.2	2.0	15.38%	4.0	48.50%	Mpdz
IP10011416.1	12	6	13	11	51	11	10.2	3.0	33.33%	2.0	18.18%	Ndrp1
IP100874973.1	13	13	7	6	51	12	10.2	0.6	4.56%	3.2	38.57%	Phgdh
IP100664176.1	10	10	13	7	51	10	10.2	0.6	5.59%	3.1	29.57%	Sept9
IP100131881.1	14	10	11	7	51	10	10.2	2.6	24.05%	2.0	22.22%	Tommm40
IP100109311.3	9	11	16	5	51	10	10.2	1.0	10.00%	5.5	53.30%	Ubln2
IP100848866.1	9	15	12	3	51	12	10.2	3.0	25.00%	5.2	57.74%	Cbr1
IP100119785.2	8	13	9	11	50	9	10.0	2.6	26.46%	1.2	11.95%	Col4a2
IP100115620.1	8	10	22	0	50	10	10.0	1.2	12.37%	11.0	103.27%	ENSMUSG0000058126
IP100648927.1	12	6	13	7	50	12	10.0	3.5	34.64%	3.2	30.14%	Erb2ip
IP100136350.3	12	10	5	11	50	11	10.0	1.2	10.19%	3.8	40.56%	Mapre3
IP100121322.2	10	7	13	13	50	10	10.0	1.7	21.65%	3.5	31.49%	Mtap4
IP100318204.5	10	14	7	7	50	10	10.0	2.0	16.67%	2.9	33.31%	Mut
IP100828670.2	10	7	11	13	50	10	10.0	1.5	17.63%	2.0	18.18%	

IP00380983.3	17	17	13	3	0	50	13	10.0	2.3	14.74%	6.8	127.63%	Myel2
IP00122353.1	13	11	7	15	4	50	11	10.0	3.1	29.57%	5.7	65.61%	Myo6
IP00120030.1	10	8	8	12	12	50	10	10.0	1.2	13.32%	2.3	21.65%	Otub1
IP00339916.9	14	9	10	10	7	50	10	10.0	2.6	24.05%	1.7	19.25%	Ptxna2
IP00351827.6	8	8	9	11	14	50	9	10.0	0.6	6.93%	2.5	22.21%	Prdx1
IP00132208.1	12	10	11	7	10	50	10	10.0	1.0	9.09%	2.1	22.30%	Tagln3
IP00420745.7	14	11	12	7	6	50	11	10.0	1.5	12.39%	3.2	38.57%	Ubxnd8
IP00228693.4	12	10	11	8	9	50	10	10.0	1.0	9.09%	1.5	16.37%	Unc13a
IP00757097.2	16	8	12	8	5	49	8	9.8	4.0	33.33%	3.5	42.14%	Abhd6
IP00468380.4	8	16	9	6	10	49	9	9.8	4.4	39.63%	2.1	24.98%	Ak3
IP00125939.2	11	11	10	8	9	49	10	9.8	0.6	5.41%	1.0	11.11%	Ak3l1
IP00314191.6	11	12	7	10	9	49	10	9.8	2.6	26.46%	1.5	17.63%	Ap3d1
IP00331549.1	13	8	11	11	6	49	11	9.8	2.5	23.59%	2.9	30.93%	Chchd6
IP00322760.7	10	8	9	8	14	49	9	9.8	1.0	11.11%	3.2	31.11%	Cox6a1
IP00626950.2	7	18	9	11	4	49	9	9.8	5.9	51.70%	3.6	45.07%	Grim2
IP00321320.3	14	8	14	13	0	49	13	9.8	3.5	28.87%	7.8	86.78%	LOC100045332
IP00117929.1	11	5	5	9	19	49	9	9.8	3.5	49.49%	7.2	65.56%	Mtx2
IP00313222.5	14	9	16	8	2	49	9	9.8	3.6	27.74%	7.0	81.04%	Nono
IP00118978.4	12	13	0	14	10	49	12	9.8	7.2	86.81%	7.2	90.14%	Prkca
IP00108685.3	11	4	11	15	8	49	11	9.8	4.0	46.63%	3.5	30.99%	Psat1
IP00230003.7	6	14	15	7	7	49	7	9.8	4.9	42.28%	4.6	47.78%	Ptplad1
IP00129253.2	12	14	10	13	0	49	12	9.8	2.0	16.67%	6.8	88.79%	Sf3b3
IP00662342.3	13	17	8	5	6	49	8	9.8	4.5	35.60%	1.5	24.12%	Sic27a1
IP00120097.1	12	14	11	7	5	49	11	9.8	1.5	12.39%	3.1	39.85%	Stxbp5l
IP00225961.5	8	8	9	15	9	49	9	9.8	0.6	6.93%	3.5	31.49%	Vcan
IP0011181.1	17	7	8	5	11	48	8	9.6	5.5	51.63%	3.0	37.50%	5230400G24Rik
IP00225209.1	16	12	9	3	8	48	9	9.6	3.5	28.47%	3.2	48.22%	Abcg2
IP00338452.3	9	11	10	11	7	48	10	9.6	1.0	10.00%	2.1	22.30%	Bphl
IP00467661.3	16	10	6	11	5	48	10	9.6	5.0	47.19%	3.2	43.83%	EG629383
IP00136251.1	16	7	9	10	6	48	9	9.6	4.7	44.30%	2.1	24.98%	Entpd1
IP00118191.2	14	9	7	12	6	48	9	9.6	3.6	36.06%	3.2	38.57%	Exoc4
IP00121788.1	10	11	8	14	5	48	10	9.6	1.5	15.80%	4.6	50.92%	Gng3
IP00315135.3	10	14	11	5	8	48	10	9.6	2.1	17.84%	3.0	37.50%	Ly75
IP00420185.1	10	14	6	9	9	48	9	9.6	4.0	40.00%	1.7	21.65%	Mpst



IP00224740.6	12	7	7	6	16	48	7	9.6	2.9	33.31%	5.5	56.97%	Ppp2r2a
IP00116394.1	14	7	9	12	6	48	9	9.6	3.6	36.06%	3.0	33.33%	Ranbp5
IP00331490.5	11	19	6	8	4	48	8	9.6	6.6	54.65%	2.0	33.33%	Rpl12
IP00111151.2	10	6	10	17	5	48	10	9.6	2.3	26.65%	6.0	56.51%	Vwa1
IP00221769.5	13	7	9	6	12	47	9	9.4	3.1	31.60%	3.0	33.33%	Car4
IP00109044.8	11	8	11	8	9	47	9	9.4	1.7	17.32%	1.5	16.37%	Cpne9
IP00118821.2	11	7	15	7	7	47	7	9.4	4.0	36.36%	4.6	47.78%	Cpt2
IP00134621.3	13	13	11	6	4	47	11	9.4	1.2	9.36%	3.6	51.51%	Dbt
IP00115708.1	10	11	16	5	5	47	10	9.4	3.2	26.06%	6.4	73.28%	Ddx3x
IP00121627.3	14	7	7	10	9	47	9	9.4	4.0	43.30%	1.5	17.63%	Eps15l1
IP00311443.1	13	12	9	6	7	47	9	9.4	2.1	18.37%	1.5	20.83%	Gucy1a2
IP00316740.4	10	9	12	10	6	47	10	9.4	1.5	14.78%	3.1	32.73%	Kif5b
IP00109672.3	6	8	12	11	10	47	10	9.4	3.1	35.25%	1.0	9.09%	Ogt
IP00468691.3	5	14	16	9	3	47	9	9.4	5.9	50.22%	6.5	69.71%	Pdcd6ip
IP00119131.2	12	9	12	9	5	47	9	9.4	1.7	15.75%	3.5	40.52%	Pde10a
IP00387421.1	19	9	6	9	4	47	9	9.4	6.8	60.06%	2.5	39.74%	Rpl7
IP00856692.1	10	12	11	9	5	47	10	9.4	1.0	9.09%	3.1	36.66%	Scn1a
IP00474157.4	9	9	9	11	9	47	9	9.4	0.0	0.00%	1.2	11.95%	Sic12a6
IP00468960.1	11	8	8	14	6	47	8	9.4	1.7	19.25%	4.2	44.61%	Srgap3
IP00133399.1	9	9	14	7	8	47	9	9.4	2.9	27.06%	3.8	39.16%	Trap1
IP00116254.1	13	6	11	8	8	46	8	9.2	3.6	36.06%	1.7	19.25%	Ccdc141
IP00751834.1	7	17	9	9	4	46	9	9.2	5.3	48.10%	2.9	39.36%	Eef1g
IP00323483.3	6	12	13	7	8	46	8	9.2	3.8	36.64%	3.2	34.44%	Map2k1
IP00308484.3	12	11	10	6	7	46	10	9.2	1.0	9.09%	2.1	27.15%	Nbea
IP00135475.3	8	8	12	18	0	46	8	9.2	2.3	24.74%	9.2	91.65%	Nid2
IP00719950.2	11	12	10	6	7	46	10	9.2	1.0	9.09%	2.1	27.15%	Prodh
IP00556814.3	11	11	14	10	0	46	11	9.2	1.7	14.43%	7.2	90.14%	Rab6
IP00336291.6	9	16	7	9	5	46	9	9.2	4.7	44.30%	2.0	28.57%	Rpl6
IP00620804.2	10	12	10	10	4	46	10	9.2	1.2	10.83%	3.5	43.30%	Sf3b1
IP00112414.1	11	6	6	12	11	46	11	9.2	2.9	37.65%	3.2	33.25%	Sic17a6
IP00129268.1	9	12	10	11	4	46	10	9.2	1.5	14.78%	3.8	45.43%	Sntb2
IP00130535.1	11	6	12	11	6	46	11	9.2	3.2	33.25%	3.2	33.25%	Tpp2
IP00315794.1	13	10	7	8	8	46	8	9.2	3.0	30.00%	0.6	7.53%	Txndc14
IP00224697.1	7	13	13	6	6	45	7	9.0	3.5	31.49%	4.0	48.50%	1110031B06Rik

IP00626660.3	8	17	8	5	7	45	8	9.0	5.2	47.24%	1.5	22.91%	Acaa1b
IP00129512.3	14	7	12	6	6	45	7	9.0	3.6	32.78%	3.5	43.30%	Agpat3
IP00875405.1	7	15	5	12	6	45	7	9.0	5.3	58.79%	3.8	49.38%	Arpc3
IP00312244.3	7	19	9	10	0	45	9	9.0	6.4	55.11%	5.5	86.96%	Bat5
IP00828649.1	11	5	12	13	4	45	11	9.0	3.8	40.56%	4.9	51.03%	Cadm4
IP00313750.3	11	11	8	11	4	45	11	9.0	1.7	17.32%	3.5	45.81%	Capza1
IP00118173.1	12	10	8	9	6	45	9	9.0	2.0	20.00%	1.5	19.92%	Cdpt
IP00126085.1	7	8	18	12	0	45	8	9.0	6.1	55.30%	9.2	91.65%	Des
IP00831546.1	7	15	11	8	4	45	8	9.0	4.0	36.36%	3.5	45.81%	Dgke
IP00387374.1	8	15	9	5	8	45	8	9.0	3.8	35.49%	2.1	28.39%	Erlin2
IP00279079.1	9	10	10	11	5	45	10	9.0	0.6	5.97%	3.2	37.09%	Fbxo41
IP00129959.2	12	9	10	5	9	45	9	9.0	1.5	14.78%	2.6	33.07%	Ppa2
IP00114667.1	10	8	11	8	8	45	8	9.0	1.5	15.80%	1.7	19.25%	Psd3
IP00278804.1	11	8	12	8	6	45	8	9.0	2.1	20.15%	3.1	35.25%	Rras2
IP00119896.2	9	11	12	9	4	45	9	9.0	1.5	14.32%	4.0	48.50%	Ruvbl1
IP00127172.3	8	8	11	11	7	45	8	9.0	1.7	19.25%	2.3	23.89%	Sept2
IP00121623.1	8	12	10	6	9	45	9	9.0	2.0	20.00%	2.1	24.98%	Shank2
IP00654342.2	10	17	10	8	0	45	10	9.0	4.0	32.77%	5.3	88.19%	Slc7a1
IP00121576.2	7	7	11	11	9	45	9	9.0	2.3	27.71%	1.2	11.17%	Tomm22
IP00626994.3	8	10	14	8	5	45	8	9.0	3.1	28.64%	4.6	50.92%	Txndc5
IP00137194.1	7	8	9	9	12	45	9	9.0	1.0	12.50%	1.7	17.32%	Wdr1
IP00133224.1	7	12	12	10	3	44	10	8.8	2.9	27.94%	4.7	56.71%	2810459M11Rik
IP00652316.4	8	13	6	10	7	44	8	8.8	3.6	40.06%	2.1	27.15%	Acox1
IP00775948.1	13	5	8	10	8	44	8	8.8	4.0	46.63%	1.2	13.32%	Adam23
IP00126716.3	10	12	9	8	5	44	9	8.8	1.5	14.78%	2.1	28.39%	Arvcf
IP00756786.1	9	10	11	8	6	44	9	8.8	1.0	10.00%	2.5	30.20%	Atp9a
IP00132762.1	9	11	9	8	7	44	9	8.8	1.2	11.95%	1.0	12.50%	Calb1
IP00470004.3	11	9	12	9	3	44	9	8.8	1.5	14.32%	4.6	57.28%	Cct8
IP00116112.3	10	9	12	9	4	44	9	8.8	1.5	14.78%	4.0	48.50%	Cd38
IP00266188.6	8	13	8	9	6	44	8	8.8	2.9	29.86%	1.5	19.92%	Coro1c
IP00162949.1	13	5	11	9	6	44	9	8.8	4.2	43.07%	2.5	29.04%	Dnaja2
IP00754880.3	10	11	9	8	6	44	9	8.8	1.0	10.00%	1.5	19.92%	Dnajb6
IP00331552.4	6	10	8	17	3	44	8	8.8	2.0	25.00%	7.1	76.01%	Eif4a3
IP00121038.2	9	11	12	8	4	44	9	8.8	1.5	14.32%	4.0	50.00%	Endod1

IP00467530.3	7	13	10	8	6	44	8	8.8	3.0	30.00%	2.0	25.00%	Fermt2
IP00228343.4	7	15	11	8	3	44	8	8.8	4.0	36.36%	4.0	55.11%	llvbl
IP00410819.3	10	5	11	12	6	44	10	8.8	3.2	37.09%	3.2	33.25%	Klc1
IP00225307.1	7	11	12	6	8	44	8	8.8	2.6	26.46%	3.1	35.25%	Mpp1
IP00230035.7	6	10	10	12	6	44	10	8.8	2.3	26.65%	3.1	32.73%	N28178
IP00120066.5	11	9	6	6	12	44	9	8.8	2.5	29.04%	3.5	43.30%	Pecam1
IP00331163.9	11	13	10	6	4	44	10	8.8	1.5	13.48%	3.1	45.83%	Ptprg
IP00130339.3	10	7	9	11	7	44	9	8.8	1.5	17.63%	2.0	22.22%	Sh3glb2
IP00462251.3	5	9	12	8	10	44	9	8.8	3.5	40.52%	2.0	20.00%	Smpd3
IP00133985.1	11	6	11	8	7	43	8	8.6	2.9	30.93%	2.1	24.02%	Clasp2
IP00387227.1	9	7	11	7	9	43	9	8.6	2.0	22.22%	2.0	22.22%	Daam1
IP00307966.3	3	14	9	8	9	43	9	8.6	5.5	63.55%	0.6	6.66%	Dctn2
IP00119024.3	7	7	9	10	10	43	9	8.6	1.2	15.06%	0.6	5.97%	Fabp5
IP00622837.1	7	11	12	11	2	43	11	8.6	2.6	26.46%	5.5	66.09%	Hnrnpu
IP00265467.3	8	11	12	7	5	43	8	8.6	2.1	20.15%	3.6	45.07%	Lpgat1
IP00313899.4	6	11	9	10	7	43	9	8.6	2.5	29.04%	1.5	17.63%	Ndufa6
IP00124820.2	8	12	8	8	7	43	8	8.6	2.3	24.74%	0.6	7.53%	Ppt1
IP00662157.2	11	12	8	6	6	43	8	8.6	2.1	20.15%	1.2	17.32%	Sbf1
IP00224210.5	6	8	12	9	8	43	8	8.6	3.1	35.25%	2.1	21.53%	Src
IP00323822.1	7	12	7	6	11	43	7	8.6	2.9	33.31%	2.6	33.07%	Vappb
IP00471372.2	10	13	9	6	4	42	9	8.4	2.1	19.52%	2.5	39.74%	0710008K08Rik
IP00276926.3	9	13	6	10	4	42	9	8.4	3.5	37.63%	3.1	45.83%	2310035C23Rik
IP00654178.1	10	8	11	5	8	42	8	8.4	1.5	15.80%	3.0	37.50%	Ace
IP00469268.5	6	10	9	7	10	42	9	8.4	2.1	24.98%	1.5	17.63%	Acol7
IP00330039.3	14	5	9	7	7	42	7	8.4	4.5	48.31%	1.2	15.06%	Cacnb4
IP00129535.1	9	5	7	8	13	42	8	8.4	2.0	28.57%	3.2	34.44%	Ddah1
IP00113377.1	8	13	11	6	4	42	8	8.4	2.5	23.59%	3.6	51.51%	Erp29
IP00857195.1	13	10	9	10	0	42	10	8.4	2.1	19.52%	5.5	86.96%	Fxyd6
IP00331066.6	7	8	9	10	8	42	8	8.4	1.0	12.50%	1.0	11.11%	Ggtl3
IP00224575.1	13	8	8	5	8	42	8	8.4	2.9	29.86%	1.7	24.74%	Gpx1
IP00457485.2	12	7	11	12	0	42	11	8.4	2.6	26.46%	6.7	86.85%	Hnrnpa2b1
IP00118963.1	13	13	13	3	0	42	13	8.4	0.0	0.00%	6.8	127.63%	Krt76
IP00411109.4	10	9	9	8	6	42	9	8.4	0.6	6.19%	1.5	19.92%	Nap111
IP00627008.2	9	8	7	11	7	42	8	8.4	1.0	12.50%	2.3	27.71%	Nf1

IP100658782.4	12	8	9	11	2	42	9	8.4	2.1	21.53%	4.7	64.44%	Pabpc1
IP100265568.6	10	7	5	12	8	42	8	8.4	2.5	34.32%	3.5	42.14%	Pafah1b2
IP100387392.4	9	8	6	11	8	42	8	8.4	1.5	19.92%	2.5	30.20%	Pdxx
IP100134961.1	13	8	9	7	5	42	8	8.4	2.6	26.46%	2.0	28.57%	Prkosh
IP10011501.1	6	13	8	9	6	42	8	8.4	3.6	40.06%	1.5	19.92%	Ptprz1
IP100137618.1	9	9	7	11	6	42	9	8.4	1.2	13.86%	2.6	33.07%	Sars
IP100720013.2	9	10	6	8	9	42	9	8.4	2.1	24.98%	1.5	19.92%	Scamp1
IP100125807.3	11	11	3	14	3	42	11	8.4	4.6	55.43%	6.4	95.26%	Sic23a2
IP100120295.1	13	9	7	8	5	42	8	8.4	3.1	31.60%	1.5	22.91%	Stxbp3a
IP100664670.4	8	15	7	9	3	42	8	8.4	4.4	43.59%	3.1	48.24%	Surf4
IP100331318.2	9	4	7	14	7	41	7	8.2	2.5	37.75%	4.0	43.30%	0910001A06RIK
IP100465999.5	9	0	9	13	10	41	9	8.2	5.2	86.60%	2.1	19.52%	2900073G15RIK
IP100776230.1	13	8	6	6	8	41	8	8.2	3.6	40.06%	1.2	17.32%	Aoc3
IP100130661.1	6	15	8	6	6	41	6	8.2	4.7	48.89%	1.2	17.32%	Atrgdia
IP100474125.3	14	12	5	10	0	41	10	8.2	4.7	45.73%	5.0	100.00%	Alp6ap1
IP100230194.5	11	8	9	11	2	41	9	8.2	1.5	16.37%	4.7	64.44%	Clptm1
IP100555125.1	11	6	10	11	3	41	10	8.2	2.6	29.40%	4.4	54.49%	Ddb1
IP100153642.1	13	12	7	5	4	41	7	8.2	3.2	30.14%	1.5	28.64%	Epb4.1
IP100458870.3	10	10	13	6	2	41	10	8.2	1.7	15.75%	5.6	79.54%	Eprs
IP100113845.1	9	8	6	13	5	41	8	8.2	1.5	19.92%	4.4	54.49%	Gpd11
IP100876001.1	11	11	0	19	0	41	11	8.2	6.4	86.60%	11.0	173.21%	Gstm7
IP100117214.2	9	11	5	8	8	41	8	8.2	3.1	36.66%	1.7	24.74%	Hip1
IP100115949.1	9	4	11	7	10	41	9	8.2	3.6	45.07%	2.1	22.30%	Homer1
IP100407863.5	11	7	8	7	8	41	8	8.2	2.1	24.02%	0.6	7.53%	Hsdl2
IP100221540.1	8	11	8	11	3	41	8	8.2	1.7	19.25%	4.0	55.11%	Htt
IP100118832.1	8	11	8	9	5	41	8	8.2	1.7	19.25%	2.1	28.39%	Igsf21
IP100467338.4	8	9	6	7	11	41	8	8.2	1.5	19.92%	2.6	33.07%	Kalrn
IP100122139.3	12	5	13	8	3	41	8	8.2	4.4	43.59%	5.0	62.50%	Ligl1
IP100465880.4	11	6	6	7	11	41	7	8.2	2.9	37.65%	2.6	33.07%	LOC100044630
IP100857354.1	12	4	11	8	6	41	8	8.2	4.4	48.43%	2.5	30.20%	Mapre2
IP100762435.2	7	11	11	9	3	41	9	8.2	2.3	23.89%	4.2	54.30%	Prdx4
IP100115429.1	12	7	5	9	8	41	8	8.2	3.6	45.07%	2.1	28.39%	Psma2
IP100123534.1	9	11	9	8	4	41	9	8.2	1.2	11.95%	2.6	37.80%	Rhog
IP100459309.2	10	9	7	6	9	41	9	8.2	1.5	17.63%	1.5	20.83%	Sic25a25

IP00163011.2	10	9	9	9	7	6	41	9	8.2	0.6	6.19%	1.5	20.83%	Steap3
IP00121218.5	9	5	10	7	7	10	41	9	8.2	2.6	33.07%	1.7	19.25%	Tmed10
IP00117167.2	11	8	9	8	8	4	40	8	8.0	1.5	16.37%	2.6	37.80%	Ablim1
IP00119130.1	9	7	8	11	5	5	40	8	8.0	1.0	12.50%	3.0	37.50%	Add3
IP00227236.2	8	5	8	9	10	10	40	8	8.0	1.7	24.74%	1.0	11.11%	Caskin1
IP00624098.1	10	7	7	7	3	9	40	7	8.0	1.7	21.65%	1.2	15.06%	Cldn11
IP00129362.1	11	10	9	3	7	7	40	9	8.0	1.0	10.00%	3.1	48.24%	Dagla
IP00321386.2	12	4	7	9	8	8	40	8	8.0	4.0	52.71%	1.0	12.50%	Erc2
IP00850694.2	8	12	11	4	5	5	40	8	8.0	2.1	20.15%	3.8	56.79%	Fech
IP00762862.3	10	8	3	9	9	10	40	9	8.0	3.6	51.51%	3.8	51.63%	Mfn2
IP00172234.3	10	10	9	9	2	2	40	9	8.0	0.6	5.97%	4.0	60.62%	Nol5
IP00283511.1	11	0	5	7	17	17	40	7	8.0	5.5	103.27%	6.4	66.51%	Pea15a
IP00649277.2	15	8	6	6	5	5	40	6	8.0	4.7	48.89%	0.6	10.19%	Ppp2f5c
IP00170387.1	5	12	8	15	0	0	40	8	8.0	3.5	42.14%	7.5	97.90%	Rangap1
IP00133218.3	7	4	9	10	10	10	40	9	8.0	2.5	37.75%	0.6	5.97%	Sez6l2
IP00336807.4	7	11	6	6	5	8	40	7	8.0	2.0	22.22%	2.1	28.39%	Srgap2
IP00761259.1	12	7	6	6	8	8	39	7	7.8	3.2	38.57%	1.2	17.32%	2310014G06Rik
IP00128760.1	10	7	6	9	7	7	39	7	7.8	2.1	27.15%	1.5	20.83%	Coro2b
IP00108150.1	9	9	6	9	6	6	39	9	7.8	1.7	21.65%	1.7	24.74%	Dclk1
IP00121105.2	6	7	13	13	6	0	39	7	7.8	3.8	43.68%	7.5	86.60%	EG331392
IP00111271.1	11	8	11	6	3	3	39	8	7.8	1.7	17.32%	4.0	60.62%	Gsn
IP00331092.7	9	6	12	7	5	5	39	7	7.8	3.0	33.33%	3.6	45.07%	Madd
IP00135186.1	11	8	7	8	8	5	39	8	7.8	2.1	24.02%	1.5	22.91%	Me1
IP00624843.3	9	9	6	7	7	8	39	8	7.8	1.7	21.65%	1.0	14.29%	Mirps36
IP00828221.2	5	9	10	11	11	4	39	9	7.8	2.6	33.07%	3.8	45.43%	Nup160
IP00652694.1	10	11	11	5	2	2	39	10	7.8	0.6	5.41%	4.6	76.38%	Pigs
IP00132080.1	8	12	11	4	4	4	39	8	7.8	2.1	20.15%	4.0	63.81%	Ppif
IP00270877.5	7	10	10	7	5	5	39	7	7.8	1.7	19.25%	2.5	34.32%	Psemb1
IP00875943.1	9	10	10	7	3	3	39	9	7.8	0.6	5.97%	3.5	52.68%	Psm7
IP00126548.2	12	9	6	9	3	3	39	9	7.8	3.0	33.33%	3.0	50.00%	Sec22b
IP00785449.1	12	8	7	9	3	3	39	8	7.8	2.6	29.40%	3.1	48.24%	Tspan2
IP00849793.1	9	5	10	7	7	7	38	7	7.6	2.6	33.07%	1.7	21.65%	Acadm
IP00343557.1	9	6	10	2	11	11	38	9	7.6	2.1	24.98%	4.9	64.34%	Adam10
IP00420734.1	6	8	8	10	6	6	38	8	7.6	1.2	15.75%	2.0	25.00%	Cct6a

IP100830314.1	8	9	6	8	7	38	8	7.6	1.5	19.92%	1.0	14.29%	Cpne5
IP100128341.1	9	10	4	9	6	38	9	7.6	3.2	41.93%	2.5	39.74%	Cyb5r1
IP100315280.5	6	11	10	6	5	38	6	7.6	2.6	29.40%	2.6	37.80%	Faah
IP100113539.2	9	6	9	9	5	38	9	7.6	1.7	21.65%	2.3	30.12%	Fgg
IP100881401.1	8	10	7	6	7	38	7	7.6	1.5	18.33%	0.6	8.66%	Folh1
IP100322145.10	7	8	6	12	5	38	7	7.6	1.0	14.29%	3.8	49.38%	Hkdc1
IP100356904.7	7	4	10	7	10	38	7	7.6	3.0	42.86%	1.7	19.25%	Lin7c
IP100469918.4	11	10	6	6	5	38	6	7.6	2.6	29.40%	0.6	10.19%	Ociad1
IP100880948.1	7	7	7	8	9	38	7	7.6	0.0	0.00%	1.0	12.50%	Peci
IP100415914.3	5	6	10	10	7	38	7	7.6	2.6	37.80%	1.7	19.25%	Prkar1a
IP100331463.1	9	9	11	7	2	38	9	7.6	1.2	11.95%	4.5	67.64%	Sic25a46
IP100323571.1	11	9	6	7	5	38	7	7.6	2.5	29.04%	1.0	16.67%	Sor11
IP100228253.2	8	9	9	9	5	38	8	7.6	0.6	6.66%	2.0	28.57%	Sparc1
IP100114162.3	8	6	9	10	5	38	8	7.6	1.5	19.92%	2.6	33.07%	Tpm1
IP100123557.3	8	9	4	11	6	38	8	7.6	2.6	37.80%	3.6	51.51%	Ube2l3
IP100114368.3	9	3	4	8	14	38	8	7.6	3.2	60.27%	5.0	58.08%	Ube2h
IP100116281.3	10	4	9	8	6	37	8	7.4	3.2	41.93%	1.5	19.92%	Arhgef12
IP100120199.1	10	6	12	5	4	37	6	7.4	3.1	32.73%	4.4	62.27%	Capn2
IP100469338.4	4	12	7	9	5	37	7	7.4	4.0	52.71%	2.0	28.57%	Efr3b
IP100128945.1	13	5	4	10	5	37	5	7.4	4.9	67.27%	3.2	50.76%	Fkbp2
IP100869381.1	7	12	5	6	7	37	7	7.4	3.6	45.07%	1.0	16.67%	Grhpr
IP100331146.5	6	8	13	8	2	37	7	7.4	3.6	40.06%	5.5	71.84%	LOC100048622
IP100225575.5	10	7	4	9	7	37	7	7.4	3.0	42.86%	2.5	37.75%	Mccc1
IP100466185.4	9	11	5	6	6	37	6	7.4	3.1	36.66%	0.6	10.19%	Mccc2
IP100109033.3	8	10	7	6	6	37	7	7.4	1.5	18.33%	0.6	9.12%	Necap1
IP100276222.1	5	10	3	11	8	37	8	7.4	3.6	60.09%	4.0	55.11%	Nlgn3
IP100117986.2	9	11	5	5	7	37	7	7.4	3.1	36.66%	1.2	20.38%	Nucb1
IP100648709.2	14	3	3	6	11	37	6	7.4	6.4	95.26%	4.0	60.62%	Pak1
IP100318841.4	7	5	7	9	9	37	7	7.4	1.2	18.23%	1.2	13.86%	Pin1
IP100128436.1	7	7	6	9	8	37	7	7.4	0.6	8.66%	1.5	19.92%	Ppp2r5e
IP100114377.1	9	5	8	9	6	37	8	7.4	2.1	28.39%	1.5	19.92%	Ptprn2
IP100754572.2	4	13	8	5	7	37	7	7.4	4.5	54.11%	1.5	22.91%	Rab5b
IP100271166.4	0	16	8	13	0	37	8	7.4	8.0	100.00%	6.6	93.68%	Rac2
IP100380814.1	9	8	6	7	7	37	7	7.4	1.5	19.92%	0.6	8.66%	Sic25a1

IP100420394.4	8	7	7	7	7	7	7	8	37	7	7.4	0.6	7.87%	0.6	7.87%	0.6	7.87%	Synpo
IP100109588.4	8	9	9	7	7	4	37	8	37	8	7.4	0.6	6.66%	2.5	37.75%	2.5	37.75%	Thnsl1
IP100125521.3	10	3	3	7	14	7	37	7	37	7	7.4	4.0	75.78%	5.6	69.60%	5.6	69.60%	Tpp1
IP100126796.3	9	3	10	8	7	4	37	8	37	8	7.4	3.8	51.63%	1.5	18.33%	1.5	18.33%	Vps13a
IP100673886.4	7	13	5	8	4	4	37	7	37	7	7.4	4.2	49.96%	2.1	36.74%	2.1	36.74%	Wdr37
IP100408646.5	8	8	9	7	4	8	36	8	36	8	7.2	0.6	6.93%	2.5	37.75%	2.5	37.75%	1500005102Rik
IP100762919.2	6	9	10	3	8	8	36	8	36	8	7.2	2.1	24.98%	3.6	51.51%	3.6	51.51%	Adcy9
IP100653300.1	9	7	5	9	6	6	36	7	36	7	7.2	2.0	28.57%	2.1	31.22%	2.1	31.22%	Asrgl1
IP100153903.6	10	9	8	6	3	3	36	7	36	7	7.2	1.0	11.11%	2.5	44.41%	2.5	44.41%	Cox7a2
IP100225533.1	6	14	9	7	0	8	36	7	36	7	7.2	4.0	41.81%	4.7	88.61%	4.7	88.61%	Ddx5
IP100466128.3	12	3	6	7	8	8	36	7	36	7	7.2	4.6	65.47%	1.0	14.29%	1.0	14.29%	Epha4
IP100762185.2	9	8	9	10	0	0	36	9	36	9	7.2	0.6	6.66%	5.5	86.96%	5.5	86.96%	Fn1
IP100420931.1	8	6	8	5	9	9	36	8	36	8	7.2	1.2	15.75%	2.1	28.39%	2.1	28.39%	Grid2
IP100330363.8	8	4	11	6	7	7	36	7	36	7	7.2	3.5	45.81%	2.6	33.07%	2.6	33.07%	Krt1
IP100420363.2	10	6	8	5	7	7	36	7	36	7	7.2	2.0	25.00%	1.5	22.91%	1.5	22.91%	Ktn1
IP100420467.1	12	6	4	9	5	5	36	6	36	6	7.2	4.2	56.77%	2.6	44.10%	2.6	44.10%	Odz4
IP100625729.2	8	10	8	7	3	3	36	8	36	8	7.2	1.2	13.32%	2.6	44.10%	2.6	44.10%	Pitpna
IP100857742.1	6	4	11	9	6	6	36	6	36	6	7.2	3.6	51.51%	2.5	29.04%	2.5	29.04%	Pthr2
IP100308187.1	13	3	6	9	5	5	36	6	36	6	7.2	5.1	69.98%	2.1	31.22%	2.1	31.22%	Scn1
IP100411157.4	7	11	7	8	3	3	36	7	36	7	7.2	2.3	27.71%	2.6	44.10%	2.6	44.10%	Tmem33
IP100118022.1	7	8	8	6	7	7	36	7	36	7	7.2	0.6	7.53%	1.0	14.29%	1.0	14.29%	Ubr4
IP100648288.1	6	6	10	8	6	6	36	6	36	6	7.2	2.3	31.49%	2.0	25.00%	2.0	25.00%	Vgf
IP100750536.1	8	11	9	6	2	2	36	8	36	8	7.2	1.5	16.37%	3.5	61.97%	3.5	61.97%	Xpo1
IP100221725.2	7	8	7	8	5	5	35	7	35	7	7.0	0.6	7.87%	1.5	22.91%	1.5	22.91%	Ap1g1
IP100515370.1	8	9	6	5	7	7	35	7	35	7	7.0	1.5	19.92%	1.0	16.67%	1.0	16.67%	Apoa1bp
IP100330193.3	13	6	5	8	3	3	35	6	35	6	7.0	4.4	54.49%	2.5	47.19%	2.5	47.19%	Cacna2d3
IP100115626.1	9	5	5	11	5	5	35	5	35	5	7.0	2.3	36.46%	3.5	49.49%	3.5	49.49%	Cap2
IP100318496.1	8	4	9	9	5	5	35	8	35	8	7.0	2.6	37.80%	2.3	30.12%	2.3	30.12%	Cfl2
IP100466610.5	11	6	2	8	8	8	35	8	35	8	7.0	4.5	71.20%	3.5	57.74%	3.5	57.74%	Chl1
IP100466919.7	9	6	12	4	4	4	35	6	35	6	7.0	3.0	33.33%	4.6	69.28%	4.6	69.28%	Cln6
IP100828479.1	6	5	6	10	8	8	35	6	35	6	7.0	0.6	10.19%	2.0	25.00%	2.0	25.00%	Clic6
IP100318522.3	6	6	9	5	9	9	35	6	35	6	7.0	1.7	24.74%	2.3	30.12%	2.3	30.12%	Eif4a1
IP100653643.3	13	6	6	3	7	7	35	6	35	6	7.0	4.0	48.50%	2.1	39.03%	2.1	39.03%	Enpp1
IP100153702.2	6	11	11	5	2	2	35	6	35	6	7.0	2.9	30.93%	4.6	76.38%	4.6	76.38%	Emp1

IP100230084.5	11	3	9	5	7	35	7	7.0	4.2	54.30%	2.0	28.57%	Gnb211
IP001122559.3	12	5	7	7	4	35	7	7.0	3.6	45.07%	1.7	28.87%	Gbn
IP001133249.1	4	8	10	7	6	35	7	7.0	3.1	41.66%	2.1	27.15%	Gstk1
IP00118447.1	6	4	14	4	7	35	6	7.0	5.3	66.14%	5.1	61.58%	Hin2
IP001128632.6	7	13	10	5	0	35	7	7.0	3.0	30.00%	5.0	100.00%	Krt14
IP001133234.1	8	7	6	9	5	35	7	7.0	1.0	14.29%	2.1	31.22%	Lxn
IP001131143.1	10	8	7	5	5	35	7	7.0	1.5	18.33%	1.2	20.38%	Mcart1
IP00221739.1	7	5	7	11	5	35	7	7.0	1.2	18.23%	3.1	39.85%	Mecp2
IP001122396.1	6	4	8	11	6	35	6	7.0	2.0	33.33%	2.5	30.20%	Ndufb11
IP001131204.1	3	10	6	7	9	35	7	7.0	3.5	55.45%	1.5	20.83%	Ndufb6
IP00317740.5	7	9	7	8	4	35	7	7.0	1.2	15.06%	2.1	32.87%	Padi2
IP00420806.3	8	7	7	6	7	35	7	7.0	0.6	7.87%	0.6	8.66%	Pck2
IP00223274.2	11	8	4	6	6	35	6	7.0	3.5	45.81%	1.2	21.65%	Pcmt1
IP00321308.4	7	9	8	6	5	35	7	7.0	1.0	12.50%	1.5	24.12%	Prom1
IP00407222.4	5	13	8	6	3	35	6	7.0	4.0	46.63%	2.5	44.41%	Rps15
IP00881758.1	4	13	9	4	5	35	5	7.0	4.5	52.03%	2.6	44.10%	Tjp2
IP001153950.1	6	10	9	5	5	35	6	7.0	2.1	24.98%	2.3	36.46%	Tst
IP001122312.2	8	3	5	10	9	35	8	7.0	2.5	47.19%	2.6	33.07%	Zadh2
IP00115977.1	7	4	10	3	10	34	7	6.8	3.0	42.86%	4.0	52.71%	2210010N04Rik
IP00652758.1	4	10	10	6	4	34	6	6.8	3.5	43.30%	3.1	45.83%	Akr7a5
IP00775806.2	9	8	7	3	7	34	7	6.8	1.0	12.50%	2.3	40.75%	Aldh7a1
IP00116563.2	7	5	6	11	5	34	6	6.8	1.0	16.67%	3.2	43.83%	Ap3b2
IP00608114.2	7	15	7	3	2	34	7	6.8	4.6	47.78%	2.6	66.14%	Calu
IP00338785.3	9	6	8	8	3	34	8	6.8	1.5	19.92%	2.9	45.58%	Cmpk
IP00420426.1	9	7	9	6	3	34	7	6.8	1.2	13.86%	3.0	50.00%	Cpt1a
IP001124286.2	7	9	6	5	7	34	7	6.8	1.5	20.83%	1.0	16.67%	EG545878
IP00330599.1	9	9	7	9	0	34	9	6.8	1.2	13.86%	4.7	88.61%	Ehd4
IP00222759.3	6	7	9	5	7	34	7	6.8	1.5	20.83%	2.0	28.57%	Eif4a2
IP00118880.5	7	11	8	6	2	34	7	6.8	2.1	24.02%	3.1	57.28%	Ggt1
IP001128859.2	13	4	4	6	7	34	6	6.8	5.2	74.23%	1.5	26.96%	Lmf2
IP00270326.1	10	5	5	9	5	34	5	6.8	2.9	43.30%	2.3	36.46%	Pls3
IP001123762.2	10	6	5	4	9	34	6	6.8	2.6	37.80%	2.6	44.10%	Ppap2b
IP001130530.1	11	4	7	5	7	34	7	6.8	3.5	47.89%	1.2	18.23%	Psmd1
IP00226872.1	7	7	4	9	7	34	7	6.8	1.7	28.87%	2.5	37.75%	Ptpn5



IP00136207.6	10	8	9	4	3	34	8	6.8	1.0	11.11%	3.2	60.27%	Pura
IP00403682.2	6	10	8	10	0	34	8	6.8	2.0	25.00%	5.3	88.19%	Rpl7a
IP00881197.1	5	11	8	8	2	34	8	6.8	3.0	37.50%	3.5	57.74%	Rps17
IP00131895.4	13	6	6	6	3	34	6	6.8	4.0	48.50%	1.7	34.64%	Sel1l
IP00223527.1	10	4	7	6	7	34	7	6.8	3.0	42.86%	0.6	8.66%	Sema7a
IP00318671.2	2	7	9	7	9	34	7	6.8	3.6	60.09%	1.2	13.86%	Skp1a
IP00457659.2	5	8	6	4	11	34	6	6.8	1.5	24.12%	3.6	51.51%	Slc1a1
IP00874376.1	7	9	6	5	7	34	7	6.8	1.5	20.83%	1.0	16.67%	Tom1
IP00132734.1	0	6	12	12	4	34	6	6.8	6.0	100.00%	4.6	49.49%	Tpm3
IP00399905.5	8	10	4	8	4	34	8	6.8	3.1	41.66%	2.3	43.30%	Usp14
IP00169622.3	9	6	11	3	4	33	6	6.6	2.5	29.04%	4.4	72.65%	Abi1
IP00330464.5	6	10	5	5	7	33	6	6.6	2.6	37.80%	1.2	20.38%	Agpat1
IP00125180.9	7	7	3	10	6	33	7	6.6	2.3	40.75%	3.5	55.45%	Al427515
IP00221828.1	5	11	6	8	3	33	6	6.6	3.2	43.83%	2.5	44.41%	Aldh18a1
IP00118795.1	9	7	8	3	6	33	7	6.6	1.0	12.50%	2.5	44.41%	Ahrgef2
IP00752390.3	6	3	2	15	7	33	6	6.6	2.1	56.77%	6.6	81.97%	Cd9
IP00124051.1	6	13	8	6	0	33	6	6.6	3.6	40.06%	4.2	89.21%	Dcakd
IP00138217.2	12	8	7	2	4	33	7	6.6	2.6	29.40%	2.5	58.08%	Drp2
IP00113521.5	7	7	7	5	7	33	7	6.6	0.0	0.00%	1.2	18.23%	Dync1l12
IP00165854.3	8	3	6	7	9	33	7	6.6	2.5	44.41%	1.5	20.83%	Efhf2
IP00653706.1	8	5	4	7	9	33	7	6.6	2.1	36.74%	2.5	37.75%	Freq
IP00776324.1	7	6	6	9	5	33	6	6.6	0.6	9.12%	2.1	31.22%	Git1
IP00130546.1	8	6	7	12	0	33	7	6.6	1.0	14.29%	6.0	95.17%	Gnb5
IP00134058.3	7	6	6	9	5	33	6	6.6	0.6	9.12%	2.1	31.22%	Hist1h1b
IP00336313.2	7	10	12	4	0	33	7	6.6	2.5	26.03%	6.1	114.56%	Inf2
IP00380354.1	7	7	9	2	8	33	7	6.6	1.2	15.06%	3.8	59.78%	Lphn3
IP00132696.2	14	7	6	4	2	33	6	6.6	4.4	48.43%	2.0	50.00%	Lrrc8a
IP00221857.1	7	8	12	4	2	33	7	6.6	2.6	29.40%	5.3	88.19%	Mtdh
IP00378764.6	9	8	7	4	5	33	7	6.6	1.0	12.50%	1.5	28.64%	Pgrmc2
IP00127930.1	12	2	7	5	7	33	7	6.6	5.0	71.43%	1.2	18.23%	Pitrm1
IP00330857.5	8	8	8	5	4	33	8	6.6	0.0	0.00%	2.1	36.74%	Rgs6
IP00625848.2	8	3	14	3	5	33	5	6.6	5.5	66.09%	5.9	79.90%	Sic30a1
IP00269408.1	5	6	10	7	5	33	6	6.6	2.6	37.80%	2.5	34.32%	Sorbs2
IP00121051.3	7	7	4	8	7	33	7	6.6	1.7	28.87%	2.1	32.87%	Wasf1

IP100341322.3	7	9	7	7	2	32	7	6.4	1.2	15.06%	2.9	54.13%	A330104H05Rik
IP100319652.1	6	4	8	5	9	32	6	6.4	2.0	33.33%	2.1	28.39%	Abr
IP100129350.1	7	7	5	7	6	32	7	6.4	1.2	18.23%	1.0	16.67%	Acat2
IP100380338.1	6	5	8	5	8	32	6	6.4	1.5	24.12%	1.7	24.74%	Aldh3a2
IP100408059.4	6	7	4	5	10	32	6	6.4	1.5	26.96%	3.2	50.76%	Ap3m2
IP100223070.3	6	9	8	6	3	32	6	6.4	1.5	19.92%	2.5	44.41%	Apool
IP100468685.2	9	7	6	7	3	32	7	6.4	1.5	20.83%	2.1	39.03%	Cacna1a
IP100330941.3	4	3	5	12	8	32	5	6.4	1.0	25.00%	3.5	42.14%	Csrp1
IP100466570.4	4	3	5	12	8	32	5	6.4	1.0	25.00%	3.5	42.14%	Ctnna3
IP100330094.4	5	7	8	7	5	32	7	6.4	1.5	22.91%	1.5	22.91%	D1Erd53e
IP100319231.4	10	4	5	8	5	32	5	6.4	3.2	50.76%	1.7	28.87%	Dlgap3
IP100109482.3	7	2	10	8	5	32	7	6.4	4.0	63.81%	2.5	32.83%	Fads2
IP100798483.1	8	5	8	5	6	32	6	6.4	1.7	24.74%	1.5	24.12%	Fcgrt
IP100130102.4	5	4	8	8	7	32	7	6.4	2.1	36.74%	0.6	7.53%	Gabrb3
IP100403618.4	5	11	6	5	5	32	5	6.4	3.2	43.83%	0.6	10.83%	Gng12
IP100279010.6	10	7	5	6	4	32	6	6.4	2.5	34.32%	1.0	20.00%	Gucy1b3
IP100122522.1	3	10	5	7	7	32	7	6.4	3.6	60.09%	1.2	18.23%	Hebp1
IP100132314.1	0	16	9	0	7	32	7	6.4	8.0	96.25%	4.7	88.61%	HK2
IP100225534.3	8	9	8	5	2	32	8	6.4	0.6	6.93%	3.0	60.00%	Hnrpk
IP100124829.5	6	5	9	7	5	32	6	6.4	2.1	31.22%	2.0	28.57%	Konc1
IP100135068.1	6	9	5	9	3	32	6	6.4	2.1	31.22%	3.1	53.91%	Kond2
IP100118676.3	10	5	6	6	5	32	6	6.4	2.6	37.80%	0.6	10.19%	Lman2
IP100624175.1	9	11	8	4	0	32	8	6.4	1.5	16.37%	4.0	100.00%	LOC672959
IP100408796.3	8	7	5	4	8	32	7	6.4	1.5	22.91%	2.1	36.74%	Mink1
IP100153381.1	8	8	6	5	5	32	6	6.4	1.2	15.75%	0.6	10.83%	Plcb4
IP100170126.4	11	7	7	0	7	32	7	6.4	2.3	27.71%	4.0	86.60%	Ppp2r2d
IP100124640.4	8	5	4	7	8	32	7	6.4	2.1	36.74%	2.1	32.87%	Ran
IP100310247.7	5	11	5	8	3	32	5	6.4	3.5	49.49%	2.5	47.19%	Reep5
IP100854952.1	8	7	5	9	3	32	7	6.4	1.5	22.91%	3.1	53.91%	Rock2
IP100469942.2	11	9	8	4	0	32	8	6.4	1.5	16.37%	4.0	100.00%	Rps5
IP100461626.1	6	10	9	4	3	32	6	6.4	2.1	24.98%	3.2	60.27%	Sec63
IP100351206.5	10	7	7	8	0	32	7	6.4	1.7	21.65%	4.4	87.18%	Tm9sf2
IP100114945.1	6	2	7	15	2	32	6	6.4	2.6	52.92%	6.6	81.97%	Tmpo
IP100622240.4	7	9	6	6	4	32	6	6.4	1.5	20.83%	1.2	21.65%	Ubqln1

IP00133403.1	10	8	11	3	0	32	8	6.4	1.5	15.80%	5.7	121.85%	Ugp2
IP00308149.1	11	5	7	6	2	31	6	6.2	3.1	39.85%	2.6	52.92%	Adcy6
IP00347634.2	4	10	7	4	6	31	6	6.2	3.0	42.86%	1.5	26.96%	Apoa2
IP00227140.1	8	5	7	4	7	31	7	6.2	1.5	22.91%	1.7	28.87%	Arpc4
IP00409336.4	9	9	5	6	2	31	6	6.2	2.3	30.12%	2.1	48.04%	Cbara1
IP00114962.1	13	3	4	5	6	31	5	6.2	5.5	82.61%	1.0	20.00%	Cdc42bpb
IP00395038.3	5	4	9	6	7	31	6	6.2	2.6	44.10%	1.5	20.83%	Clybl
IP00120083.1	9	5	4	5	8	31	5	6.2	2.6	44.10%	2.1	36.74%	Cnksr2
IP00473728.1	2	11	10	3	5	31	5	6.2	4.9	64.34%	3.6	60.09%	Comtd1
IP00153227.5	9	7	6	5	4	31	6	6.2	1.5	20.83%	1.0	20.00%	Cpm
IP00115231.2	13	4	4	4	6	31	4	6.2	5.2	74.23%	1.2	24.74%	ENSMUSG00000073431
IP00114389.4	5	8	9	3	6	31	6	6.2	2.1	28.39%	3.0	50.00%	Hexb
IP00649763.1	5	11	7	6	2	31	6	6.2	3.1	39.85%	2.6	52.92%	Hnrnp1
IP00830701.1	3	5	10	6	7	31	6	6.2	3.6	60.09%	2.1	27.15%	Hprt1
IP00409918.1	6	5	4	9	7	31	6	6.2	1.0	20.00%	2.5	37.75%	Itns1
IP00380331.1	6	6	11	5	3	31	6	6.2	2.9	37.65%	4.2	65.74%	LOC100044842
IP00322312.3	6	7	5	7	6	31	6	6.2	1.0	16.67%	1.0	16.67%	M6pr
IP00123014.1	8	8	8	7	0	31	8	6.2	0.0	0.00%	4.4	87.18%	Mipol1
IP00420244.3	7	7	7	5	5	31	7	6.2	0.0	0.00%	1.2	20.38%	Ociad2
IP00223060.1	8	8	8	7	0	31	8	6.2	0.0	0.00%	4.4	87.18%	Prps1
IP00122015.1	6	3	11	7	4	31	6	6.2	4.0	60.62%	3.5	47.89%	Psmc2
IP00315593.4	10	7	2	6	6	31	6	6.2	4.0	63.81%	2.3	49.49%	Psmc3
IP00420651.3	7	8	12	4	0	31	7	6.2	2.6	29.40%	6.1	114.56%	Rasip1
IP00120715.1	6	7	8	7	3	31	7	6.2	1.0	14.29%	2.6	44.10%	Serpib1a
IP00272690.2	7	0	7	9	8	31	7	6.2	4.0	86.60%	1.0	12.50%	Trim2
IP00137173.1	8	7	10	3	3	31	7	6.2	1.5	18.33%	4.0	75.78%	Txndc4
IP00222515.5	8	4	9	5	5	31	5	6.2	2.6	37.80%	2.3	36.46%	Ube2o
IP00751337.3	9	4	7	4	6	30	6	6.0	2.5	37.75%	1.5	26.96%	Abi2
IP00170307.1	6	3	8	6	7	30	6	6.0	2.5	44.41%	1.0	14.29%	Apoe
IP00135085.4	5	6	4	8	7	30	6	6.0	1.0	20.00%	2.1	32.87%	Bcan
IP00110049.5	6	8	5	6	5	30	6	6.0	1.5	24.12%	0.6	10.83%	Calb2
IP00135640.1	9	6	8	3	4	30	6	6.0	1.5	19.92%	2.6	52.92%	Ckap5
IP00266836.1	4	10	5	6	5	30	5	6.0	3.2	50.76%	0.6	10.83%	Cpne4
IP00108980.2	9	7	3	6	5	30	6	6.0	3.1	48.24%	1.5	32.73%	D430041D05Rik

IP100122862.4	10	4	10	3	3	30	4	6.0	3.5	43.30%	4.0	75.78%	Efr3a
IP100130804.1	9	4	6	6	5	30	6	6.0	2.5	39.74%	0.6	10.19%	Entpd2
IP100851092.1	6	4	9	8	3	30	6	6.0	2.5	39.74%	3.2	48.22%	Ephb2
IP100114342.1	5	6	5	7	7	30	6	6.0	0.6	10.83%	1.2	18.23%	Gpr3711
IP100224151.5	9	0	9	7	5	30	7	6.0	5.2	86.60%	2.0	28.57%	H2-Q1
IP100128201.1	17	0	13	0	0	30	0	6.0	8.9	88.88%	7.5	173.21%	Hist1H1c
IP100136498.1	5	7	11	7	0	30	7	6.0	3.1	39.85%	5.6	92.80%	Irgm
IP100381365.1	4	7	6	8	5	30	6	6.0	1.5	26.96%	1.5	24.12%	Pdk1
IP100221569.1	6	2	7	6	9	30	6	6.0	2.6	52.92%	1.5	20.83%	Pdcp
IP100377680.4	9	5	5	5	6	30	5	6.0	2.3	36.46%	0.6	10.83%	Plxnd1
IP100117454.1	7	6	7	6	4	30	6	6.0	0.6	8.66%	1.5	26.96%	Psmid1
IP100131870.3	10	5	5	10	0	30	5	6.0	2.9	43.30%	5.0	100.00%	Rasa3
IP100776023.1	8	4	4	10	4	30	4	6.0	2.3	43.30%	3.5	57.74%	Sdhc
IP100876558.1	12	10	4	4	0	30	4	6.0	4.2	48.04%	2.3	86.60%	Slc44a2
IP100653535.2	9	5	6	4	6	30	6	6.0	2.1	31.22%	1.2	21.65%	Timm44
IP100153421.1	3	0	11	14	2	30	3	6.0	5.7	121.85%	6.2	69.39%	Tpr
IP100828513.1	7	7	9	2	4	29	7	5.8	1.2	15.06%	3.6	72.11%	Akr1a4
IP100816935.3	6	9	8	4	2	29	6	5.8	1.5	19.92%	3.1	65.47%	App
IP100312507.2	11	3	2	0	13	29	3	5.8	4.9	92.49%	7.0	140.00%	Atf8a
IP100121319.1	6	12	4	3	4	29	4	5.8	4.2	56.77%	0.6	15.75%	Armc10
IP100230133.5	5	8	6	3	7	29	6	5.8	1.5	24.12%	2.1	39.03%	Ech1
IP100138691.6	5	3	7	4	10	29	5	5.8	2.0	40.00%	3.0	42.86%	ENSMUSG00000071280
IP100471022.3	9	3	4	11	2	29	4	5.8	3.2	60.27%	4.7	83.40%	Gak
IP100126338.5	7	5	9	3	5	29	5	5.8	2.0	28.57%	3.1	53.91%	Hax1
IP100229834.2	8	10	7	4	0	29	7	5.8	1.5	18.33%	3.5	95.78%	Kir77
IP100111117.5	8	3	4	6	8	29	6	5.8	2.6	52.92%	2.0	33.33%	Mpp5
IP100137792.1	0	0	0	29	0	29	0	5.8	0.0	#DIV/0!	16.7	173.21%	Myh4
IP100655156.3	10	5	6	5	3	29	5	5.8	2.6	37.80%	1.5	32.73%	Nap114
IP100284806.8	6	7	6	8	2	29	6	5.8	0.6	9.12%	3.1	57.28%	Nars
IP100128567.2	10	3	7	4	5	29	5	5.8	3.5	52.68%	1.5	28.64%	Ncln
IP100131406.1	5	8	9	3	4	29	5	5.8	2.1	28.39%	3.2	60.27%	Ospl8
IP100654967.3	2	9	8	6	4	29	6	5.8	3.8	59.78%	2.0	33.33%	Paics
IP100133428.3	8	5	4	8	4	29	5	5.8	2.1	36.74%	2.3	43.30%	Ppfia3
IP100130253.4	6	5	6	7	5	29	6	5.8	0.6	10.19%	1.0	16.67%	Ppp2r4

IP100131871.1	7	8	5	9	0	29	7	5.8	1.5	22.91%	4.5	96.63%	Psmb4
IP100312374.6	8	5	8	5	3	29	5	5.8	1.7	24.74%	2.5	47.19%	Psmc4
IP100756908.2	0	0	15	14	0	29	0	5.8	8.7	173.21%	8.4	86.76%	Rhoc
IP100125652.1	7	5	10	5	2	29	5	5.8	2.5	34.32%	4.0	71.32%	Rpl10a
IP100224093.2	8	10	9	2	0	29	8	5.8	1.0	11.11%	4.7	128.89%	Rps4x
IP100453829.3	8	4	8	4	5	29	5	5.8	2.3	34.64%	2.1	36.74%	Sic12a4
IP100853981.1	11	7	4	5	2	29	5	5.8	3.5	47.89%	1.5	41.66%	Sic44a1
IP100112139.1	6	7	7	7	2	29	7	5.8	0.6	8.66%	2.9	54.13%	Tbc1d24
IP100230766.4	9	5	7	4	4	29	5	5.8	2.0	28.57%	1.7	34.64%	Tenc1
IP100222852.3	5	10	8	2	4	29	5	5.8	2.5	32.83%	3.1	65.47%	Timm50
IP100330634.1	11	6	5	4	3	29	5	5.8	3.2	43.83%	1.0	25.00%	Tmem30a
IP100224070.2	8	5	5	9	2	29	5	5.8	1.7	28.87%	3.5	65.85%	Wars
IP100128285.1	4	8	4	7	5	28	5	5.6	2.3	43.30%	1.5	28.64%	1110020P15RIK
IP100331745.2	4	6	9	2	7	28	6	5.6	2.5	39.74%	3.6	60.09%	1300010F03RIK
IP100551236.3	6	9	4	4	5	28	5	5.6	2.5	39.74%	0.6	13.32%	Adcy2
IP100338327.1	7	6	10	2	3	28	6	5.6	2.1	27.15%	4.4	87.18%	Arpc5l
IP100230171.1	6	5	7	7	3	28	6	5.6	1.0	16.67%	2.3	40.75%	Atp11c
IP100857720.1	8	3	6	6	5	28	6	5.6	2.5	44.41%	0.6	10.19%	Csnk2a2
IP100112377.1	5	9	10	4	0	28	5	5.6	2.6	33.07%	5.0	107.85%	Eif3a
IP100387377.1	5	13	10	0	0	28	5	5.6	4.0	43.30%	5.8	173.21%	Eif3f
IP100337992.1	4	4	4	7	9	28	4	5.6	0.0	0.00%	2.5	37.75%	Etl4
IP100120113.5	15	3	3	7	0	28	3	5.6	6.9	98.97%	3.5	105.36%	Hsd17b11
IP100310764.7	5	2	7	8	6	28	6	5.6	2.5	53.93%	1.0	14.29%	Inpp5a
IP100460653.2	6	3	6	6	7	28	6	5.6	1.7	34.64%	0.6	9.12%	Kcnma1
IP100132993.1	11	5	6	6	0	28	6	5.6	3.2	43.83%	3.5	86.60%	Krt2
IP100269661.1	6	6	10	6	0	28	6	5.6	2.3	31.49%	5.0	94.37%	Lin7a
IP100831445.1	8	2	4	4	10	28	4	5.6	3.1	65.47%	3.5	57.74%	Ndrp3
IP100225634.6	3	6	9	2	8	28	6	5.6	3.0	50.00%	3.8	59.78%	Nt5c
IP100127495.3	6	3	5	7	7	28	6	5.6	1.5	32.73%	1.2	18.23%	Nup210
IP100869394.1	7	3	6	9	3	28	6	5.6	2.1	39.03%	3.0	50.00%	Pcbp1
IP100649005.1	4	4	5	8	7	28	5	5.6	0.6	13.32%	1.5	22.91%	Pcdh1
IP100108844.1	7	6	6	7	2	28	6	5.6	0.6	9.12%	2.6	52.92%	Pip4k2a
IP100653079.2	9	2	4	6	7	28	6	5.6	3.6	72.11%	1.5	26.96%	Pmpca
IP100119346.1	8	2	4	6	8	28	6	5.6	3.1	65.47%	2.0	33.33%	Pripsap2

PI00316650.1	8	6	5	4	5	28	5	5.6	1.5	24.12%	0.6	12.37%	Psmc5
PI00877214.1	6	7	8	7	0	28	7	5.6	1.0	14.29%	4.4	87.18%	Rplp1
PI00133608.3	7	8	6	7	0	28	7	5.6	1.0	14.29%	3.8	87.37%	Ruvbl2
PI00402913.1	6	5	7	5	5	28	5	5.6	1.0	16.67%	1.2	20.38%	Sic25a27
PI00118723.3	4	4	5	4	11	28	4	5.6	0.6	13.32%	3.8	56.79%	Syng1
PI00108431.8	6	6	5	7	4	28	6	5.6	0.6	10.19%	1.5	28.64%	Ttyh3
PI00115530.1	6	5	8	6	2	27	6	5.4	1.5	24.12%	3.1	57.28%	1110020G09Rik
PI00308938.6	5	4	7	6	5	27	5	5.4	1.5	28.64%	1.0	16.67%	1500032D16Rik
PI00278230.5	6	7	9	2	3	27	6	5.4	1.5	20.83%	3.8	81.13%	2410166I05Rik
PI00229680.5	6	2	8	9	2	27	6	5.4	3.1	57.28%	3.8	59.78%	Aars
PI00229030.1	8	8	6	5	0	27	6	5.4	1.2	15.75%	3.2	87.67%	Abcc9
PI00848443.1	5	3	6	8	5	27	5	5.4	1.5	32.73%	1.5	24.12%	Ache
PI00136334.1	5	3	6	6	7	27	6	5.4	1.5	32.73%	0.6	9.12%	Acol2
PI00462140.1	7	2	8	7	3	27	7	5.4	3.2	56.73%	2.6	44.10%	Atp1f1
PI00318428.4	3	12	6	3	3	27	3	5.4	4.6	65.47%	1.7	43.30%	Bcap31
PI00845689.1	6	4	5	7	5	27	5	5.4	1.0	20.00%	1.2	20.38%	Cd200
PI00308813.1	5	4	5	7	6	27	5	5.4	0.6	12.37%	1.0	16.67%	D2hgdh
PI00405756.8	7	6	4	3	7	27	6	5.4	1.5	26.96%	2.1	44.61%	Dnaja3
PI00133034.3	5	6	7	3	6	27	6	5.4	1.0	16.67%	2.1	39.03%	Dock4
PI00676841.1	10	0	8	9	0	27	8	5.4	5.3	88.19%	4.9	87.05%	Ephb3
PI00874555.1	4	9	6	5	3	27	5	5.4	2.5	39.74%	1.5	32.73%	Epn1
PI00120264.1	4	4	7	7	5	27	5	5.4	1.7	34.64%	1.2	18.23%	Fga
PI00620097.2	4	5	4	7	7	27	5	5.4	0.6	13.32%	1.7	28.87%	Gabrb2
PI00114613.3	7	5	5	7	3	27	5	5.4	1.2	20.38%	2.0	40.00%	Gdap1
PI00127204.1	5	11	5	4	2	27	5	5.4	3.5	49.49%	1.5	41.66%	Gng4
PI00406030.2	5	6	5	5	6	27	5	5.4	0.6	10.83%	0.6	10.83%	Ipo7
PI00785231.1	7	3	4	10	3	27	4	5.4	2.1	44.61%	3.8	66.81%	Kif2a
PI00120914.1	9	4	3	6	5	27	5	5.4	3.2	60.27%	1.5	32.73%	Mtx1
PI00227838.4	4	10	5	6	2	27	5	5.4	3.2	50.76%	2.1	48.04%	Nup133
PI00115089.2	7	3	4	6	7	27	6	5.4	2.1	44.61%	1.5	26.96%	Olfm1
PI00331283.2	5	5	8	3	6	27	5	5.4	1.7	28.87%	2.5	44.41%	Pdk2
PI00331394.3	7	10	6	4	0	27	6	5.4	2.1	27.15%	3.1	91.65%	Ppp2r5d
PI00404474.1	8	2	6	9	2	27	6	5.4	3.1	57.28%	3.5	61.97%	Sf3a1
PI00153501.5	12	0	15	0	0	27	0	5.4	7.9	88.19%	8.7	173.21%	Syt5

IP100387318.3	5	9	4	9	0	27	5	5.4	2.6	44.10%	4.5	104.06%	Trim28
IP100648214.1	8	6	8	5	0	27	6	5.4	1.2	15.75%	4.0	93.26%	Wdr47
IP100132529.2	7	8	6	4	2	27	6	5.4	1.0	14.29%	2.0	50.00%	Xpo7
IP100318545.3	4	7	5	5	5	26	5	5.2	1.5	28.64%	0.0	0.00%	1500001M20Rik
IP100648981.1	7	7	6	4	2	26	6	5.2	0.6	8.66%	2.0	50.00%	Acaca
IP100749720.1	7	4	4	8	3	26	4	5.2	1.7	34.64%	2.6	52.92%	AK2
IP100337930.4	4	4	8	4	6	26	4	5.2	2.3	43.30%	2.0	33.33%	Ap1m1
IP100466878.1	7	5	5	5	4	26	5	5.2	1.2	20.38%	0.6	12.37%	Bcam
IP100776252.1	4	9	5	4	4	26	4	5.2	2.6	44.10%	0.6	13.32%	Cacna1e
IP100875550.1	9	4	2	7	4	26	4	5.2	3.6	72.11%	2.5	58.08%	Centa1
IP100762081.1	8	4	0	4	10	26	4	5.2	4.0	100.00%	5.0	107.85%	Coro1a
IP100775807.1	7	4	8	5	2	26	5	5.2	2.1	32.87%	3.0	60.00%	F11r
IP100775975.1	6	5	6	4	5	26	5	5.2	0.6	10.19%	1.0	20.00%	Glo1
IP100278611.6	4	5	5	7	5	26	5	5.2	0.6	12.37%	1.2	20.38%	Hhahl
IP100126343.1	8	0	18	0	0	26	0	5.2	9.0	104.06%	10.4	173.21%	Hspa1a
IP100135977.3	4	4	6	4	8	26	4	5.2	1.2	24.74%	2.0	33.33%	Kcna1
IP100475378.4	8	7	6	2	3	26	6	5.2	1.0	14.29%	2.1	56.77%	Lyn
IP100467127.4	6	5	6	5	4	26	5	5.2	0.6	10.19%	1.0	20.00%	Magi1
IP100464256.2	5	9	8	0	4	26	5	5.2	2.1	28.39%	4.0	100.00%	Mapk3
IP100320378.3	6	9	8	0	3	26	6	5.2	1.5	19.92%	4.0	110.22%	Naga
IP100129164.1	10	2	0	0	14	26	2	5.2	5.3	132.29%	8.1	173.21%	Ndrgr4
IP100135698.1	8	4	6	8	0	26	6	5.2	2.0	33.33%	4.2	89.21%	Numa1
IP100315808.3	4	6	0	8	8	26	6	5.2	3.1	91.65%	4.6	86.60%	Park7
IP100137206.1	4	6	5	8	3	26	5	5.2	1.0	20.00%	2.5	47.19%	Pdcd6
IP100130949.2	5	5	5	5	6	26	5	5.2	0.0	0.00%	0.6	10.83%	Pex14
IP100226563.4	5	4	5	7	5	26	5	5.2	0.6	12.37%	1.2	20.38%	Pgd
IP100876541.1	8	5	5	6	2	26	5	5.2	1.7	28.87%	2.1	48.04%	Pgs1
IP100664442.3	8	4	4	7	3	26	4	5.2	2.3	43.30%	2.1	44.61%	Pip4k2b
IP100123129.1	8	4	4	7	3	26	4	5.2	2.3	43.30%	2.1	44.61%	Praf2
IP10022125.1	7	4	7	5	3	26	5	5.2	1.7	28.87%	2.0	40.00%	Psmc3
IP100227260.4	9	4	9	4	0	26	4	5.2	2.9	39.36%	4.5	104.06%	Rpl3
IP100122297.5	4	3	6	4	9	26	4	5.2	1.5	35.25%	2.5	39.74%	Scg5
IP100121362.2	4	8	9	5	0	26	5	5.2	2.6	37.80%	4.5	96.63%	Snrpd1
IP100187407.4	7	8	7	4	0	26	7	5.2	0.6	7.87%	3.5	95.78%	Srpb

IP00336318.1	8	6	6	3	3	26	6	5.2	1.2	17.32%	1.7	43.30%	Tdrkh
IP00125813.1	6	10	6	4	0	26	6	5.2	2.3	31.49%	3.1	91.65%	Tmed9
IP00132600.1	7	6	7	3	3	26	6	5.2	0.6	8.66%	2.3	53.29%	Tmem43
IP00410904.1	7	3	2	6	8	26	6	5.2	2.6	66.14%	3.1	57.28%	Txndc13
IP00622540.2	0	7	7	8	4	26	7	5.2	4.0	86.60%	2.1	32.87%	Wasf3
IP00130322.5	5	6	3	9	3	26	5	5.2	1.5	32.73%	3.5	69.28%	Wfs1
IP00323554.3	8	4	5	6	2	25	5	5.0	2.1	36.74%	2.1	48.04%	Actr10
IP00123183.3	11	2	5	4	3	25	4	5.0	4.6	76.38%	1.0	25.00%	AI464131
IP0032128.3	9	3	5	4	4	25	4	5.0	3.1	53.91%	0.6	13.32%	Akap12
IP00227832.8	4	3	9	6	3	25	4	5.0	3.2	60.27%	3.0	50.00%	Agp1
IP00262198.6	3	4	11	0	7	25	4	5.0	4.4	72.65%	5.6	92.80%	Asna1
IP00624653.1	3	9	2	7	4	25	4	5.0	3.8	81.13%	2.5	58.08%	Cndp2
IP00114256.2	5	7	4	6	3	25	5	5.0	1.5	28.64%	1.5	35.25%	Cyp29
IP00132194.3	2	9	7	3	4	25	4	5.0	3.6	60.09%	2.1	44.61%	Dhrs7
IP00137601.2	5	5	6	5	4	25	5	5.0	0.6	10.83%	1.0	20.00%	Dlgap1
IP00720005.1	6	7	6	6	0	25	6	5.0	0.6	9.12%	3.5	86.60%	ENSMUSG00000059179
IP0066885.2	8	7	6	4	0	25	6	5.0	1.0	14.29%	3.1	91.65%	ENSMUSG00000060419
IP00227035.2	9	6	7	0	3	25	6	5.0	1.5	20.83%	3.5	105.36%	Epb4.2
IP00667787.1	8	2	6	7	2	25	6	5.0	3.1	57.28%	2.6	52.92%	Eps15
IP00121013.1	0	10	8	7	0	25	7	5.0	5.3	88.19%	4.4	87.18%	Finc
IP00459642.1	8	4	5	3	5	25	5	5.0	2.1	36.74%	1.2	26.65%	Gcs1
IP00344142.4	4	5	8	6	2	25	5	5.0	2.1	36.74%	3.1	57.28%	Glb1
IP00853823.2	6	4	4	4	7	25	4	5.0	1.2	24.74%	1.7	34.64%	Grm1
IP00624863.1	6	7	6	6	0	25	6	5.0	0.6	9.12%	3.5	86.60%	Hnrpa3
IP00344734.3	6	7	4	5	3	25	5	5.0	1.5	26.96%	1.0	25.00%	Hrsp12
IP00274222.2	6	3	6	5	5	25	5	5.0	1.7	34.64%	0.6	10.83%	Itga3
IP00221921.5	5	2	7	5	6	25	5	5.0	2.5	53.93%	1.0	16.67%	Kpna4
IP00117264.1	6	7	6	6	0	25	6	5.0	0.6	9.12%	3.5	86.60%	LOC545592
IP00136475.1	6	7	7	5	0	25	6	5.0	0.6	8.66%	3.6	90.14%	March5
IP00848939.1	7	5	5	4	4	25	5	5.0	1.2	20.38%	0.6	13.32%	mt-Alp8
IP00828826.1	4	7	7	4	3	25	4	5.0	1.7	28.87%	2.1	44.61%	Ncstn
IP00321734.7	6	6	5	5	3	25	5	5.0	0.6	10.19%	1.2	26.65%	Nos1
IP00127596.1	4	6	6	5	4	25	5	5.0	1.2	21.65%	1.0	20.00%	Nsdhl
IP00111045.1	7	5	6	3	4	25	5	5.0	1.0	16.67%	1.5	35.25%	Ocln



IP100115544.1	5	7	5	6	2	25	5	5.0	1.2	20.38%	2.1	48.04%	Pex5l
IP100469227.2	7	5	5	5	3	25	5	5.0	1.2	20.38%	1.2	26.65%	Pgls
IP100315463.4	5	6	4	7	3	25	5	5.0	1.0	20.00%	2.1	44.61%	Prrt3
IP100128904.1	9	4	6	6	0	25	6	5.0	2.5	39.74%	3.5	86.60%	Psmb2
IP100831223.1	2	7	6	3	7	25	6	5.0	2.6	52.92%	2.1	39.03%	Pvt1
IP100113262.1	5	5	4	7	4	25	5	5.0	0.6	12.37%	1.7	34.64%	Rdh14
IP100284521.9	8	7	6	4	0	25	6	5.0	1.0	14.29%	3.1	91.65%	Rps16
IP100457661.5	5	5	7	4	4	25	5	5.0	1.2	20.38%	1.7	34.64%	Rufy3
IP100125970.1	4	8	6	4	3	25	4	5.0	2.0	33.33%	1.5	35.25%	Slc9a1
IP100653566.1	2	3	9	6	4	24	4	4.8	3.8	81.13%	2.5	39.74%	6430704M03Rik
IP100221486.1	7	5	4	8	0	24	5	4.8	1.5	28.64%	4.0	100.00%	Atrgap1
IP100387216.1	7	3	6	5	3	24	5	4.8	2.1	39.03%	1.5	32.73%	Alp13a5
IP100785452.1	3	6	5	6	4	24	5	4.8	1.5	32.73%	1.0	20.00%	AU040829
IP100788326.1	7	2	5	4	6	24	5	4.8	2.5	53.93%	1.0	20.00%	BC026585
IP100678133.3	5	0	8	4	7	24	5	4.8	4.0	93.26%	2.1	32.87%	Bpnt1
IP100454120.1	5	7	5	5	2	24	5	4.8	1.2	20.38%	1.7	43.30%	C85492
IP100112001.1	4	5	3	5	7	24	5	4.8	1.0	25.00%	2.0	40.00%	Ccdc44
IP100831371.1	4	6	5	4	5	24	5	4.8	1.0	20.00%	0.6	12.37%	Cops3
IP100330323.1	4	7	6	2	5	24	5	4.8	1.5	26.96%	2.1	48.04%	Cops4
IP100330619.3	2	9	4	5	4	24	4	4.8	3.6	72.11%	0.6	13.32%	Cops8
IP100113738.2	3	8	5	4	4	24	4	4.8	2.5	47.19%	0.6	13.32%	Cpne2
IP100624501.2	7	5	4	4	4	24	4	4.8	1.5	28.64%	0.0	0.00%	Cryz
IP100270496.5	5	12	7	0	0	24	5	4.8	3.6	45.07%	4.0	173.21%	Dab2ip
IP100409462.2	7	5	4	4	4	24	4	4.8	1.5	28.64%	0.0	0.00%	Dgkb
IP100380817.6	2	10	5	4	3	24	4	4.8	4.0	71.32%	1.0	25.00%	Eif4h
IP100458074.3	4	6	9	5	0	24	5	4.8	2.5	39.74%	4.5	96.63%	Fis1
IP100126759.6	6	3	2	6	7	24	6	4.8	2.1	56.77%	2.6	52.92%	Folr1
IP100115668.1	7	3	7	5	2	24	5	4.8	2.3	40.75%	2.5	53.93%	Hapln4
IP100310701.4	4	4	6	6	4	24	4	4.8	1.2	24.74%	1.2	21.65%	Hdh2
IP100223415.3	5	5	6	2	6	24	5	4.8	0.6	10.83%	2.3	49.49%	Hsd1
IP100131830.1	0	24	0	0	0	24	0	4.8	13.9	173.21%	0.0	#DIV/0!	Krt6a
IP100123104.1	0	15	0	9	0	24	0	4.8	8.7	173.21%	5.2	173.21%	mCG_121688
IP100129276.1	5	5	7	5	2	24	5	4.8	1.2	20.38%	2.5	53.93%	Mlic1
IP100129264.1	8	6	5	3	2	24	5	4.8	1.5	24.12%	1.5	45.83%	Mlycd

IP100109611.1	8	7	0	6	3	24	6	4.8	4.4	87.18%	3.0	100.00%	mt-Atp6
IP100230034.5	10	3	2	5	4	24	4	4.8	4.4	87.18%	1.5	41.66%	Mtch1
IP100755161.2	6	3	11	2	2	24	3	4.8	4.0	60.62%	5.2	103.92%	Nol5a
IP100330477.4	3	7	5	7	2	24	5	4.8	2.0	40.00%	2.5	53.93%	Nup214
IP100318485.3	10	3	0	7	11	24	3	4.8	5.1	118.42%	6.4	173.21%	Opa3
IP100109142.4	6	4	2	4	5	24	5	4.8	2.0	50.00%	2.5	53.93%	Pafah1b1
IP100323349.2	5	6	6	4	3	24	5	4.8	0.6	10.19%	1.5	35.25%	Pcyox1l
IP100120080.2	4	3	3	4	10	24	4	4.8	0.6	17.32%	3.8	66.81%	Psma3
IP100553717.6	5	4	6	6	3	24	5	4.8	1.0	20.00%	1.7	34.64%	Psma7
IP100262114.1	6	3	2	13	0	24	3	4.8	2.1	56.77%	7.0	140.00%	Sf3a3
IP100137706.1	3	4	9	8	0	24	4	4.8	3.2	60.27%	4.9	87.05%	Sfrs7
IP100116554.2	6	7	6	2	3	24	6	4.8	0.6	9.12%	2.1	56.77%	Sic30a9
IP100421081.3	8	4	7	2	3	24	4	4.8	2.1	32.87%	2.6	66.14%	Snpb
IP100130640.5	7	4	5	5	3	24	5	4.8	1.5	28.64%	1.2	26.65%	Snx27
IP100130307.1	8	4	3	5	4	24	4	4.8	2.6	52.92%	1.0	25.00%	Tpm4
IP100120592.1	6	6	7	5	0	24	6	4.8	0.6	9.12%	3.6	90.14%	Ube2v2
IP100331251.1	8	3	3	4	6	24	4	4.8	2.9	61.86%	1.5	35.25%	Vps45
IP100123891.3	5	4	6	5	3	23	5	4.6	1.0	20.00%	1.5	32.73%	6330407J3Rik
IP100123709.1	7	4	8	0	4	23	4	4.6	2.1	32.87%	4.0	100.00%	Abca7
IP100849927.1	4	5	6	4	4	23	4	4.6	1.0	20.00%	1.2	24.74%	Ablim2
IP100223882.5	4	5	5	5	4	23	5	4.6	0.6	12.37%	0.6	12.37%	Acad8
IP100653743.1	9	5	5	4	0	23	5	4.6	2.3	36.46%	2.6	88.19%	Aifm3
IP100323796.3	6	4	6	5	2	23	5	4.6	1.2	21.65%	2.1	48.04%	Capn5
IP100458711.2	4	5	5	5	4	23	5	4.6	0.6	12.37%	0.6	12.37%	Clu
IP100113992.2	4	10	5	4	0	23	4	4.6	3.2	50.76%	2.6	88.19%	Crip2
IP100662277.1	5	5	4	5	4	23	5	4.6	0.6	12.37%	0.6	13.32%	Cul2
IP100114833.1	7	3	6	4	3	23	4	4.6	2.1	39.03%	1.5	35.25%	Cul3
IP100132966.3	7	4	7	3	2	23	4	4.6	1.7	28.87%	2.6	66.14%	Dhodh
IP100134137.1	7	3	5	6	2	23	5	4.6	2.0	40.00%	2.1	48.04%	Dpm1
IP100122346.2	8	8	5	0	2	23	5	4.6	1.7	24.74%	2.5	107.85%	Ehd2
IP100131584.1	4	5	8	4	2	23	4	4.6	2.1	36.74%	3.1	65.47%	Ergic1
IP100136402.2	2	7	5	5	4	23	5	4.6	2.5	53.93%	0.6	12.37%	Esd
IP100134997.3	7	2	5	4	5	23	5	4.6	2.5	53.93%	0.6	12.37%	Exoc7
IP100119116.1	5	9	5	2	2	23	5	4.6	2.3	36.46%	1.7	57.74%	Fkbp8

IP00223371.2	8	4	6	3	2	23	4	4.6	2.0	33.33%	2.1	56.77%	Fry
IP00830222.1	5	6	5	5	2	23	5	4.6	0.6	10.83%	1.7	43.30%	Gaa
IP00265107.4	5	8	4	4	2	23	4	4.6	2.1	36.74%	1.2	34.64%	Gabra1
IP00118674.7	4	3	3	2	11	23	3	4.6	0.6	17.32%	4.9	92.49%	Glrx5
IP00331519.2	5	0	4	4	10	23	4	4.6	2.6	88.19%	3.5	57.74%	Hdhd3
IP00848482.1	6	8	4	5	0	23	5	4.6	2.0	33.33%	2.6	88.19%	Hspb1
IP00230440.6	6	5	5	3	4	23	5	4.6	0.6	10.83%	1.0	25.00%	Lrig1
IP00402968.1	5	4	5	4	5	23	5	4.6	0.6	12.37%	0.6	12.37%	Ly6a
IP00331014.1	2	8	5	4	4	23	4	4.6	3.0	60.00%	0.6	13.32%	Ndufb2
IP00129423.1	5	5	3	6	4	23	5	4.6	1.2	26.65%	1.5	35.25%	Nup98
IP00223216.5	6	0	5	8	4	23	5	4.6	3.2	87.67%	2.1	36.74%	Pacs1
IP00131994.1	4	4	3	9	3	23	4	4.6	0.6	15.75%	3.5	69.28%	Pitpnm1
IP00136246.1	6	5	6	3	3	23	5	4.6	0.6	10.19%	1.7	43.30%	Pnpla8
IP00228877.1	4	4	5	5	5	23	5	4.6	0.6	13.32%	0.0	0.00%	Ppp1r9a
IP00138190.1	2	9	8	4	0	23	4	4.6	3.8	59.78%	4.0	100.00%	Psmb3
IP00116896.1	3	8	5	5	2	23	5	4.6	2.5	47.19%	1.7	43.30%	Psmd5
IP00132093.1	3	5	7	8	0	23	5	4.6	2.0	40.00%	4.4	87.18%	Pvalb
IP00229722.3	5	7	6	5	0	23	5	4.6	1.0	16.67%	3.2	87.67%	Rab3gap1
IP00124639.1	5	4	6	8	0	23	5	4.6	1.0	20.00%	4.2	89.21%	Rab3gap2
IP00112327.1	6	5	5	5	2	23	5	4.6	0.6	10.83%	1.7	43.30%	Soye1
IP00623737.2	7	7	3	2	4	23	4	4.6	2.3	40.75%	1.0	33.33%	Sec23a
IP00170128.7	8	5	5	2	3	23	5	4.6	1.7	28.87%	1.5	45.83%	Shmt2
IP00874854.1	8	5	2	5	3	23	5	4.6	3.0	60.00%	1.5	45.83%	Snd1
IP00331444.7	5	5	5	5	3	23	5	4.6	0.0	0.00%	1.2	26.65%	Sypl
IP00170213.1	7	3	4	5	4	23	4	4.6	2.1	44.61%	0.6	13.32%	Tmem16f
IP00129491.1	7	0	7	3	6	23	6	4.6	4.0	86.60%	2.1	39.03%	Txn2
IP00468674.5	6	4	5	5	3	23	5	4.6	1.0	20.00%	1.2	26.65%	Txnrd1
IP00322497.4	5	0	4	6	7	22	5	4.4	2.6	88.19%	1.5	26.96%	1810026J23Rik
IP00119680.3	7	3	6	4	2	22	4	4.4	2.1	39.03%	2.0	50.00%	Acads
IP00120085.1	4	5	5	4	4	22	4	4.4	0.6	12.37%	0.6	13.32%	Addc1
IP00662244.2	5	5	5	4	3	22	5	4.4	0.0	0.00%	1.0	25.00%	Adcy1
IP00162790.1	6	2	4	4	6	22	4	4.4	2.0	50.00%	1.2	24.74%	Agf
IP00311387.1	6	4	5	3	4	22	4	4.4	1.0	20.00%	1.0	25.00%	Agpat7
IP00310099.1	5	5	7	3	2	22	5	4.4	1.2	20.38%	2.6	66.14%	Arlfp2

PI00331644.5	7	3	7	3	2	22	3	4.4	2.3	40.75%	2.6	66.14%	Cacng8
PI00230277.3	6	4	7	5	0	22	4.4	1.5	1.5	26.96%	3.6	90.14%	Cobll1
PI00130240.1	6	4	6	3	3	22	4.4	1.2	1.2	21.65%	1.7	43.30%	D3Bwg0562e
PI00454008.1	6	2	4	7	3	22	4.4	2.0	2.0	50.00%	2.1	44.61%	Dkc1
PI00130381.1	4	6	6	4	2	22	4.4	1.2	1.2	21.65%	2.0	50.00%	Dnajc13
PI00119432.1	0	0	22	0	0	22	4.4	12.7	12.7	173.21%	12.7	173.21%	EG629557
PI00620040.3	6	4	3	3	6	22	4.4	1.5	1.5	35.25%	1.7	43.30%	Gstz1
PI00320303.1	7	4	6	5	0	22	4.4	1.5	1.5	26.96%	3.2	87.67%	Hnrpd
PI00750694.2	8	0	0	10	4	22	4.4	4.6	4.6	173.21%	5.0	107.85%	Klc2
PI00112126.1	5	4	6	4	3	22	4.4	1.0	1.0	20.00%	1.5	35.25%	Krr74
PI00659608.4	4	5	4	7	2	22	4.4	0.6	0.6	13.32%	2.5	58.08%	Mapre1
PI00132444.1	5	3	8	3	3	22	4.4	2.5	2.5	47.19%	2.9	61.86%	Mcee
PI00380280.3	3	7	5	4	3	22	4.4	2.0	2.0	40.00%	1.0	25.00%	Mcts1
PI00474631.6	5	4	6	3	4	22	4.4	1.0	1.0	20.00%	1.5	35.25%	Mllt4
PI00114866.1	6	4	7	5	0	22	4.4	1.5	1.5	26.96%	3.6	90.14%	Osbp1a
PI00312752.3	8	3	2	6	3	22	4.4	3.2	3.2	74.18%	2.1	56.77%	Pak3
PI00762331.2	5	2	5	4	6	22	4.4	1.7	1.7	43.30%	1.0	20.00%	Ppp1r9b
PI00170008.1	0	0	0	0	22	22	4.4	0.0	0.0	#DIV/0!	12.7	173.21%	Ppp2r1b
PI00108125.4	3	9	0	4	6	22	4.4	4.6	4.6	114.56%	3.1	91.65%	Psm6
PI00224073.3	5	5	6	3	3	22	4.4	0.6	0.6	10.83%	1.7	43.30%	Psmc1
PI00356063.2	8	6	2	4	2	22	4.4	3.1	3.1	57.28%	1.2	43.30%	Ptpn11
PI00221547.4	8	4	3	4	3	22	4.4	2.6	2.6	52.92%	0.6	17.32%	Pzp
PI00130624.1	5	7	8	2	0	22	4.4	1.5	1.5	22.91%	4.2	124.90%	Rpl18a
PI00132456.1	6	7	6	0	3	22	4.4	0.6	0.6	9.12%	3.0	100.00%	Sic12a9
PI00556823.3	7	6	5	2	2	22	4.4	1.0	1.0	16.67%	1.7	57.74%	Sic25a20
PI00320850.3	4	0	7	3	8	22	4.4	3.5	3.5	95.78%	2.6	44.10%	Sic30a3
PI00330958.2	5	4	6	5	2	22	4.4	1.0	1.0	20.00%	2.1	48.04%	Sic6a9
PI00459383.2	4	5	4	6	3	22	4.4	0.6	0.6	13.32%	1.5	35.25%	Six16
PI00349020.6	6	4	4	4	4	22	4.4	1.2	1.2	24.74%	0.0	0.00%	Suc1g2
PI00655150.1	5	11	6	0	0	22	4.4	3.2	3.2	43.83%	3.5	173.21%	Sv2c
PI00224775.2	5	5	3	6	3	22	4.4	1.2	1.2	26.65%	1.7	43.30%	Usmg5
PI00153959.1	6	5	3	4	3	21	4.2	1.5	1.5	32.73%	0.6	17.32%	1810027O10RIK
PI00875497.1	4	6	4	4	3	21	4.2	1.2	1.2	24.74%	0.6	15.75%	Abhd11
PI00136984.1	5	2	4	6	4	21	4.2	1.5	1.5	41.66%	1.2	24.74%	AI595406

PI00221685.1	6	3	3	4	5	21	4	4.2	1.7	43.30%	1.0	25.00%	Armc1
PI00420726.3	4	3	4	6	4	21	4	4.2	0.6	15.75%	1.2	24.74%	Atcay
PI00137460.3	4	5	6	4	2	21	4	4.2	1.0	20.00%	2.0	50.00%	Bmpr2
PI00469318.1	7	4	5	5	0	21	5	4.2	1.5	28.64%	2.9	86.60%	Cacnb3
PI00112555.3	6	2	5	5	3	21	5	4.2	2.1	48.04%	1.2	26.65%	Cdh13
PI00755309.1	3	7	3	5	3	21	3	4.2	2.3	53.29%	1.2	31.49%	Cpt1c
PI00119615.1	8	5	4	0	4	21	4	4.2	2.1	36.74%	2.3	86.60%	Crat
PI00139780.1	5	8	4	2	2	21	4	4.2	2.1	36.74%	1.2	43.30%	Dbln
PI00139259.1	3	4	4	4	6	21	4	4.2	0.6	15.75%	1.2	24.74%	Dip2b
PI00457815.3	8	8	5	0	0	21	5	4.2	1.7	24.74%	2.9	173.21%	Dnajc13
PI00331345.5	7	3	5	3	3	21	3	4.2	2.0	40.00%	1.2	31.49%	Dnajc3b
PI00225267.6	6	6	6	3	0	21	6	4.2	0.0	0.00%	3.0	100.00%	Dpy1911
PI00330695.5	4	8	4	3	2	21	4	4.2	2.3	43.30%	1.0	33.33%	EG667977
PI00108998.1	4	3	4	2	8	21	4	4.2	0.6	15.75%	3.1	65.47%	Exoc1
PI00329953.1	4	7	3	7	0	21	4	4.2	2.1	44.61%	3.5	105.36%	Fah
PI00379695.2	5	5	5	4	2	21	5	4.2	0.0	0.00%	1.5	41.66%	Gabra6
PI00323600.4	5	5	5	2	4	21	5	4.2	0.0	0.00%	1.5	41.66%	Gars
PI00120166.1	3	4	7	4	3	21	4	4.2	2.1	44.61%	2.1	44.61%	Gcat
PI00153107.3	5	8	5	0	3	21	5	4.2	1.7	28.87%	2.5	94.37%	Gpc6
PI00116850.3	6	4	3	4	4	21	4	4.2	1.5	35.25%	0.6	15.75%	Grb2
PI00420590.3	6	4	7	4	0	21	4	4.2	1.5	26.96%	3.5	95.78%	H2-K1
PI00120094.1	4	8	4	3	2	21	4	4.2	2.3	43.30%	1.0	33.33%	H2-Q10
PI00874518.1	7	4	4	2	4	21	4	4.2	1.7	34.64%	1.2	34.64%	Huwe1
PI00331064.7	4	5	4	4	4	21	4	4.2	0.6	13.32%	0.0	0.00%	Kpna6
PI00121827.4	9	0	10	2	0	21	2	4.2	5.5	86.96%	5.3	132.29%	Krt4
PI00554884.1	10	2	9	0	0	21	2	4.2	4.4	62.27%	5.2	173.21%	LOC637606
PI00315187.5	5	3	3	6	4	21	4	4.2	1.2	31.49%	1.5	35.25%	Lphn2
PI00420970.3	6	7	4	4	0	21	4	4.2	1.5	26.96%	2.3	86.60%	Mpdu1
PI00131173.2	5	7	4	3	2	21	4	4.2	1.5	28.64%	1.0	33.33%	Mpi
PI00112536.1	3	0	4	5	9	21	4	4.2	2.1	89.21%	2.6	44.10%	Mirpl12
PI00775993.1	4	4	5	8	0	21	4	4.2	0.6	13.32%	4.0	93.26%	Mthfd1
PI00308851.3	6	5	2	2	6	21	5	4.2	2.1	48.04%	2.3	69.28%	Neddd4l
PI00114246.2	5	2	2	6	6	21	5	4.2	1.7	57.74%	2.3	49.49%	Omg
PI00850889.1	5	3	3	6	4	21	4	4.2	1.2	31.49%	1.5	35.25%	Podhgc5

IP100405114.1	5	0	4	3	9	21	4	4.2	2.6	88.19%	3.2	60.27%	Pfn2
IP100320420.3	6	0	5	4	6	21	5	4.2	3.2	87.67%	1.0	20.00%	Phyhip
IP100123278.1	5	5	6	3	2	21	5	4.2	0.6	10.83%	2.1	56.77%	Pld2
IP100468633.3	5	2	4	6	4	21	4	4.2	1.5	41.66%	1.2	24.74%	Ppfa2
IP100830853.1	6	6	3	0	6	21	6	4.2	1.7	34.64%	3.0	100.00%	Ppm2c
IP100130992.1	4	5	4	5	3	21	4	4.2	0.6	13.32%	1.0	25.00%	Ppp5c
IP100119180.1	5	4	7	2	3	21	4	4.2	1.5	28.64%	2.6	66.14%	Prkce
IP100408119.6	0	0	0	10	11	21	0	4.2	0.0	#DIV/0!	6.1	86.90%	Rab6b
IP100129964.1	3	5	5	6	2	21	5	4.2	1.2	26.65%	2.1	48.04%	Rapgef2
IP100221782.3	3	5	6	3	4	21	4	4.2	1.5	32.73%	1.5	35.25%	Rogdi
IP100378768.5	5	4	4	4	4	21	4	4.2	0.6	13.32%	0.0	0.00%	Rpl8
IP100113772.1	7	5	5	4	0	21	5	4.2	1.2	20.38%	2.6	88.19%	Rps7
IP100109108.2	9	5	2	5	0	21	5	4.2	3.5	65.85%	2.5	107.85%	Rps9
IP100322749.3	3	4	6	4	4	21	4	4.2	1.5	35.25%	1.2	24.74%	Sh3gl1
IP100129572.1	5	3	3	7	3	21	3	4.2	1.2	31.49%	2.3	53.29%	Sipa111
IP100817008.1	4	6	6	2	3	21	4	4.2	1.2	21.65%	2.1	56.77%	Slc25a10
IP100222760.1	7	5	7	2	0	21	5	4.2	1.2	18.23%	3.6	120.19%	Snap23
IP100649060.1	3	6	10	2	0	21	3	4.2	3.5	55.45%	5.3	132.29%	Spcc2
IP100225301.1	5	3	6	5	2	21	5	4.2	1.5	32.73%	2.1	48.04%	Srr
IP100115522.3	7	0	4	5	5	21	5	4.2	3.5	95.78%	0.6	12.37%	Tnik
IP100470095.1	4	0	5	6	6	21	5	4.2	2.6	88.19%	0.6	10.19%	Tprgl
IP100227805.6	5	3	7	3	3	21	3	4.2	2.0	40.00%	2.3	53.29%	Ubxdl
IP100762199.2	3	5	4	5	4	21	4	4.2	1.0	25.00%	0.6	13.32%	Vps13c
IP100551454.3	20	0	0	0	0	20	0	4.0	11.5	173.21%	0.0	#DIV/0!	Abcb4
IP100132217.1	2	4	2	5	7	20	4	4.0	1.2	43.30%	2.5	53.93%	Ahcy
IP100320241.1	7	6	4	3	0	20	4	4.0	1.5	26.96%	2.1	89.21%	Ap3m1
IP100655136.3	4	4	6	4	2	20	4	4.0	1.2	24.74%	2.0	50.00%	Atrgef7
IP100110100.1	5	3	8	4	0	20	4	4.0	2.5	47.19%	4.0	100.00%	Alp13a1
IP100461290.1	4	6	0	5	5	20	5	4.0	3.1	91.65%	2.9	86.60%	Ckm
IP100473646.2	9	0	3	6	2	20	3	4.0	4.6	114.56%	2.1	56.77%	Clip2
IP100221845.1	3	5	3	4	5	20	4	4.0	1.2	31.49%	1.0	25.00%	Csnk2b
IP100314467.3	5	3	3	2	7	20	3	4.0	1.2	31.49%	2.6	66.14%	Cyld
IP100396840.3	5	5	4	4	2	20	4	4.0	0.6	12.37%	1.2	34.64%	D10Wsu52e
IP100222307.1	4	3	2	8	3	20	3	4.0	1.0	33.33%	3.2	74.18%	Dusp3

IP100396753.2	2	6	6	4	2	20	4	4.0	2.3	49.49%	2.0	50.00%	Ecsit
IP100624663.3	4	5	6	3	2	20	4	4.0	1.0	20.00%	2.1	56.77%	Egfr
IP100267983.5	7	4	3	2	4	20	4	4.0	2.1	44.61%	1.0	33.33%	Eif5a
IP100115644.2	7	5	2	4	2	20	4	4.0	2.5	53.93%	1.2	43.30%	Ephx2
IP100671847.3	5	6	5	2	2	20	5	4.0	0.6	10.83%	1.7	57.74%	Gabb2
IP100761429.1	6	3	2	4	5	20	4	4.0	2.1	56.77%	1.5	41.66%	Gba
IP100459487.3	5	5	4	3	3	20	4	4.0	0.6	12.37%	0.6	17.32%	Gng7
IP100112616.2	7	4	3	2	4	20	4	4.0	2.1	44.61%	1.0	33.33%	Grm5
IP100354363.3	4	6	6	4	0	20	4	4.0	1.2	21.65%	3.1	91.65%	Gsr
IP100776018.1	7	6	4	0	3	20	4	4.0	1.5	26.96%	2.1	89.21%	Hccs
IP100338094.6	4	4	4	5	3	20	4	4.0	0.0	0.00%	1.0	25.00%	Kcnip4
IP100421041.5	5	6	5	4	0	20	5	4.0	0.6	10.83%	2.6	88.19%	Krt42
IP100856490.1	8	3	2	5	2	20	3	4.0	3.2	74.18%	1.7	57.74%	Lamp1
IP100225378.3	4	8	4	4	0	20	4	4.0	2.3	43.30%	2.3	86.60%	LOC100043780
IP100115257.1	7	4	5	2	2	20	4	4.0	1.5	28.64%	1.7	57.74%	Lrrc8c
IP100607957.1	6	5	3	3	3	20	3	4.0	1.5	32.73%	0.0	0.00%	Ntrk3
IP100113798.2	4	2	4	4	6	20	4	4.0	1.2	34.64%	1.2	24.74%	Ppm1h
IP100750473.1	12	0	4	4	0	20	4	4.0	6.1	114.56%	2.3	86.60%	Pikag1
IP100473521.1	7	5	4	4	0	20	4	4.0	1.5	28.64%	2.3	86.60%	Ptbp1
IP100377609.2	7	8	5	0	0	20	5	4.0	1.5	22.91%	2.9	173.21%	Rasal2
IP100122499.5	7	7	0	6	0	20	6	4.0	4.0	86.60%	3.5	173.21%	Rcc1
IP100315620.2	6	7	7	0	0	20	6	4.0	0.6	8.66%	4.0	173.21%	Rpl14
IP100661338.1	4	8	4	4	0	20	4	4.0	2.3	43.30%	2.3	86.60%	Rps3a
IP100622780.1	7	4	6	0	3	20	4	4.0	1.5	26.96%	3.0	100.00%	Scfd1
IP100321922.2	4	6	5	5	0	20	5	4.0	1.0	20.00%	2.9	86.60%	Sfrs10
IP100128522.1	5	2	5	6	2	20	5	4.0	1.7	43.30%	2.1	48.04%	Strn1
IP100674554.2	10	5	2	3	0	20	3	4.0	4.0	71.32%	1.5	91.65%	Stt3b
IP100130988.1	4	5	5	4	2	20	4	4.0	0.6	12.37%	1.5	41.66%	Tnks1bp1
IP100111960.2	4	2	5	3	6	20	4	4.0	1.5	41.66%	1.5	32.73%	Twf2
IP100130758.1	5	6	0	4	4	19	4	3.8	3.2	87.67%	2.3	86.60%	2900075B16RIK
IP100315538.6	4	6	5	2	2	19	4	3.8	1.0	20.00%	1.7	57.74%	3110047P20RIK
IP100120546.1	4	3	9	3	0	19	3	3.8	3.2	60.27%	4.6	114.56%	9130404D14RIK
IP100124980.1	5	2	3	2	7	19	3	3.8	1.5	45.83%	2.6	66.14%	9330129D05RIK
IP100126500.1	9	2	6	0	2	19	2	3.8	3.5	61.97%	3.1	114.56%	Abca9

IP00119930.2	5	2	3	2	3	7	19	3	3.8	1.5	45.83%	2.6	66.14%	Acad10
IP00662721.1	4	2	4	2	4	7	19	4	3.8	1.2	34.64%	2.5	58.08%	Acss1
IP00113127.7	6	5	4	2	4	2	19	4	3.8	1.0	20.00%	1.2	43.30%	Adpgk
IP00877328.1	8	3	5	3	5	0	19	3	3.8	2.5	47.19%	2.5	94.37%	Als2
IP00471418.1	8	6	5	0	5	0	19	5	3.8	1.5	24.12%	2.9	173.21%	Ank
IP00313296.3	5	6	2	2	4	4	19	4	3.8	2.1	48.04%	1.2	43.30%	BC025575
IP00857969.1	5	4	6	4	6	0	19	4	3.8	1.0	20.00%	3.1	91.65%	Cacnb1
IP00277001.4	7	3	5	2	5	2	19	3	3.8	2.0	40.00%	1.7	57.74%	Cd99l2
IP00109764.2	3	6	3	5	3	2	19	3	3.8	1.7	43.30%	1.5	45.83%	Clpp
IP00404477.4	4	0	4	11	4	0	19	4	3.8	2.3	86.60%	5.6	111.36%	Col6a3
IP00126762.1	8	0	2	4	4	5	19	4	3.8	4.2	124.90%	1.5	41.66%	Cpd
IP00109891.1	4	3	3	5	3	4	19	4	3.8	0.6	17.32%	1.0	25.00%	Ctsa
IP00120346.3	6	5	6	2	6	0	19	5	3.8	0.6	10.19%	3.1	114.56%	Cyp51
IP00137787.3	3	5	3	3	3	5	19	3	3.8	1.2	31.49%	1.2	31.49%	D12Ertid553e
IP00462006.3	5	8	3	3	3	0	19	3	3.8	2.5	47.19%	1.7	86.60%	Dhrs4
IP00551399.3	5	4	4	4	4	2	19	4	3.8	0.6	13.32%	1.2	34.64%	Dhrs7b
IP00459033.2	4	8	4	3	4	0	19	4	3.8	2.3	43.30%	2.1	89.21%	Dnajc10
IP00115902.2	7	4	2	4	4	2	19	4	3.8	2.5	58.08%	1.2	43.30%	Dync111
IP00408734.3	3	9	5	2	5	0	19	3	3.8	3.1	53.91%	2.5	107.85%	Exoc2
IP00308609.1	0	3	6	4	6	6	19	4	3.8	3.0	100.00%	1.2	21.65%	Fahd1
IP00877291.1	7	3	4	3	4	2	19	3	3.8	2.1	44.61%	1.0	33.33%	Fkbp4
IP00758170.2	8	4	5	0	5	2	19	4	3.8	2.1	36.74%	2.5	107.85%	G6pc3
IP00263048.1	3	2	0	8	0	6	19	3	3.8	1.5	91.65%	4.2	89.21%	Gabrb1
IP00128692.1	2	6	6	5	6	0	19	5	3.8	2.3	49.49%	3.2	87.67%	Gabrd
IP00342158.4	3	2	8	0	8	6	19	3	3.8	3.2	74.18%	4.2	89.21%	Gng13
IP00126048.2	7	3	7	0	7	2	19	3	3.8	2.3	40.75%	3.6	120.19%	Gpc4
IP00116308.1	6	6	5	0	5	2	19	5	3.8	0.6	10.19%	2.5	107.85%	Gpr116
IP00652882.2	4	2	4	4	4	5	19	4	3.8	1.2	34.64%	0.6	13.32%	Grm7
IP00127707.1	4	2	7	3	3	3	19	3	3.8	2.5	58.08%	2.3	53.29%	Itgb5
IP00132604.3	6	5	4	2	4	2	19	4	3.8	1.0	20.00%	1.2	43.30%	Itm2b
IP00123059.1	3	5	6	2	6	3	19	3	3.8	1.5	32.73%	2.1	56.77%	Itm2c
IP00750671.1	6	6	4	3	4	0	19	4	3.8	1.2	21.65%	2.1	89.21%	Jtv1
IP00679159.2	4	5	6	4	6	0	19	4	3.8	1.0	20.00%	3.1	91.65%	Kona2
IP00228423.1	4	7	5	3	5	0	19	4	3.8	1.5	28.64%	2.5	94.37%	Krt17



IP100226730.1	8	3	3	3	3	2	19	3	3.8	2.9	61.86%	0.6	21.65%	Letmd1
IP100319111.3	5	3	8	3	3	0	19	3	3.8	2.5	47.19%	4.0	110.22%	Lin7b
IP100467854.1	4	7	3	3	3	2	19	3	3.8	2.1	44.61%	0.6	21.65%	LOC100047443
IP100118875.4	4	8	4	0	4	3	19	4	3.8	2.3	43.30%	2.1	89.21%	Mrs2l
IP100320687.2	4	9	0	4	4	2	19	4	3.8	4.5	104.06%	2.0	100.00%	Ndufa7
IP100322209.5	6	5	4	4	4	0	19	4	3.8	1.0	20.00%	2.3	86.60%	Pcbp2
IP100135087.3	3	8	5	0	6	3	19	3	3.8	2.5	47.19%	2.5	94.37%	Plaa
IP100119220.1	11	0	2	6	6	0	19	2	3.8	5.9	135.22%	3.1	114.56%	Pld1
IP100317074.3	3	5	3	3	2	6	19	3	3.8	1.2	31.49%	2.1	56.77%	Pnpla6
IP100648688.3	0	0	0	0	0	19	19	0	3.8	0.0	#DIV/0!	11.0	173.21%	Ppp2cb
IP100229605.2	5	2	3	5	5	4	19	4	3.8	1.5	45.83%	1.0	25.00%	Pycr2
IP100474558.7	2	6	0	6	6	5	19	5	3.8	3.1	114.56%	3.2	87.67%	Rp2h
IP100468688.4	8	4	7	0	0	0	19	4	3.8	2.1	32.87%	4.0	173.21%	Rpl5
IP100875190.2	5	4	7	3	3	0	19	4	3.8	1.5	28.64%	3.5	105.36%	Rps3
IP100674255.2	6	3	2	6	6	2	19	3	3.8	2.1	56.77%	2.3	69.28%	Sema4a
IP100222632.3	6	5	0	6	6	2	19	5	3.8	3.2	87.67%	3.1	114.56%	Slc39a12
IP100315964.1	6	0	5	8	8	0	19	5	3.8	3.2	87.67%	4.0	93.26%	Slc4a7
IP100411102.3	5	5	0	2	2	7	19	5	3.8	2.9	86.60%	3.6	120.19%	Thbd
IP100121277.1	3	5	0	2	2	9	19	3	3.8	2.5	94.37%	4.7	128.89%	Tsfm
IP100330482.1	4	9	4	4	2	0	19	4	3.8	2.9	50.94%	2.0	100.00%	U2af2
IP100321617.1	6	3	4	4	4	2	19	4	3.8	1.5	35.25%	1.2	34.64%	Vps26b
IP100648318.1	2	4	4	4	4	4	18	4	3.6	1.2	34.64%	0.0	0.00%	2400001E08Rik
IP100172189.1	5	2	4	5	5	2	18	4	3.6	1.5	41.66%	1.5	41.66%	4930572J05Rik
IP100226414.7	6	6	2	4	4	0	18	4	3.6	2.3	49.49%	2.0	100.00%	4932438A13Rik
IP100469218.1	5	7	3	0	3	3	18	3	3.6	2.0	40.00%	1.7	86.60%	Abcb6
IP100719967.1	4	2	2	6	6	4	18	4	3.6	1.2	43.30%	2.0	50.00%	Abcb8
IP100130419.1	3	3	7	3	3	2	18	3	3.6	2.3	53.29%	2.6	66.14%	Atic
IP100129187.1	5	5	2	4	4	2	18	4	3.6	1.7	43.30%	1.2	43.30%	Bat3
IP100344681.2	7	2	4	3	3	2	18	3	3.6	2.5	58.08%	1.0	33.33%	Bckdha
IP100315879.1	3	3	4	3	3	5	18	3	3.6	0.6	17.32%	1.0	25.00%	Camk4
IP100223757.4	3	6	3	4	4	2	18	3	3.6	1.7	43.30%	1.0	33.33%	Cd44
IP100323974.7	6	0	2	8	8	2	18	2	3.6	3.1	114.56%	3.5	86.60%	Cdk5
IP100133948.1	0	4	5	7	7	2	18	4	3.6	2.6	88.19%	2.5	53.93%	Chgb
IP100881287.1	4	0	5	5	5	4	18	4	3.6	2.6	88.19%	0.6	12.37%	Clasp1

IP00624881.4	7	0	5	3	3	18	3	3.6	3.6	3.6	90.14%	1.2	31.49%	Coasy
IP00881291.1	7	5	4	0	2	18	4	3.6	1.5	28.64%	28.64%	2.0	100.00%	D030074E01Rik
IP00658539.1	0	7	4	4	3	18	4	3.6	3.5	95.78%	95.78%	0.6	15.75%	Ddt
IP00134599.1	4	3	5	3	3	18	3	3.6	1.0	25.00%	25.00%	1.2	31.49%	Dlgap2
IP00136505.1	3	5	4	0	6	18	4	3.6	1.0	25.00%	25.00%	3.1	91.65%	Dlgap4
IP00112785.1	10	2	0	6	0	18	2	3.6	5.3	132.29%	132.29%	3.5	173.21%	EG620248
IP00621076.2	4	4	7	3	0	18	4	3.6	1.7	34.64%	34.64%	3.5	105.36%	Endog
IP00135267.1	5	6	2	3	2	18	3	3.6	2.1	48.04%	48.04%	0.6	24.74%	Gpc1
IP00115763.2	4	4	3	4	3	18	4	3.6	0.6	15.75%	15.75%	0.6	17.32%	H2-Ke6
IP00749655.2	6	4	0	8	0	18	4	3.6	3.1	91.65%	91.65%	4.6	173.21%	Hdac2
IP00113746.3	5	3	5	3	2	18	3	3.6	1.2	26.65%	26.65%	1.5	45.83%	Icam2
IP00378015.1	5	2	0	4	7	18	4	3.6	2.5	107.85%	107.85%	3.5	95.78%	Kif21a
IP00136618.1	8	3	5	0	2	18	3	3.6	2.5	47.19%	47.19%	2.5	107.85%	Lpcat1
IP00649958.4	3	6	6	0	3	18	3	3.6	1.7	34.64%	34.64%	3.0	100.00%	Mvp
IP00857740.1	5	5	3	5	0	18	5	3.6	1.2	26.65%	26.65%	2.5	94.37%	Nes
IP00123349.2	4	3	3	6	2	18	3	3.6	0.6	17.32%	17.32%	2.1	56.77%	Nptx1
IP00420807.3	3	7	4	2	2	18	3	3.6	2.1	44.61%	44.61%	1.2	43.30%	Pdgfrb
IP00132475.2	3	2	5	2	6	18	3	3.6	1.5	45.83%	45.83%	2.1	48.04%	Phyhlpl
IP00229184.3	5	0	5	3	5	18	5	3.6	2.9	86.60%	86.60%	1.2	26.65%	Plcb3
IP00135560.1	5	3	2	2	6	18	3	3.6	1.5	45.83%	45.83%	2.3	69.28%	Pmpcb
IP00331631.1	6	5	0	5	2	18	5	3.6	3.2	87.67%	87.67%	2.5	107.85%	Phn
IP00157497.8	5	0	6	7	0	18	5	3.6	3.2	87.67%	87.67%	3.8	87.37%	Polr2a
IP00170013.1	5	4	6	3	0	18	4	3.6	1.0	20.00%	20.00%	3.0	100.00%	Pptc7
IP00653307.1	4	3	3	4	4	18	4	3.6	0.6	17.32%	17.32%	0.6	15.75%	Psma1
IP00153144.3	5	5	0	0	8	18	5	3.6	2.9	86.60%	86.60%	4.6	173.21%	Psmid14
IP00226854.2	3	4	3	6	2	18	3	3.6	0.6	17.32%	17.32%	2.1	56.77%	Ptcd3
IP00230550.1	3	6	5	4	0	18	4	3.6	1.5	32.73%	32.73%	2.6	88.19%	Ptdss1
IP00171981.2	5	8	3	2	0	18	3	3.6	2.5	47.19%	47.19%	1.5	91.65%	Qars
IP00223759.2	8	0	4	3	3	18	3	3.6	4.0	100.00%	100.00%	0.6	17.32%	Scn2b
IP00875929.1	4	6	4	4	0	18	4	3.6	1.2	24.74%	24.74%	2.3	86.60%	Slc25a40
IP00125397.1	3	4	5	3	3	18	3	3.6	1.0	25.00%	25.00%	1.2	31.49%	Ssr4
IP00228907.3	5	6	3	0	4	18	4	3.6	1.5	32.73%	32.73%	2.1	89.21%	Tek
IP00133977.3	3	3	2	5	5	18	3	3.6	0.6	21.65%	21.65%	1.7	43.30%	Them2
IP00129968.2	5	6	4	3	0	18	4	3.6	1.0	20.00%	20.00%	2.1	89.21%	Tmem49

IP00169896.1	3	4	3	3	3	5	18	3	3.6	0.6	17.32%	1.2	31.49%	Tollip
IP00331577.4	5	2	3	3	6	2	18	3	3.6	1.5	45.83%	2.1	56.77%	Ttc35
IP00130733.3	5	2	4	2	2	5	18	4	3.6	1.5	41.66%	1.5	41.66%	Tubg1
IP00117828.1	2	5	6	5	5	0	18	5	3.6	2.1	48.04%	3.2	87.67%	Vac14
IP00338270.5	4	4	4	4	3	2	17	4	3.4	0.0	0.00%	1.0	33.33%	1110014N23RIK
IP00128323.3	3	5	4	4	3	2	17	3	3.4	1.0	25.00%	1.0	33.33%	2700060E02RIK
IP00165794.1	6	5	6	0	0	0	17	5	3.4	0.6	10.19%	3.5	173.21%	AA536749
IP00108372.1	4	5	6	2	2	0	17	4	3.4	1.0	20.00%	3.1	114.56%	Agpat5
IP00129388.1	3	6	6	2	2	0	17	3	3.4	1.7	34.64%	3.1	114.56%	Ankrd47
IP00129774.1	4	5	4	4	4	0	17	4	3.4	0.6	13.32%	2.3	86.60%	Art15
IP00131224.1	5	5	2	2	2	3	17	3	3.4	1.7	43.30%	0.6	24.74%	Aspa
IP00845690.1	3	6	2	2	3	3	17	3	3.4	2.1	56.77%	0.6	21.65%	Atp6v1g2
IP00109904.3	3	5	4	5	5	0	17	4	3.4	1.0	25.00%	2.6	88.19%	Bat1a
IP00134704.1	4	2	4	3	3	4	17	4	3.4	1.2	34.64%	0.6	15.75%	Centg1
IP00131845.1	2	3	2	4	7	3	17	3	3.4	0.6	24.74%	2.6	66.14%	Centg3
IP00459443.5	4	3	3	2	2	5	17	3	3.4	0.6	17.32%	1.5	45.83%	Cgnl1
IP00410802.2	4	5	2	3	3	3	17	3	3.4	1.5	41.66%	0.6	21.65%	Chchd3
IP00117981.1	5	6	4	2	2	0	17	4	3.4	1.0	20.00%	2.0	100.00%	Cox4i2
IP00154056.1	2	2	4	3	3	6	17	3	3.4	1.2	43.30%	1.5	35.25%	Cspg4
IP00330497.4	5	3	4	3	3	2	17	3	3.4	1.0	25.00%	1.0	33.33%	Dao1
IP00467383.2	6	0	3	4	4	4	17	4	3.4	3.0	100.00%	0.6	15.75%	Dgkz
IP00131566.2	3	5	4	5	5	0	17	4	3.4	1.0	25.00%	2.6	88.19%	EG268795
IP00110402.1	3	5	4	5	5	0	17	4	3.4	1.0	25.00%	2.6	88.19%	EG622727
IP00224689.3	0	6	4	4	5	2	17	4	3.4	3.1	91.65%	1.5	41.66%	Gam14
IP00138180.2	3	6	3	3	3	2	17	3	3.4	1.7	43.30%	0.6	21.65%	Gas7
IP00850413.1	7	5	0	2	2	3	17	3	3.4	3.6	90.14%	1.5	91.65%	Gcdh
IP00133110.1	3	4	5	3	3	2	17	3	3.4	1.0	25.00%	1.5	45.83%	Homer3
IP00120115.2	4	6	4	3	3	0	17	4	3.4	1.2	24.74%	2.1	89.21%	Krt15
IP00470956.1	2	6	0	5	5	4	17	4	3.4	3.1	114.56%	2.6	88.19%	Lingo1
IP00653571.1	5	7	5	0	0	0	17	5	3.4	1.2	20.38%	2.9	173.21%	Lman1
IP00336881.1	6	0	3	4	4	4	17	4	3.4	3.0	100.00%	0.6	15.75%	LOC100045062
IP00121190.1	5	2	4	4	4	2	17	4	3.4	1.5	41.66%	1.2	34.64%	LOC676948
IP00830443.1	6	3	3	0	0	5	17	3	3.4	1.7	43.30%	2.5	94.37%	Mag12
IP00421126.5	7	5	3	2	2	0	17	3	3.4	2.0	40.00%	1.5	91.65%	Mcam

IP100111359.2	4	4	4	5	0	17	4	3.4	0.0	0.00%	2.6	88.19%	Mertk
IP100625441.3	6	4	3	2	2	17	3	3.4	1.5	35.25%	0.6	24.74%	Mon2
IP100221616.5	5	3	2	3	4	17	3	3.4	1.5	45.83%	1.0	33.33%	Mpp3
IP100124610.1	3	5	3	3	3	17	3	3.4	1.2	31.49%	0.0	0.00%	Ndufa2
IP100622878.4	4	7	4	2	0	17	4	3.4	1.7	34.64%	2.0	100.00%	Ndufb3
IP100129792.1	4	0	5	6	2	17	4	3.4	2.6	88.19%	2.1	48.04%	Nup93
IP100222457.2	4	8	2	3	0	17	3	3.4	3.1	65.47%	1.5	91.65%	Nupl1
IP100658499.1	9	2	2	0	4	17	2	3.4	4.0	93.26%	2.0	100.00%	Pcsk2
IP100124742.3	3	4	6	0	4	17	4	3.4	1.5	35.25%	3.1	91.65%	Ppid
IP100330176.4	7	4	3	3	0	17	3	3.4	2.1	44.61%	1.7	86.60%	Ppox
IP100132786.1	5	4	3	5	0	17	4	3.4	1.0	25.00%	2.5	94.37%	Prr6
IP100816974.1	4	2	3	4	4	17	4	3.4	1.0	33.33%	0.6	15.75%	Psm6
IP100608044.4	5	3	5	0	4	17	4	3.4	1.2	26.65%	2.6	88.19%	Rgs7bp
IP100125776.1	4	5	4	0	4	17	4	3.4	0.6	13.32%	2.3	86.60%	Rho12
IP100274656.6	4	3	8	2	0	17	3	3.4	2.6	52.92%	4.2	124.90%	Snrpd2
IP100138089.4	5	4	4	4	0	17	4	3.4	0.6	13.32%	2.3	86.60%	Sorbs3
IP100762636.2	4	4	4	3	2	17	4	3.4	0.0	0.00%	1.0	33.33%	Spg7
IP100404595.3	5	4	4	4	0	17	4	3.4	0.6	13.32%	2.3	86.60%	Tm9sf4
IP100315302.5	5	6	3	3	0	17	3	3.4	1.5	32.73%	1.7	86.60%	Ttrap
IP100187338.2	3	5	4	3	2	17	3	3.4	1.0	25.00%	1.0	33.33%	Ubx2
IP100648438.1	6	3	5	2	0	16	3	3.2	1.5	32.73%	2.5	107.85%	2310056P07Rik
IP100153740.1	4	3	3	3	3	16	3	3.2	0.6	17.32%	0.0	0.00%	Akr1b3
IP100468696.3	3	0	6	2	5	16	3	3.2	3.0	100.00%	2.1	48.04%	Aplp2
IP100461277.4	3	4	4	5	0	16	4	3.2	0.6	15.75%	2.6	88.19%	Bcr
IP100411079.2	5	5	2	4	0	16	4	3.2	1.7	43.30%	2.0	100.00%	Cad
IP100338458.2	7	0	7	2	0	16	2	3.2	4.0	86.60%	3.6	120.19%	Cam1
IP100467430.2	3	5	4	4	0	16	4	3.2	1.0	25.00%	2.3	86.60%	Capns1
IP100323114.3	7	4	5	0	0	16	4	3.2	1.5	28.64%	2.9	173.21%	Copa
IP100421223.3	2	4	4	3	3	16	3	3.2	1.2	34.64%	0.6	17.32%	Cops2
IP100762033.2	2	7	3	0	4	16	3	3.2	2.6	66.14%	2.1	89.21%	Cplx1
IP100308706.4	3	2	5	3	3	16	3	3.2	1.5	45.83%	1.2	31.49%	Cred1
IP100875256.1	3	5	6	2	0	16	3	3.2	1.5	32.73%	3.1	114.56%	Cugbp2
IP100137424.2	2	4	4	4	2	16	4	3.2	1.2	34.64%	1.2	34.64%	Ddah2
IP100110760.1	4	5	4	3	0	16	4	3.2	0.6	13.32%	2.1	89.21%	Ddx17

IP100124890.1	7	0	5	0	4	16	4	3.2	3.6	90.14%	2.6	88.19%	Dock3
IP100319135.4	4	4	8	0	0	16	4	3.2	2.3	43.30%	4.6	173.21%	Dpp4
IP100130324.1	3	4	6	3	0	16	3	3.2	1.5	35.25%	3.0	100.00%	Edg1
IP100124979.2	16	0	0	0	0	16	0	3.2	9.2	173.21%	0.0	#DIV/0!	EG627998
IP100225023.3	2	6	6	2	0	16	2	3.2	2.3	49.49%	3.1	114.56%	Fdps
IP100109705.3	6	6	4	4	0	16	4	3.2	1.2	21.65%	2.3	173.21%	Gabra3
IP100114878.1	3	2	2	5	4	16	3	3.2	0.6	24.74%	1.5	41.66%	Gmfb
IP100134820.1	4	5	4	3	0	16	4	3.2	0.6	13.32%	2.1	89.21%	Gmppb
IP100648654.1	5	4	3	4	0	16	4	3.2	1.0	25.00%	2.1	89.21%	Gprc5b
IP100128319.1	4	4	2	3	3	16	3	3.2	1.2	34.64%	0.6	21.65%	Gpx4
IP100877279.1	3	7	4	2	0	16	3	3.2	2.1	44.61%	2.0	100.00%	Hnrph1
IP100128389.3	2	5	5	2	2	16	2	3.2	1.7	43.30%	1.7	57.74%	Htra2
IP100659932.4	4	3	5	4	0	16	4	3.2	1.0	25.00%	2.6	88.19%	Impdh2
IP100224219.1	5	0	2	3	6	16	3	3.2	2.5	107.85%	2.1	56.77%	Inpp4a
IP100464166.2	7	0	0	0	7	16	2	3.2	4.0	173.21%	3.6	120.19%	Insr
IP100551411.5	4	2	3	7	0	16	3	3.2	1.0	33.33%	3.5	105.36%	Itih5
IP100461339.1	5	6	5	0	0	16	5	3.2	0.6	10.83%	2.9	173.21%	Jak1
IP100480558.4	4	0	6	6	0	16	4	3.2	3.1	91.65%	3.5	86.60%	Konc3
IP100649329.3	8	0	4	4	0	16	4	3.2	4.0	100.00%	2.3	86.60%	Krt8
IP100137848.1	5	3	2	6	0	16	3	3.2	1.5	45.83%	3.1	114.56%	Lcp1
IP100828952.1	5	0	9	2	0	16	2	3.2	4.5	96.63%	4.7	128.89%	LOC100039924
IP100421276.1	4	0	4	3	5	16	4	3.2	2.3	86.60%	1.0	25.00%	LOC100048676
IP100349401.3	5	3	4	2	2	16	3	3.2	1.0	25.00%	1.2	43.30%	Lrpap1
IP100230365.5	4	4	4	4	0	16	4	3.2	0.0	0.00%	2.3	86.60%	Lrrc47
IP100114818.1	2	3	3	8	0	16	3	3.2	0.6	21.65%	4.0	110.22%	Ltf
IP100830749.1	2	2	5	4	3	16	3	3.2	1.7	57.74%	1.0	25.00%	Map2k4
IP100113043.1	4	6	6	0	0	16	4	3.2	1.2	21.65%	3.5	173.21%	Mars
IP100869359.1	4	6	2	2	2	16	2	3.2	2.0	50.00%	0.0	0.00%	Naalad2
IP100135512.1	6	2	2	3	3	16	3	3.2	2.3	69.28%	0.6	21.65%	Nfs1
IP100322610.5	5	3	5	0	3	16	3	3.2	1.2	26.65%	2.5	94.37%	Nlr3l1
IP100121767.1	6	6	4	0	0	16	4	3.2	1.2	21.65%	2.3	173.21%	Nipsnap3a
IP100675933.2	3	2	4	4	3	16	3	3.2	1.0	33.33%	0.6	15.75%	Plxnc1
IP100407413.2	8	4	4	0	0	16	4	3.2	2.3	43.30%	2.3	173.21%	Ppic
IP100119224.1	2	6	6	2	0	16	2	3.2	2.3	49.49%	3.1	114.56%	Rps14

IP100131428.1	4	5	4	0	3	16	4	3.2	0.6	13.32%	2.1	89.21%	Sec14I2
IP100649438.1	3	5	6	2	0	16	3	3.2	1.5	32.73%	3.1	114.56%	Slc7a2
IP100330679.2	8	4	0	0	4	16	4	3.2	4.0	100.00%	2.3	173.21%	Slc7a5
IP100395047.1	6	3	4	3	0	16	3	3.2	1.5	35.25%	2.1	89.21%	Stt3a
IP100110684.1	4	2	6	2	2	16	2	3.2	2.0	50.00%	2.3	69.28%	Stx6
IP100108883.3	7	2	3	2	2	16	2	3.2	2.6	66.14%	0.6	24.74%	Suox
IP100125853.1	5	3	4	2	2	16	3	3.2	1.0	25.00%	1.2	43.30%	Tmem1
IP100128826.1	4	4	5	3	0	16	4	3.2	0.6	13.32%	2.5	94.37%	Top1
IP100137400.2	2	4	4	4	2	16	4	3.2	1.2	34.64%	1.2	34.64%	Tpd5l2
IP100122251.1	3	3	5	0	5	16	3	3.2	1.2	31.49%	2.9	86.60%	Trpm3
IP100798576.2	7	4	0	5	0	16	4	3.2	3.5	95.78%	2.9	173.21%	Ttr
IP100125778.3	5	4	5	0	2	16	4	3.2	0.6	12.37%	2.5	107.85%	Txndc12
IP100672213.3	4	0	4	4	4	16	4	3.2	2.3	86.60%	0.0	0.00%	Tyki
IP100875818.1	4	5	4	0	3	16	4	3.2	0.6	13.32%	2.1	89.21%	Wdfy1
IP100415908.4	6	0	2	8	0	16	2	3.2	3.1	114.56%	4.2	124.90%	Ykt6
IP100468236.3	5	0	5	2	3	15	3	3.0	2.9	86.60%	1.5	45.83%	2310022B05Rik
IP100315504.1	5	4	2	2	2	15	2	3.0	1.5	41.66%	0.0	0.00%	2810405K02Rik
IP100755315.5	6	6	3	0	0	15	3	3.0	1.7	34.64%	1.7	173.21%	Aco1
IP100123519.6	7	2	2	2	2	15	2	3.0	2.9	78.73%	0.0	0.00%	Acsl5
IP100776142.1	2	4	0	5	4	15	4	3.0	2.0	100.00%	2.6	88.19%	Adam11
IP100223818.2	2	4	3	4	2	15	3	3.0	1.0	33.33%	1.0	33.33%	AI427122
IP100420480.5	4	2	0	4	5	15	4	3.0	2.0	100.00%	2.6	88.19%	Ankfy1
IP100125771.3	3	6	0	3	3	15	3	3.0	3.0	100.00%	1.7	86.60%	Anxa4
IP100172906.2	4	3	2	2	4	15	3	3.0	1.0	33.33%	1.2	43.30%	Arf6
IP100655041.2	6	2	3	0	4	15	3	3.0	2.1	56.77%	2.1	89.21%	Arsa
IP100283862.6	3	0	6	3	3	15	3	3.0	3.0	100.00%	1.7	43.30%	Birnh
IP100116072.7	4	2	3	4	2	15	3	3.0	1.0	33.33%	1.0	33.33%	Cacng2
IP100170363.1	5	4	2	4	0	15	4	3.0	1.5	41.66%	2.0	100.00%	Cdh11
IP100108811.1	0	6	0	4	5	15	4	3.0	3.5	173.21%	2.6	88.19%	Cds1
IP100112719.1	6	0	4	3	2	15	3	3.0	3.1	91.65%	1.0	33.33%	Chchd4
IP100109603.1	5	5	5	0	0	15	5	3.0	0.0	0.00%	2.9	173.21%	Chid1
IP100322698.7	2	3	4	3	3	15	3	3.0	1.0	33.33%	0.6	17.32%	Cttn
IP100127560.1	3	2	3	4	3	15	3	3.0	0.6	21.65%	0.6	17.32%	Cyb5
IP100125971.1	3	3	7	2	0	15	3	3.0	2.3	53.29%	3.6	120.19%	Degs1

IP00314204.5	5	4	4	0	2	15	4	3.0	0.6	13.32%	2.0	100.00%	Dhcr7
IP00119058.1	2	4	5	4	0	15	4	3.0	1.5	41.66%	2.6	88.19%	EG240853
IP00322150.8	8	0	7	0	0	15	0	3.0	4.4	87.18%	4.0	173.21%	EG268809
IP00222113.2	5	2	5	3	0	15	3	3.0	1.7	43.30%	2.5	94.37%	Elmo1
IP00153468.1	5	3	0	5	2	15	3	3.0	2.5	94.37%	2.5	107.85%	Elmod1
IP00309285.5	3	3	5	0	4	15	3	3.0	1.2	31.49%	2.6	88.19%	Eng
IP00316469.4	4	4	5	2	0	15	4	3.0	0.6	13.32%	2.5	107.85%	Erlin1
IP00133066.3	3	3	2	5	2	15	3	3.0	0.6	21.65%	1.7	57.74%	Exoc3
IP00331524.7	3	2	4	6	0	15	3	3.0	1.0	33.33%	3.1	91.65%	Fabp7
IP00848909.1	5	3	3	2	2	15	3	3.0	1.2	31.49%	0.6	24.74%	Fmo1
IP00124479.2	2	5	4	4	0	15	4	3.0	1.5	41.66%	2.3	86.60%	Grik2
IP00187450.6	7	4	0	4	0	15	4	3.0	3.5	95.78%	2.3	173.21%	Hdac1
IP00112335.1	5	4	3	3	0	15	3	3.0	1.0	25.00%	1.7	86.60%	Iars
IP00124389.1	5	2	2	3	3	15	3	3.0	1.7	57.74%	0.6	21.65%	Iitfg3
IP00140061.1	2	7	2	2	2	15	2	3.0	2.9	78.73%	0.0	0.00%	Lars2
IP00653636.1	6	0	3	4	2	15	3	3.0	3.0	100.00%	1.0	33.33%	Lgi2
IP00119913.1	3	3	3	3	3	15	3	3.0	0.0	0.00%	0.0	0.00%	Lrrc57
IP00664030.3	4	6	0	5	0	15	4	3.0	3.1	91.65%	2.9	173.21%	Man2c1
IP00119320.4	2	6	7	0	0	15	2	3.0	2.6	52.92%	4.0	173.21%	Map2k2
IP00555004.3	4	5	4	2	0	15	4	3.0	0.6	13.32%	2.0	100.00%	Mfsd2
IP00126120.1	2	4	5	0	4	15	4	3.0	1.5	41.66%	2.6	88.19%	Nfu1
IP00880644.1	5	3	5	0	2	15	3	3.0	1.2	26.65%	2.5	107.85%	Nlgn1
IP00652358.5	3	3	3	4	2	15	3	3.0	0.0	0.00%	1.0	33.33%	Nup188
IP00227460.1	5	4	4	2	0	15	4	3.0	0.6	13.32%	2.0	100.00%	Phr1
IP00606944.2	4	4	3	4	0	15	4	3.0	0.6	15.75%	2.1	89.21%	Pisd
IP00398833.1	3	7	5	0	0	15	3	3.0	2.0	40.00%	2.9	173.21%	Plcxd3
IP00134746.5	6	6	0	0	3	15	3	3.0	3.5	86.60%	1.7	173.21%	Pltp
IP00330303.7	3	3	3	6	0	15	3	3.0	0.0	0.00%	3.0	100.00%	Psp1
IP00624883.2	3	3	0	3	6	15	3	3.0	1.7	86.60%	3.0	100.00%	Psm4
IP00411164.4	2	3	4	4	2	15	3	3.0	1.0	33.33%	1.2	34.64%	Psmc6
IP00117416.1	4	3	3	3	2	15	3	3.0	0.6	17.32%	0.6	21.65%	Psmd12
IP00117771.1	7	3	3	2	0	15	3	3.0	2.3	53.29%	1.5	91.65%	Ptpn9
IP00310635.3	3	5	2	3	2	15	3	3.0	1.5	45.83%	0.6	24.74%	Rab12
IP00153813.4	2	3	4	4	2	15	3	3.0	1.0	33.33%	1.2	34.64%	Rab27b

IP00221577.1	5	2	2	3	3	15	3	3.0	1.7	57.74%	0.6	21.65%	Rab31
IP00553357.1	5	4	4	2	0	15	4	3.0	0.6	13.32%	2.0	100.00%	Rbm39
IP00408957.3	0	6	9	0	0	15	0	3.0	4.6	91.65%	5.2	173.21%	Rhd
IP00459331.2	2	4	6	3	0	15	3	3.0	2.0	50.00%	3.0	100.00%	Scamp3
IP00124298.2	6	3	4	2	0	15	3	3.0	1.5	35.25%	2.0	100.00%	Scrib
IP00128699.3	7	0	6	0	2	15	2	3.0	3.8	87.37%	3.1	114.56%	Sec31a
IP00129323.1	3	4	6	0	2	15	3	3.0	1.5	35.25%	3.1	114.56%	Slc25a15
IP00875553.1	6	4	0	2	3	15	3	3.0	3.1	91.65%	1.5	91.65%	Sorcs2
IP00756446.1	0	4	4	7	0	15	4	3.0	2.3	86.60%	3.5	95.78%	Srrm2
IP00120923.2	4	6	5	0	0	15	4	3.0	1.0	20.00%	2.9	173.21%	Stab1
IP00114840.1	0	5	2	4	4	15	4	3.0	2.5	107.85%	1.2	34.64%	Stx7
IP00377731.2	8	0	2	0	5	15	2	3.0	4.2	124.90%	2.5	107.85%	Susd2
IP00453688.1	3	4	6	2	0	15	3	3.0	1.5	35.25%	3.1	114.56%	Tagln2
IP00754723.2	3	2	5	3	2	15	3	3.0	1.5	45.83%	1.5	45.83%	Tceb2
IP00112128.1	3	3	4	0	5	15	3	3.0	0.6	17.32%	2.6	88.19%	Timm10
IP00312113.1	2	3	3	3	4	15	3	3.0	0.6	21.65%	0.6	17.32%	Traf3
IP00122740.2	4	3	4	4	0	15	4	3.0	0.6	15.75%	2.3	86.60%	Trpv2
IP00420172.1	5	3	3	4	0	15	3	3.0	1.2	31.49%	2.1	89.21%	Tspan7
IP00751703.1	5	2	3	5	0	15	3	3.0	1.5	45.83%	2.5	94.37%	Usp7
IP00134378.8	5	2	3	4	0	14	3	2.8	1.5	45.83%	2.1	89.21%	2610101N10Rik
IP00153103.2	6	0	4	4	0	14	4	2.8	3.1	91.65%	2.3	86.60%	4933403F05Rik
IP00453819.2	5	3	0	3	3	14	3	2.8	2.5	94.37%	1.7	86.60%	Abhd3
IP00309207.5	4	0	0	5	5	14	4	2.8	2.3	173.21%	2.9	86.60%	Acp6
IP00469012.3	4	3	3	2	2	14	3	2.8	0.6	17.32%	0.6	24.74%	Agps
IP00759876.2	2	0	8	4	0	14	2	2.8	4.2	124.90%	4.0	100.00%	AI118078
IP00307991.1	4	5	0	2	3	14	3	2.8	2.6	88.19%	1.5	91.65%	Alad
IP00875145.1	3	4	3	4	0	14	3	2.8	0.6	17.32%	2.1	89.21%	Anxa7
IP00226958.3	0	2	5	7	0	14	2	2.8	2.5	107.85%	3.6	90.14%	Appl1
IP00170121.1	5	3	3	3	0	14	3	2.8	1.2	31.49%	1.7	86.60%	Arhgap23
IP00136496.1	3	5	3	3	0	14	3	2.8	1.2	31.49%	1.7	86.60%	Alp6ap2
IP00672924.1	6	3	2	3	0	14	3	2.8	2.1	56.77%	1.5	91.65%	Blvrb
IP00808053.2	2	4	3	3	2	14	3	2.8	1.0	33.33%	0.6	21.65%	Ccbl2
IP00553414.3	6	0	2	4	2	14	2	2.8	3.1	114.56%	1.2	43.30%	Cdc42bpa
IP00454000.3	6	0	4	0	4	14	4	2.8	3.1	91.65%	2.3	86.60%	Comt



IP00136655.1	5	0	4	0	5	14	4	2.8	2.6	88.19%	2.6	88.19%	2.6	88.19%	Ccq5
IP00314205.1	0	6	4	2	2	14	2	2.8	3.1	91.65%	3.1	91.65%	1.2	43.30%	Crb2
IP00319965.3	2	0	7	2	3	14	2	2.8	3.6	120.19%	2.6	66.14%	2.6	66.14%	Crk
IP00230139.5	3	4	4	0	3	14	3	2.8	0.6	15.75%	2.1	89.21%	2.1	89.21%	Crkl
IP00876368.1	5	3	2	4	0	14	3	2.8	1.5	45.83%	2.0	100.00%	2.0	100.00%	Cryab
IP00110659.2	3	0	4	4	3	14	3	2.8	2.1	89.21%	0.6	15.75%	0.6	15.75%	Cspg5
IP00230566.4	5	0	5	4	0	14	4	2.8	2.9	86.60%	2.6	88.19%	2.6	88.19%	Cul4b
IP00134484.1	3	4	2	2	3	14	3	2.8	1.0	33.33%	0.6	24.74%	0.6	24.74%	Dctn4
IP00380739.5	6	4	4	4	0	14	4	2.8	1.2	24.74%	2.3	173.21%	2.3	173.21%	Dnpep
IP00165799.1	5	2	2	2	5	14	2	2.8	1.7	57.74%	2.5	107.85%	2.5	107.85%	Dock1
IP00111320.7	0	0	10	0	4	14	0	2.8	5.8	173.21%	5.0	107.85%	5.0	107.85%	Dpy30
IP00112641.1	3	3	2	2	6	14	3	2.8	0.6	21.65%	3.1	114.56%	3.1	114.56%	Dynlrb1
IP00270376.6	4	5	3	3	2	14	3	2.8	1.0	25.00%	1.5	91.65%	1.5	91.65%	Emb
IP00320157.8	7	4	3	3	0	14	3	2.8	2.1	44.61%	1.7	173.21%	1.7	173.21%	ENSMUSG00000074407
IP00331284.1	5	2	2	2	5	14	2	2.8	1.7	57.74%	2.5	107.85%	2.5	107.85%	Fibp
IP00119478.1	4	4	0	0	6	14	4	2.8	2.3	86.60%	3.5	173.21%	3.5	173.21%	Galc
IP00330161.2	6	6	0	0	2	14	2	2.8	3.5	86.60%	1.2	173.21%	1.2	173.21%	Hexa
IP00322245.3	2	5	4	4	3	14	3	2.8	1.5	41.66%	2.1	89.21%	2.1	89.21%	Hsd11b1
IP00460720.3	4	4	0	0	3	14	3	2.8	2.3	86.60%	1.7	86.60%	1.7	86.60%	Ilk
IP00406464.2	2	2	3	2	5	14	2	2.8	0.6	24.74%	1.5	45.83%	1.5	45.83%	Itgb3
IP00857771.1	3	3	2	3	3	14	3	2.8	0.6	21.65%	0.6	21.65%	0.6	21.65%	Kcnab1
IP00109420.2	0	6	4	4	4	14	4	2.8	3.1	91.65%	2.3	86.60%	2.3	86.60%	Kcnj3
IP00648186.4	3	3	3	3	3	14	3	2.8	0.0	0.00%	0.6	21.65%	0.6	21.65%	Lgi3
IP00154057.3	4	0	5	0	5	14	4	2.8	2.6	88.19%	2.9	86.60%	2.9	86.60%	Macrocl1
IP00133557.1	3	4	3	3	2	14	3	2.8	0.6	17.32%	0.6	24.74%	0.6	24.74%	Nedd4
IP00331314.6	0	3	4	3	4	14	3	2.8	2.1	89.21%	0.6	15.75%	0.6	15.75%	Nit2
IP00480307.1	6	0	5	3	0	14	3	2.8	3.2	87.67%	2.5	94.37%	2.5	94.37%	Nup107
IP00222763.1	3	0	3	5	3	14	3	2.8	1.7	86.60%	1.2	31.49%	1.2	31.49%	Odz2
IP00126184.7	4	3	4	3	0	14	3	2.8	0.6	15.75%	2.1	89.21%	2.1	89.21%	Osbp
IP00114232.1	2	3	4	4	5	14	3	2.8	1.0	33.33%	2.6	88.19%	2.6	88.19%	Palm2-akap2
IP00309262.3	2	6	4	4	2	14	2	2.8	2.0	50.00%	2.0	100.00%	2.0	100.00%	Pelp1
IP00788331.1	5	2	0	0	4	14	3	2.8	2.5	107.85%	2.1	89.21%	2.1	89.21%	Pld3
IP00129685.2	4	4	6	0	0	14	4	2.8	1.2	24.74%	3.5	173.21%	3.5	173.21%	Pm20d1
IP00111902.1	3	4	4	4	3	14	3	2.8	0.6	15.75%	2.1	89.21%	2.1	89.21%	Ppme1

IP00453823.3	5	5	4	0	0	14	4	2.8	0.6	12.37%	2.3	173.21%	Preb
IP00624896.4	5	0	3	3	3	14	3	2.8	2.5	94.37%	0.0	0.00%	Prkaa1
IP00125602.1	4	3	3	4	0	14	3	2.8	0.6	17.32%	2.1	89.21%	Pipsap1
IP00137087.5	3	5	4	0	2	14	3	2.8	1.0	25.00%	2.0	100.00%	Psmid8
IP00720103.1	0	5	0	9	0	14	0	2.8	2.9	173.21%	5.2	173.21%	Pygl
IP00112822.2	2	5	7	0	0	14	2	2.8	2.5	53.93%	4.0	173.21%	Rars
IP00137668.1	2	3	5	4	0	14	3	2.8	1.5	45.83%	2.6	88.19%	Slc4a2
IP00223055.2	2	3	2	5	2	14	2	2.8	0.6	24.74%	1.7	57.74%	Smc1a
IP00123817.1	3	2	4	2	3	14	3	2.8	1.0	33.33%	1.0	33.33%	Stim1
IP00480432.2	5	0	5	2	2	14	2	2.8	2.9	86.60%	1.7	57.74%	Syt12
IP00761607.1	3	2	4	5	0	14	3	2.8	1.0	33.33%	2.6	88.19%	Tanc2
IP00816904.2	6	3	2	3	3	14	3	2.8	2.1	56.77%	1.5	91.65%	Tppp3
IP00170221.2	7	3	0	3	0	13	3	2.6	3.5	105.36%	1.7	173.21%	1810049H19RIK
IP00828271.3	6	2	2	3	0	13	2	2.6	2.3	69.28%	1.5	91.65%	4933425L03RIK
IP00470963.3	5	0	0	2	6	13	2	2.6	2.9	173.21%	3.1	114.56%	Abhd10
IP00118892.6	5	2	3	3	0	13	3	2.6	1.5	45.83%	1.7	86.60%	Acot11
IP00111286.1	3	5	2	0	3	13	3	2.6	1.5	45.83%	1.5	91.65%	Acst3
IP00330186.3	2	0	2	5	4	13	2	2.6	1.2	86.60%	1.5	41.66%	Aldh9a1
IP00323683.2	8	0	3	2	0	13	2	2.6	4.0	110.22%	1.5	91.65%	Anxa11
IP00606508.1	6	2	3	2	2	13	2	2.6	2.1	56.77%	1.5	91.65%	Arfgef2
IP00387192.4	3	0	2	3	5	13	3	2.6	1.5	91.65%	1.5	45.83%	Asph
IP00129908.1	5	3	3	0	2	13	3	2.6	1.2	31.49%	1.5	91.65%	Atp2c1
IP00379876.2	0	7	6	0	0	13	0	2.6	3.8	87.37%	3.5	173.21%	Atp6v0a4
IP00132958.1	3	3	2	5	0	13	3	2.6	0.6	21.65%	2.5	107.85%	Bcas1
IP00110598.1	5	2	3	0	3	13	3	2.6	1.5	45.83%	1.7	86.60%	Begain
IP00468079.3	2	6	3	0	0	13	2	2.6	2.1	56.77%	1.5	91.65%	Cbr4
IP00454030.3	3	3	0	3	4	13	3	2.6	1.7	86.60%	2.1	89.21%	Cdh5
IP00309844.4	4	0	4	2	3	13	3	2.6	2.3	86.60%	1.0	33.33%	Clpb
IP00808443.2	7	0	4	0	2	13	2	2.6	3.5	95.78%	2.0	100.00%	Clstn1
IP00755183.1	3	0	7	3	0	13	3	2.6	3.5	105.36%	3.5	105.36%	Cnpy2
IP00224561.1	8	2	3	0	0	13	2	2.6	3.2	74.18%	1.7	173.21%	Cttnal1
IP00321744.3	3	0	4	4	2	13	3	2.6	2.1	89.21%	1.2	34.64%	D10Bwg1379e
IP00652811.1	3	4	4	0	2	13	3	2.6	0.6	15.75%	2.0	100.00%	D15Wsu169e
IP00404338.2	2	3	6	2	0	13	2	2.6	2.1	56.77%	3.1	114.56%	Dars

IP00464299.6	3	3	2	3	2	13	3	2.6	0.6	21.65%	0.6	24.74%	Dbc1
IP00125687.1	5	0	2	2	4	13	2	2.6	2.5	107.85%	1.2	43.30%	Dguok
IP00127450.1	2	3	0	4	4	13	3	2.6	1.5	91.65%	2.3	86.60%	Diras2
IP00131309.1	3	0	3	5	2	13	3	2.6	1.7	86.60%	1.5	45.83%	Drq2
IP00123295.2	3	3	3	0	4	13	3	2.6	0.0	0.00%	2.1	89.21%	Dync112
IP00312767.1	3	5	0	5	0	13	3	2.6	2.5	94.37%	2.9	173.21%	EG243872
IP00225192.1	3	6	2	2	0	13	2	2.6	2.1	56.77%	1.2	86.60%	Enpp2
IP00323479.2	7	0	3	3	0	13	3	2.6	3.5	105.36%	1.7	86.60%	ENSMUSG00000060566
IP00177205.1	13	0	0	0	0	13	0	2.6	7.5	173.21%	0.0	#DIV/0!	ENSMUSG00000072460
IP00109335.1	0	4	0	9	0	13	0	2.6	2.3	173.21%	5.2	173.21%	Ephb4
IP00129299.3	3	0	3	4	3	13	3	2.6	1.7	86.60%	0.6	17.32%	Erc1
IP00403814.4	6	3	2	2	0	13	2	2.6	2.1	56.77%	1.2	86.60%	F3
IP00284444.5	2	3	3	3	2	13	3	2.6	0.6	21.65%	0.6	21.65%	Fmn2
IP00226726.1	3	0	5	5	0	13	3	2.6	2.5	94.37%	2.9	86.60%	Fmo2
IP00877180.1	3	2	2	3	3	13	3	2.6	0.6	24.74%	0.6	21.65%	Grit
IP00330860.2	2	2	2	3	4	13	2	2.6	0.0	0.00%	1.0	33.33%	Lanc1
IP00623570.3	5	2	0	6	0	13	2	2.6	2.5	107.85%	3.5	173.21%	LOC100043391
IP00226275.2	3	5	0	5	0	13	3	2.6	2.5	94.37%	2.9	173.21%	LOC674419
IP00653345.2	3	5	0	5	0	13	3	2.6	2.5	94.37%	2.9	173.21%	LOC677073
IP00133270.1	5	0	3	2	3	13	3	2.6	2.5	94.37%	0.6	21.65%	Map4k4
IP00121338.1	4	0	4	3	2	13	3	2.6	2.3	86.60%	1.0	33.33%	Mapk8ip3
IP00461469.3	2	3	4	0	4	13	3	2.6	1.0	33.33%	2.3	86.60%	Mlf1
IP00626860.1	2	3	4	4	0	13	3	2.6	1.0	33.33%	2.3	86.60%	Mta1
IP00119283.1	2	4	4	3	0	13	3	2.6	1.2	34.64%	2.1	89.21%	Ncan
IP00869450.1	4	2	3	4	0	13	3	2.6	1.0	33.33%	2.1	89.21%	Ncl
IP00110806.3	0	4	4	5	0	13	4	2.6	2.3	86.60%	2.6	88.19%	Npc1
IP00222749.1	6	3	4	0	0	13	3	2.6	1.5	35.25%	2.3	173.21%	Pam
IP00655040.1	5	2	2	4	0	13	2	2.6	1.7	57.74%	2.0	100.00%	Pon2
IP00469735.4	5	2	2	4	0	13	2	2.6	1.7	57.74%	2.0	100.00%	Prep
IP00378120.2	6	0	3	4	0	13	3	2.6	3.0	100.00%	2.1	89.21%	Prpf19
IP00132278.1	2	3	3	2	3	13	3	2.6	0.6	21.65%	0.6	21.65%	Pipn23
IP00129015.1	5	0	0	5	3	13	3	2.6	2.9	173.21%	2.5	94.37%	Rasgrf2
IP00109501.1	5	2	3	0	3	13	3	2.6	1.5	45.83%	1.7	86.60%	Rnh1
IP00113362.2	0	5	2	3	3	13	3	2.6	2.5	107.85%	0.6	21.65%	Sdk2

IP00109212.3	5	0	4	2	2	13	2	2.6	2.6	88.19%	1.2	43.30%	Selm
IP00469426.6	2	2	2	2	5	13	2	2.6	0.0	0.00%	1.7	57.74%	Serpina1a
IP00138274.1	3	2	5	3	0	13	3	2.6	1.5	45.83%	2.5	94.37%	Sfrs1
IP00128915.4	5	0	5	3	0	13	3	2.6	2.9	86.60%	2.5	94.37%	Sfrs3
IP00653847.1	0	5	5	3	0	13	3	2.6	2.9	86.60%	2.5	94.37%	Sfrs3
IP00408171.4	2	4	2	3	2	13	2	2.6	1.2	43.30%	0.6	24.74%	Sgpl1
IP00116729.2	5	4	0	2	2	13	2	2.6	2.6	88.19%	1.2	86.60%	Slc16a2
IP00121736.1	0	3	5	5	0	13	3	2.6	2.5	94.37%	2.9	86.60%	Slc5a6
IP00128818.2	4	4	3	2	0	13	3	2.6	0.6	15.75%	1.5	91.65%	Slc6a5
IP00224729.1	4	2	4	3	0	13	3	2.6	1.2	34.64%	2.1	89.21%	Smarcc2
IP00875363.1	3	5	2	3	0	13	3	2.6	1.5	45.83%	1.5	91.65%	Smu1
IP00111258.3	3	4	3	3	0	13	3	2.6	0.6	17.32%	1.7	86.60%	Snrpn
IP00124291.1	3	0	3	4	3	13	3	2.6	1.7	86.60%	0.6	17.32%	Spock2
IP00278348.1	4	6	3	3	0	13	3	2.6	1.5	35.25%	1.7	173.21%	Tbc1d10b
IP00110262.4	2	3	2	4	2	13	2	2.6	0.6	24.74%	1.2	43.30%	Tgm2
IP00407951.1	5	3	3	2	0	13	3	2.6	1.2	31.49%	1.5	91.65%	Tmem16k
IP00331555.2	2	2	6	3	0	13	2	2.6	2.3	69.28%	3.0	100.00%	Tmod1
IP00135659.1	2	0	6	0	5	13	2	2.6	3.1	114.56%	3.2	87.67%	Tom40l
IP00626501.3	2	6	0	3	2	13	2	2.6	3.1	114.56%	1.5	91.65%	Tpt1
IP00115599.6	4	2	4	3	0	13	3	2.6	1.2	34.64%	2.1	89.21%	Twf1
IP00875177.1	8	3	0	0	2	13	2	2.6	4.0	110.22%	1.2	173.21%	Vps16
IP00135311.2	6	2	2	2	0	12	2	2.4	2.3	69.28%	1.2	86.60%	1810047C23RIK
IP00322302.1	6	0	6	0	0	12	0	2.4	3.5	86.60%	3.5	173.21%	1810074P20RIK
IP00230113.5	2	2	5	0	3	12	2	2.4	1.7	57.74%	2.5	94.37%	4930402E16RIK
IP00137658.3	3	3	3	3	0	12	3	2.4	0.0	0.00%	1.7	86.60%	4933407N01RIK
IP00330838.6	3	4	2	0	3	12	3	2.4	1.0	33.33%	1.5	91.65%	Abca2
IP00133719.1	0	4	4	4	0	12	4	2.4	2.3	86.60%	2.3	86.60%	Acp2
IP00117731.1	2	3	4	3	0	12	3	2.4	1.0	33.33%	2.1	89.21%	Ak5
IP00225049.1	3	4	2	3	0	12	3	2.4	1.0	33.33%	1.5	91.65%	Ankrd25
IP00323406.1	2	7	3	0	0	12	2	2.4	2.6	66.14%	1.7	173.21%	Aup1
IP00124372.3	3	3	4	0	2	12	3	2.4	0.6	17.32%	2.0	100.00%	Bai3
IP00123624.8	5	3	0	0	4	12	3	2.4	2.5	94.37%	2.3	173.21%	Ccdc90b
IP00225288.3	4	0	3	5	0	12	3	2.4	2.1	89.21%	2.5	94.37%	Cdc5l
IP00875283.1	4	5	0	3	0	12	3	2.4	2.6	88.19%	1.7	173.21%	Chd4

IP100124047.1	3	4	2	0	3	12	3	2.4	1.0	33.33%	1.5	91.65%	Cln3
IP100881051.1	5	2	2	3	0	12	2	2.4	1.7	57.74%	1.5	91.65%	Clnn
IP100849185.1	0	2	2	8	0	12	2	2.4	1.2	86.60%	4.2	124.90%	Cops6
IP100134353.3	5	3	2	0	2	12	2	2.4	1.5	45.83%	1.2	86.60%	Coro1b
IP100465946.1	3	5	4	0	0	12	3	2.4	1.0	25.00%	2.3	173.21%	Cox4nb
IP100515319.2	3	0	7	2	0	12	2	2.4	3.5	105.36%	3.6	120.19%	Dip2a
IP100877240.1	3	3	4	2	0	12	3	2.4	0.6	17.32%	2.0	100.00%	Dnajb11
IP100309183.3	4	2	4	0	2	12	2	2.4	1.2	34.64%	2.0	100.00%	Dock7
IP100109389.1	4	4	4	0	0	12	4	2.4	0.0	0.00%	2.3	173.21%	Ece1
IP100278942.4	0	12	0	0	0	12	0	2.4	6.9	173.21%	0.0	#DIV/0!	EG432987
IP100454081.1	4	0	4	4	0	12	4	2.4	2.3	86.60%	2.3	86.60%	EG668144
IP100649115.1	0	0	7	5	0	12	0	2.4	4.0	173.21%	3.6	90.14%	ENSMUSG00000027694
IP100387427.3	5	0	2	2	3	12	2	2.4	2.5	107.85%	0.6	24.74%	Fads1
IP100761526.3	3	4	3	0	2	12	3	2.4	0.6	17.32%	1.5	91.65%	Fcho2
IP100459742.2	0	4	0	6	2	12	2	2.4	2.3	173.21%	3.1	114.56%	G6pdx
IP100133562.1	4	0	5	0	3	12	3	2.4	2.6	88.19%	2.5	94.37%	Impa1
IP100623897.3	4	3	3	0	2	12	3	2.4	0.6	17.32%	1.5	91.65%	Isoc1
IP100132218.2	2	0	5	3	2	12	2	2.4	2.5	107.85%	1.5	45.83%	Kif1a
IP100753928.1	4	0	4	0	4	12	4	2.4	2.3	86.60%	2.3	86.60%	Kif5a
IP100170297.3	0	4	5	3	0	12	3	2.4	2.6	88.19%	2.5	94.37%	Krt13
IP100170084.1	3	3	2	4	0	12	3	2.4	0.6	21.65%	2.0	100.00%	Mpp7
IP100331614.7	3	4	2	0	3	12	3	2.4	1.0	33.33%	1.5	91.65%	Mirf
IP100108098.1	0	6	2	4	0	12	2	2.4	3.1	114.56%	2.0	100.00%	Novel
IP100331342.6	2	2	2	2	4	12	2	2.4	0.0	0.00%	1.2	43.30%	Pde1a
IP100719884.1	0	6	4	0	2	12	2	2.4	3.1	91.65%	2.0	100.00%	Phactr1
IP100469788.1	6	0	0	3	3	12	3	2.4	3.5	173.21%	1.7	86.60%	Ppa1
IP100331094.3	0	3	6	3	0	12	3	2.4	3.0	100.00%	3.0	100.00%	Ppp1r12a
IP100467495.4	5	0	0	4	3	12	3	2.4	2.9	173.21%	2.1	89.21%	Prps2
IP100605093.2	4	6	0	2	0	12	2	2.4	3.1	91.65%	1.2	173.21%	Plges3
IP100311064.2	5	2	5	0	0	12	2	2.4	1.7	43.30%	2.9	173.21%	Plk2
IP100761693.4	4	2	3	0	3	12	3	2.4	1.0	33.33%	1.7	86.60%	Plprj
IP100128608.2	3	0	2	2	5	12	2	2.4	1.5	91.65%	1.7	57.74%	Rab4b
IP100120943.3	2	4	4	0	2	12	2	2.4	1.2	34.64%	2.0	100.00%	Rap1gap
IP100113243.1	4	8	0	0	0	12	0	2.4	4.0	100.00%	0.0	#DIV/0!	Rps10

IP100653931.1	0	2	4	4	2	12	2	2.4	2.0	100.00%	1.2	34.64%	Rps19
IP100347255.5	2	2	3	3	2	12	2	2.4	0.6	24.74%	0.6	21.65%	Sf3b4
IP100130444.1	0	8	4	0	0	12	0	2.4	4.0	100.00%	2.3	173.21%	Sic25a42
IP100464293.2	4	5	0	3	0	12	3	2.4	2.6	88.19%	1.7	173.21%	Sic6a20
IP100453771.2	3	4	0	0	5	12	3	2.4	2.1	89.21%	2.9	173.21%	Sic9a6
IP100121430.2	3	3	2	2	2	12	2	2.4	0.6	21.65%	0.0	0.00%	Spr
IP100556688.1	2	2	3	5	0	12	2	2.4	0.6	24.74%	2.5	94.37%	Thop1
IP100135277.1	4	4	2	2	0	12	2	2.4	1.2	34.64%	1.2	86.60%	Tmem10
IP100228830.3	6	0	6	0	0	12	0	2.4	3.5	86.60%	3.5	173.21%	Tmem11
IP100113731.2	2	2	3	2	3	12	2	2.4	0.6	24.74%	0.6	21.65%	Tmod3
IP100453692.4	0	0	7	5	0	12	0	2.4	4.0	173.21%	3.6	90.14%	Ube2v1
IP100387491.1	2	3	2	2	3	12	2	2.4	0.6	24.74%	0.6	24.74%	Uqcc
IP100228236.6	4	0	6	2	0	12	2	2.4	3.1	91.65%	3.1	114.56%	Usp6nl
IP100874398.1	0	7	5	0	0	12	0	2.4	3.6	90.14%	2.9	173.21%	Vps25
IP100110224.1	2	3	4	3	0	12	3	2.4	1.0	33.33%	2.1	89.21%	Vit1b
IP100135975.1	4	5	0	3	0	12	3	2.4	2.6	88.19%	1.7	173.21%	Xtrp3s1
IP100464312.4	4	5	2	0	0	11	2	2.2	1.5	41.66%	1.2	173.21%	1700012G19RIK
IP100387333.4	0	4	0	0	7	11	0	2.2	2.3	173.21%	4.0	173.21%	2010309E21RIK
IP100465761.5	2	3	4	2	0	11	2	2.2	1.0	33.33%	2.0	100.00%	2410014A08RIK
IP100396685.1	6	3	2	0	0	11	2	2.2	2.1	56.77%	1.2	173.21%	Acsi4
IP100457976.2	0	11	0	0	0	11	0	2.2	6.4	173.21%	0.0	#DIV/0!	Actn3
IP100153448.3	0	4	7	0	0	11	0	2.2	3.5	95.78%	4.0	173.21%	Adcy8
IP100265701.2	4	0	0	3	4	11	3	2.2	2.3	173.21%	2.1	89.21%	Ahsa1
IP100133612.1	2	2	5	2	0	11	2	2.2	1.7	57.74%	2.5	107.85%	Anxa3
IP100129096.1	4	7	0	0	0	11	0	2.2	3.5	95.78%	0.0	#DIV/0!	Arl1
IP100229600.2	0	0	4	5	2	11	2	2.2	2.3	173.21%	1.5	41.66%	Arl2
IP100132349.3	3	4	4	0	0	11	3	2.2	0.6	15.75%	2.3	173.21%	Arsb
IP100133531.1	5	0	0	2	4	11	2	2.2	2.9	173.21%	2.0	100.00%	Cacybp
IP100463909.3	2	0	4	5	0	11	2	2.2	2.0	100.00%	2.6	88.19%	Car14
IP100121390.1	5	2	0	0	4	11	2	2.2	2.5	107.85%	2.3	173.21%	Ccdc132
IP100474508.1	3	2	0	3	3	11	3	2.2	1.5	91.65%	1.7	86.60%	Ccdc136
IP100420789.1	2	4	0	2	3	11	2	2.2	2.0	100.00%	1.5	91.65%	Cd93
IP100468859.3	6	0	2	0	3	11	2	2.2	3.1	114.56%	1.5	91.65%	Copb1
IP100108960.3	4	0	0	4	3	11	3	2.2	2.3	173.21%	2.1	89.21%	Cops5

PI00469333.1	2	4	2	3	0	11	2	2.2	1.2	43.30%	1.5	91.65%	Cxadr
PI00462949.4	4	5	0	0	2	11	2	2.2	2.6	88.19%	1.2	173.21%	Daglb
PI00881344.1	5	2	0	2	2	11	2	2.2	2.5	107.85%	1.2	86.60%	Ddef1
PI00622160.3	2	3	0	3	3	11	3	2.2	1.5	91.65%	1.7	86.60%	Dnajb4
PI00399943.3	4	4	3	0	0	11	3	2.2	0.6	15.75%	1.7	173.21%	Eef1d
PI00762291.2	0	3	4	2	2	11	2	2.2	2.1	89.21%	1.2	43.30%	Elmo2
PI00312148.3	0	11	0	0	0	11	0	2.2	6.4	173.21%	0.0	#DIV/0!	ENSMUSG00000052469
PI00137243.1	7	0	0	4	0	11	0	2.2	4.0	173.21%	2.3	173.21%	ENSMUSG00000061062
PI00850281.1	4	3	0	0	4	11	3	2.2	2.1	89.21%	2.3	173.21%	Ero1l
PI00463367.2	4	0	2	0	5	11	2	2.2	2.0	100.00%	2.5	107.85%	Ethe1
PI00345373.4	4	0	4	3	0	11	3	2.2	2.3	86.60%	2.1	89.21%	Fmnl2
PI00553419.3	4	5	2	0	0	11	2	2.2	1.5	41.66%	1.2	173.21%	Fvt1
PI00117896.3	3	0	4	4	0	11	3	2.2	2.1	89.21%	2.3	86.60%	Gc
PI00113655.1	6	5	0	0	0	11	0	2.2	3.2	87.67%	0.0	#DIV/0!	Gfm1
PI00116432.1	4	0	3	4	0	11	3	2.2	2.1	89.21%	2.1	89.21%	Gjb6
PI00406596.4	2	4	3	2	0	11	2	2.2	1.0	33.33%	1.5	91.65%	Glr3
PI00117424.1	2	2	4	3	0	11	2	2.2	1.2	43.30%	2.1	89.21%	Grps
PI00556850.1	7	0	0	4	0	11	0	2.2	4.0	173.21%	2.3	173.21%	Gna12
PI00380195.1	2	2	5	0	2	11	2	2.2	1.7	57.74%	2.5	107.85%	Grpel1
PI00225123.3	2	4	2	3	0	11	2	2.2	1.2	43.30%	1.5	91.65%	Herc2
PI00845605.1	3	3	2	3	0	11	3	2.2	0.6	21.65%	1.5	91.65%	Hnrnc
PI00421140.2	6	0	0	3	2	11	2	2.2	3.5	173.21%	1.5	91.65%	Ktbbd11
PI00469123.1	3	2	3	3	0	11	3	2.2	0.6	21.65%	1.7	86.60%	Lars
PI00110426.1	2	2	3	4	0	11	2	2.2	0.6	24.74%	2.1	89.21%	Lbr
PI00228967.4	2	5	0	4	0	11	2	2.2	2.5	107.85%	2.3	173.21%	Lipe
PI00133776.1	3	2	0	3	3	11	3	2.2	1.5	91.65%	1.7	86.60%	LOC100047450
PI00132184.1	0	6	5	0	0	11	0	2.2	3.2	87.67%	2.9	173.21%	Mapbpi
PI00169998.2	0	5	4	0	2	11	2	2.2	2.6	88.19%	2.0	100.00%	Mirps30
PI00117087.1	5	0	4	2	0	11	2	2.2	2.6	88.19%	2.0	100.00%	OTTMUSG00000018617
PI00126172.1	5	2	0	4	0	11	2	2.2	2.5	107.85%	2.3	173.21%	Plod3
PI00312018.6	4	0	3	2	2	11	2	2.2	2.1	89.21%	0.6	24.74%	Prkar1b
PI00230368.1	2	2	3	2	2	11	2	2.2	0.6	24.74%	0.6	24.74%	Psmb5
PI00349032.2	3	3	3	2	0	11	3	2.2	0.0	0.00%	1.5	91.65%	Ptgrn
PI00454035.2	4	0	2	2	3	11	2	2.2	2.0	100.00%	0.6	24.74%	Plk2b

IP100137336.1	4	0	3	4	0	4	0	0	11	3	2.2	2.1	89.21%	2.1	89.21%	Rab11fp5
IP100875492.1	4	0	4	0	3	2.2	2.3	86.60%	11	3	2.2	2.1	89.21%	2.1	89.21%	Rab4a
IP100410796.3	2	2	4	3	0	2.2	1.2	43.30%	11	2	2.2	1.2	89.21%	2.1	89.21%	Rmnd1
IP100853911.1	0	5	3	3	0	2.2	2.5	94.37%	11	3	2.2	2.5	86.60%	1.7	86.60%	Rpl23
IP100317902.3	3	4	2	2	0	2.2	1.0	33.33%	11	2	2.2	1.0	86.60%	1.2	86.60%	Rpl30
IP100121254.1	3	0	4	4	0	2.2	2.1	89.21%	11	3	2.2	2.1	86.60%	2.3	86.60%	Rpl35a
IP100119094.1	0	4	5	2	0	2.2	2.6	88.19%	11	2	2.2	2.6	107.85%	2.5	107.85%	Rpl9
IP100625262.4	11	0	0	0	0	2.2	6.4	173.21%	11	0	2.2	6.4	#DIV/0!	0.0	#DIV/0!	Rps2
IP100380795.4	2	3	4	2	0	2.2	1.0	33.33%	11	2	2.2	1.0	100.00%	2.0	100.00%	Rps20
IP100134506.1	3	3	0	3	2	2.2	1.7	86.60%	11	3	2.2	1.7	91.65%	1.5	91.65%	Sar1a
IP100753595.1	5	0	0	2	4	2.2	2.9	173.21%	11	2	2.2	2.9	100.00%	2.0	100.00%	Sec61a1
IP100225715.7	4	0	4	3	0	2.2	2.3	86.60%	11	3	2.2	2.3	89.21%	2.1	89.21%	Sep15
IP100660262.4	4	0	3	4	0	2.2	2.1	89.21%	11	3	2.2	2.1	89.21%	2.1	89.21%	Sf3b2
IP100877242.1	4	2	2	0	3	2.2	1.2	43.30%	11	2	2.2	1.2	91.65%	1.5	91.65%	Sic12a7
IP100222935.3	0	6	3	2	0	2.2	3.0	100.00%	11	2	2.2	3.0	91.65%	1.5	91.65%	Sic22a8
IP100387261.3	7	0	4	0	0	2.2	3.5	95.78%	11	0	2.2	3.5	173.21%	2.3	173.21%	Smc3
IP100624988.4	3	0	5	3	0	2.2	2.5	94.37%	11	3	2.2	2.5	94.37%	2.5	94.37%	Snrpd3
IP100469184.1	3	3	3	2	0	2.2	0.0	0.00%	11	3	2.2	0.0	91.65%	1.5	91.65%	St13
IP100409035.5	3	2	2	4	0	2.2	0.6	24.74%	11	2	2.2	0.6	100.00%	2.0	100.00%	Stub1
IP100622815.2	0	4	6	3	2	2.2	3.5	173.21%	11	2	2.2	3.5	56.77%	2.1	56.77%	Syt3
IP100620180.3	2	4	5	0	0	2.2	1.5	41.66%	11	2	2.2	1.5	173.21%	2.9	173.21%	Tars
IP100154012.1	0	11	0	0	0	2.2	6.4	173.21%	11	0	2.2	6.4	#DIV/0!	0.0	#DIV/0!	Tcp10c
IP100228106.3	0	4	5	0	2	2.2	2.6	88.19%	11	2	2.2	2.6	107.85%	2.5	107.85%	Them4
IP100663802.1	0	2	3	3	0	2.2	1.5	91.65%	11	3	2.2	1.5	0.00%	0.0	0.00%	Thsd7a
IP100131894.1	4	4	3	0	0	2.2	0.6	15.75%	11	3	2.2	0.6	173.21%	1.7	173.21%	Tmem35
IP100665513.3	2	7	0	0	2	2.2	3.6	120.19%	11	2	2.2	3.6	173.21%	1.2	173.21%	Tmem48
IP100755697.2	3	3	3	2	0	2.2	0.0	0.00%	11	3	2.2	0.0	91.65%	1.5	91.65%	Tpd52
IP100154081.1	3	2	6	0	0	2.2	2.1	56.77%	11	2	2.2	2.1	173.21%	3.5	173.21%	Tspan14
IP100122529.1	5	2	2	2	0	2.2	1.7	57.74%	11	2	2.2	1.7	86.60%	1.2	86.60%	Unc5b
IP100762525.2	5	4	2	0	0	2.2	1.5	41.66%	11	2	2.2	1.5	173.21%	1.2	173.21%	Vcam1
IP100222462.4	3	3	2	3	0	2.2	0.6	21.65%	11	3	2.2	0.6	91.65%	1.5	91.65%	Vps33a
IP100462445.2	2	3	3	3	0	2.2	0.6	21.65%	11	3	2.2	0.6	86.60%	1.7	86.60%	Vps53
IP100675236.1	2	0	3	3	3	2.2	1.5	91.65%	11	3	2.2	1.5	0.00%	0.0	0.00%	Xpnpep1
IP100170027.1	3	3	0	0	4	2.0	1.7	86.60%	10	3	2.0	1.7	173.21%	2.3	173.21%	0610038F07Rik



IP100321477.1	3	0	4	3	0	10	3	2.0	2.1	89.21%	2.1	89.21%	2310008M10Rik
IP100674493.2	6	0	4	0	0	10	0	2.0	3.1	91.65%	2.3	173.21%	2600009E05Rik
IP100222135.3	3	0	3	2	2	10	2	2.0	1.7	86.60%	0.6	24.74%	2610110G12Rik
IP100133122.3	3	3	2	0	2	10	2	2.0	0.6	21.65%	1.2	86.60%	5730446C15Rik
IP100830849.1	5	3	0	2	0	10	2	2.0	2.5	94.37%	1.2	173.21%	8430419L09Rik
IP100109253.1	2	4	4	0	0	10	2	2.0	1.2	34.64%	2.3	173.21%	Abca3
IP100453776.1	0	5	3	2	0	10	2	2.0	2.5	94.37%	1.5	91.65%	Addk4
IP100404653.4	3	0	3	4	0	10	3	2.0	1.7	86.60%	2.1	89.21%	Adcy3
IP100113052.1	3	0	2	5	0	10	2	2.0	1.5	91.65%	2.5	107.85%	Adh5
IP100463126.1	2	0	3	5	0	10	2	2.0	1.5	91.65%	2.5	94.37%	Al314180
IP100123540.1	3	0	2	5	0	10	2	2.0	1.5	91.65%	2.5	107.85%	Anxa1
IP100127237.1	0	0	0	5	5	10	0	2.0	0.0	#DIV/0!	2.9	86.60%	Atp5g1
IP100875944.1	0	0	0	5	5	10	0	2.0	0.0	#DIV/0!	2.9	86.60%	Atp5g2
IP100278498.10	3	4	0	3	0	10	3	2.0	2.1	89.21%	1.7	173.21%	B230339M05Rik
IP100649969.1	2	4	2	2	0	10	2	2.0	1.2	43.30%	1.2	86.60%	BC033915
IP100620325.2	4	2	0	2	2	10	2	2.0	2.0	100.00%	1.2	86.60%	Cab39
IP100275149.4	6	0	2	2	0	10	2	2.0	3.1	114.56%	1.2	86.60%	Caacn1b
IP100187396.2	4	3	0	3	0	10	3	2.0	2.1	89.21%	1.7	173.21%	Caacn2
IP100828535.1	2	2	4	0	2	10	2	2.0	1.2	43.30%	2.0	100.00%	Cbr3
IP100169979.2	2	4	2	2	0	10	2	2.0	1.2	43.30%	1.2	86.60%	Cit
IP100227585.6	3	0	4	3	0	10	3	2.0	2.1	89.21%	2.1	89.21%	Clint1
IP100113386.1	3	2	3	0	2	10	2	2.0	0.6	21.65%	1.5	91.65%	Cnpy4
IP100377299.1	0	5	3	2	0	10	2	2.0	2.5	94.37%	1.5	91.65%	Cntn3
IP100869424.2	0	4	2	2	2	10	2	2.0	2.0	100.00%	0.0	0.00%	Cox15
IP100756257.1	0	4	6	0	0	10	0	2.0	3.1	91.65%	3.5	173.21%	Csnk1d
IP100377496.5	0	7	3	0	0	10	0	2.0	3.5	105.36%	1.7	173.21%	D6Wsu176e
IP100346911.5	4	2	4	0	0	10	2	2.0	1.2	34.64%	2.3	173.21%	Daam2
IP100817029.1	2	4	4	0	0	10	2	2.0	1.2	34.64%	2.3	173.21%	Der1
IP100222228.5	2	4	0	4	0	10	2	2.0	2.0	100.00%	2.3	173.21%	Dhx15
IP100229528.5	4	0	2	4	0	10	2	2.0	2.0	100.00%	2.0	100.00%	Dmn
IP100137731.5	0	0	10	0	0	10	0	2.0	5.8	173.21%	5.8	173.21%	EG666974
IP100127228.3	4	0	4	2	0	10	2	2.0	2.3	86.60%	2.0	100.00%	Eif3c
IP100309437.6	0	6	0	4	0	10	0	2.0	3.5	173.21%	2.3	173.21%	Eif5
IP100225732.4	0	4	3	0	3	10	3	2.0	2.1	89.21%	1.7	86.60%	Enah



PI00457544.1	0	5	2	3	0	10	2	2.0	2.5	107.85%	1.5	91.65%	Pllp
PI00126719.1	3	3	2	2	0	10	2	2.0	0.6	21.65%	1.2	86.60%	Pnpt1
PI00857273.1	2	2	4	0	2	10	2	2.0	1.2	43.30%	2.0	100.00%	Ppp2r5a
PI00620988.1	4	0	0	2	4	10	2	2.0	2.3	173.21%	2.0	100.00%	Prosc
PI00620460.4	4	3	0	0	3	10	3	2.0	2.1	89.21%	1.7	173.21%	Psmd13
PI00110460.2	5	0	0	0	5	10	0	2.0	2.9	173.21%	2.9	173.21%	Ptpmt1
PI00407917.1	4	4	0	0	2	10	2	2.0	2.3	86.60%	1.2	173.21%	Ptprn
PI00116908.1	5	3	0	0	2	10	2	2.0	2.5	94.37%	1.2	173.21%	Rab22a
PI00119111.2	4	0	4	2	0	10	2	2.0	2.3	86.60%	2.0	100.00%	Rab9b
PI00463195.2	3	4	3	0	0	10	3	2.0	0.6	17.32%	1.7	173.21%	Reck
PI00222044.1	2	0	3	3	5	10	2	2.0	1.5	91.65%	2.5	94.37%	Rgs14
PI00318012.6	2	3	0	0	3	10	2	2.0	1.5	91.65%	1.5	91.65%	Rrbp1
PI00270915.4	4	0	4	2	0	10	2	2.0	2.3	86.60%	2.0	100.00%	Sdcbp
PI00133132.1	6	0	2	0	2	10	2	2.0	3.1	114.56%	1.2	86.60%	Sdf2
PI00113713.1	4	4	2	2	0	10	2	2.0	1.2	34.64%	1.2	173.21%	Sdpr
PI00875295.1	3	0	2	2	2	10	2	2.0	1.5	91.65%	0.6	24.74%	Setd7
PI00115604.1	2	0	3	2	3	10	2	2.0	1.5	91.65%	0.6	21.65%	Slc39a10
PI00274140.4	3	0	3	3	4	10	3	2.0	1.7	86.60%	2.1	89.21%	Slc4a3
PI00153136.1	0	5	3	3	2	10	2	2.0	2.5	94.37%	1.5	91.65%	Slc5a3
PI00130343.2	3	0	3	3	0	10	3	2.0	1.7	86.60%	2.1	89.21%	Slc7a10
PI00877215.1	3	4	0	0	3	10	3	2.0	2.1	89.21%	1.7	173.21%	Snx2
PI00221865.1	4	2	2	0	2	10	2	2.0	1.2	43.30%	1.2	86.60%	Sparc
PI00317376.1	5	0	2	3	0	10	2	2.0	2.5	107.85%	1.5	91.65%	Spryd4
PI00463297.4	4	4	0	0	2	10	2	2.0	2.3	86.60%	1.2	173.21%	Sri
PI00135131.2	2	0	3	5	0	10	2	2.0	1.5	91.65%	2.5	94.37%	Stxbp6
PI00283156.1	3	5	0	2	0	10	2	2.0	2.5	94.37%	1.2	173.21%	Tapt1
PI00317794.5	0	4	0	0	3	10	3	2.0	2.3	173.21%	1.7	86.60%	Tmed4
PI00122032.4	6	0	2	2	0	10	2	2.0	3.1	114.56%	1.2	86.60%	Tmem143
PI00318750.4	0	2	3	5	0	10	2	2.0	1.5	91.65%	2.5	94.37%	Uba5
PI00762371.2	4	0	6	0	0	10	0	2.0	3.1	91.65%	3.5	173.21%	Ubtf
PI00113394.5	3	5	2	0	0	10	2	2.0	1.5	45.83%	1.2	173.21%	Ugt8a
PI00126551.1	0	2	3	3	2	10	2	2.0	1.5	91.65%	0.6	21.65%	Vwf
PI00750771.2	6	0	4	0	0	10	0	2.0	3.1	91.65%	2.3	173.21%	Wdfy3
PI00620960.3	0	6	0	0	4	10	0	2.0	3.5	173.21%	2.3	173.21%	Wdr57



IP100625248.3	0	2	4	3	0	9	2	1.8	2.0	100.00%	2.1	89.21%	Heph
IP100132960.2	5	0	4	0	0	9	0	1.8	2.6	88.19%	2.3	173.21%	Igf2r
IP100116744.1	4	2	0	3	0	9	2	1.8	2.0	100.00%	1.7	173.21%	Ikkap
IP100345682.6	2	3	0	4	0	9	2	1.8	1.5	91.65%	2.3	173.21%	Ilf2
IP100857213.1	3	3	3	0	0	9	3	1.8	0.0	0.00%	1.7	173.21%	Ilyna1
IP100469768.2	3	4	0	2	0	9	2	1.8	2.1	89.21%	1.2	173.21%	Itpka
IP100132478.1	0	4	5	0	0	9	0	1.8	2.6	88.19%	2.9	173.21%	Krt16
IP100115475.1	2	0	2	5	0	9	2	1.8	1.2	86.60%	2.5	107.85%	Krt18
IP100131688.2	5	0	2	2	0	9	2	1.8	2.5	107.85%	1.2	86.60%	Leng4
IP100127358.1	4	5	0	0	0	9	0	1.8	2.6	88.19%	0.0	#DIV/0!	LOC100039828
IP100751129.1	4	0	2	3	0	9	2	1.8	2.0	100.00%	1.5	91.65%	Lrrc4b
IP100454019.2	4	0	0	0	5	9	0	1.8	2.3	173.21%	2.9	173.21%	Map2k1ip1
IP100165849.3	0	2	3	0	4	9	2	1.8	1.5	91.65%	2.1	89.21%	Mir16
IP100473215.6	3	2	2	2	0	9	2	1.8	0.6	24.74%	1.2	86.60%	Mir14
IP100126501.1	0	3	4	2	0	9	2	1.8	2.1	89.21%	2.0	100.00%	Mirps22
IP100265471.2	5	0	0	4	0	9	0	1.8	2.9	173.21%	2.3	173.21%	Mtap1s
IP100119886.1	2	0	4	0	3	9	2	1.8	2.0	100.00%	2.1	89.21%	Nkain4
IP100131498.1	3	0	0	4	0	9	2	1.8	1.7	173.21%	2.0	100.00%	Nup62
IP100120344.2	6	3	0	0	0	9	0	1.8	3.0	100.00%	0.0	#DIV/0!	Pak2
IP100275992.3	2	2	3	2	0	9	2	1.8	0.6	24.74%	1.5	91.65%	Pbef1
IP100346420.7	3	2	4	0	0	9	2	1.8	1.0	33.33%	2.3	173.21%	Pcdh10
IP100122973.1	4	0	3	0	2	9	2	1.8	2.1	89.21%	1.5	91.65%	Pdzd8
IP100226729.1	0	3	2	2	2	9	2	1.8	1.5	91.65%	0.0	0.00%	Pic1
IP100379391.2	5	2	2	0	0	9	2	1.8	1.7	57.74%	1.2	173.21%	Poidip2
IP100421124.1	3	2	0	4	0	9	2	1.8	1.5	91.65%	2.3	173.21%	Ppapdc2
IP100169699.4	0	0	0	7	2	9	0	1.8	0.0	#DIV/0!	3.6	120.19%	Ppfia4
IP100133411.1	4	0	0	5	0	9	0	1.8	2.3	173.21%	2.9	173.21%	Prmt8
IP100262303.2	0	4	5	0	0	9	0	1.8	2.6	88.19%	2.9	173.21%	Psma8
IP100319880.1	0	5	4	0	0	9	0	1.8	2.6	88.19%	2.3	173.21%	Ptdss2
IP100351246.3	3	0	3	0	3	9	3	1.8	1.7	86.60%	1.7	86.60%	Ptgs1
IP100229274.1	0	5	4	0	0	9	0	1.8	2.6	88.19%	2.3	173.21%	Pxmp2
IP100409345.3	6	0	3	0	0	9	0	1.8	3.0	100.00%	1.7	173.21%	Rad23b
IP100875742.1	4	0	0	3	2	9	2	1.8	2.3	173.21%	1.5	91.65%	Rbbp9
IP100330480.1	4	3	2	0	0	9	2	1.8	1.0	33.33%	1.2	173.21%	Rpl18

IP00130304.1	4	2	0	3	0	0	0	0	9	2	1.8	2.0	100.00%	1.7	173.21%	Rps11
IP00453792.2	0	2	5	2	0	0	0	0	9	2	1.8	2.5	107.85%	2.5	107.85%	Saps3
IP00671073.3	0	0	6	3	0	0	0	0	9	0	1.8	3.5	173.21%	3.0	100.00%	Sic14a1
IP00229516.1	0	2	4	0	3	0	0	0	9	2	1.8	2.0	100.00%	2.1	89.21%	Smpd2
IP00271059.2	2	3	2	2	0	0	0	0	9	2	1.8	0.6	24.74%	1.2	86.60%	Snx1
IP00421282.6	4	2	3	0	0	0	0	0	9	2	1.8	1.0	33.33%	1.7	173.21%	Six4a
IP00135563.1	3	0	0	3	3	0	0	0	9	3	1.8	1.7	173.21%	1.7	86.60%	Synj1
IP00132148.1	5	4	0	0	0	0	0	0	9	0	1.8	2.6	88.19%	0.0	#DIV/0!	Tardbp
IP00399961.2	0	5	4	0	0	0	0	0	9	0	1.8	2.6	88.19%	2.3	173.21%	Tars2
IP00127176.3	0	2	3	2	2	0	0	0	9	2	1.8	1.5	91.65%	0.6	24.74%	Tcigr1
IP00466999.1	2	0	3	2	2	0	0	0	9	2	1.8	1.5	91.65%	0.6	24.74%	Tfam
IP00387302.4	3	2	2	2	0	0	0	0	9	2	1.8	0.6	24.74%	1.2	86.60%	Txn1
IP00153862.2	0	0	9	0	0	0	0	0	9	0	1.8	5.2	173.21%	5.2	173.21%	Ube1y1
IP00130597.3	2	4	3	0	0	0	0	0	9	2	1.8	1.0	33.33%	1.7	173.21%	Usp39
IP00222382.4	4	5	0	0	0	0	0	0	9	0	1.8	2.6	88.19%	0.0	#DIV/0!	Xpr1
IP00830169.1	3	2	0	3	0	0	0	0	8	2	1.6	1.5	91.65%	1.7	173.21%	2210010C04Rik
IP00816914.1	3	0	3	2	0	0	0	0	8	2	1.6	1.7	86.60%	1.5	91.65%	2610528K11Rik
IP00222263.3	0	0	0	5	3	0	0	0	8	0	1.6	0.0	#DIV/0!	2.5	94.37%	3930401K13Rik
IP00411074.1	3	5	0	0	0	0	0	0	8	0	1.6	2.5	94.37%	0.0	#DIV/0!	4931406C07Rik
IP00277362.6	2	4	2	0	0	0	0	0	8	2	1.6	1.2	43.30%	1.2	173.21%	5730419I09Rik
IP00187288.6	0	3	0	2	0	0	0	0	8	2	1.6	1.7	173.21%	1.5	91.65%	A230051G13Rik
IP00127464.1	3	2	0	3	0	0	0	0	8	2	1.6	1.5	91.65%	1.7	173.21%	Actl6b
IP00775791.1	0	2	2	2	2	0	0	0	8	2	1.6	1.2	86.60%	0.0	0.00%	Adrbk2
IP00320208.3	3	0	3	0	0	0	0	0	8	2	1.6	1.7	86.60%	1.5	91.65%	Aldh3b1
IP00314788.5	2	0	2	2	2	0	0	0	8	2	1.6	1.2	86.60%	0.0	0.00%	Arl6ip1
IP00116843.1	4	0	2	0	0	0	0	0	8	2	1.6	2.0	100.00%	1.2	86.60%	Armet
IP00605842.1	3	3	2	0	0	0	0	0	8	2	1.6	0.6	21.65%	1.2	173.21%	Atp5e
IP00123202.1	2	3	3	0	0	0	0	0	8	2	1.6	0.6	21.65%	1.7	173.21%	Bak1
IP00410756.2	0	4	4	0	0	0	0	0	8	0	1.6	2.3	86.60%	2.3	173.21%	Birc6
IP00172328.3	2	4	0	0	0	0	0	0	8	2	1.6	2.0	100.00%	1.2	173.21%	Cacng7
IP00122494.1	5	0	0	3	0	0	0	0	8	0	1.6	2.9	173.21%	1.7	173.21%	Carhsp1
IP00470162.5	0	5	3	0	0	0	0	0	8	0	1.6	2.5	94.37%	1.7	173.21%	Cav2
IP00114401.1	3	0	3	2	0	0	0	0	8	2	1.6	1.7	86.60%	1.5	91.65%	Ccdc56
IP00311509.1	0	4	4	0	0	0	0	0	8	0	1.6	2.3	86.60%	2.3	173.21%	Cln2







IP00318048.5	2	0	3	3	0	8	2	1.6	1.5	91.65%	1.7	86.60%	Rnpep
IP00654420.1	4	0	2	0	2	8	2	1.6	2.0	100.00%	1.2	86.60%	Robo2
IP00225335.1	4	0	4	0	0	8	0	1.6	2.3	86.60%	2.3	173.21%	Rps18
IP00380162.3	3	0	2	3	0	8	2	1.6	1.5	91.65%	1.5	91.65%	Rps6ka1
IP00153633.1	0	5	0	3	0	8	0	1.6	2.9	173.21%	1.7	173.21%	Scn1b
IP00224957.1	2	0	2	2	2	8	2	1.6	1.2	86.60%	0.0	0.00%	Sdc3
IP00874719.1	4	0	2	2	0	8	2	1.6	2.0	100.00%	1.2	86.60%	Sec11c
IP00313306.5	0	4	2	0	2	8	2	1.6	2.0	100.00%	1.2	86.60%	Sec24c
IP00225761.5	3	0	0	5	0	8	0	1.6	1.7	173.21%	2.9	173.21%	Sfrs14
IP00331507.8	0	5	3	0	0	8	0	1.6	2.5	94.37%	1.7	173.21%	Sic33a1
IP00125992.1	2	0	0	3	3	8	2	1.6	1.2	173.21%	1.7	86.60%	Smcp1
IP00462191.3	3	0	2	3	0	8	2	1.6	1.5	91.65%	1.5	91.65%	Snrpa1
IP00467809.2	0	5	0	0	3	8	0	1.6	2.9	173.21%	1.7	173.21%	Sort1
IP00474407.2	4	2	0	2	0	8	2	1.6	2.0	100.00%	1.2	173.21%	Srp54a
IP00120798.1	0	0	8	0	0	8	0	1.6	4.6	173.21%	4.6	173.21%	Ssr1
IP00380220.2	2	0	3	3	0	8	2	1.6	1.5	91.65%	1.7	86.60%	Stam
IP00165711.3	2	2	4	0	0	8	2	1.6	1.2	43.30%	2.3	173.21%	Sugt1
IP00315488.1	0	4	0	2	2	8	2	1.6	2.3	173.21%	1.2	86.60%	Synpr
IP00387504.1	0	0	4	4	0	8	0	1.6	2.3	173.21%	2.3	86.60%	Tex10
IP00474530.4	0	3	0	2	3	8	2	1.6	1.7	173.21%	1.5	91.65%	Timm22
IP00116746.1	2	2	0	2	2	8	2	1.6	1.2	86.60%	1.2	86.60%	Tmem132b
IP00111600.1	0	0	0	6	2	8	0	1.6	0.0	#DIV/0!	3.1	114.56%	Tmsb4x
IP00114099.2	0	4	0	0	4	8	0	1.6	2.3	173.21%	2.3	173.21%	Tspan9
IP00222443.1	2	2	0	4	0	8	2	1.6	1.2	86.60%	2.3	173.21%	Ttll12
IP00230429.4	0	3	3	2	0	8	2	1.6	1.7	86.60%	1.5	91.65%	Txnl1
IP00311493.5	3	0	0	3	2	8	2	1.6	1.7	173.21%	1.5	91.65%	Uhrf1bp1l
IP00330263.3	4	0	0	4	0	8	0	1.6	2.3	173.21%	2.3	173.21%	Unc84a
IP00753292.3	4	2	2	0	0	8	2	1.6	1.2	43.30%	1.2	173.21%	Vdldr
IP00221767.1	0	2	3	0	3	8	2	1.6	1.5	91.65%	1.7	86.60%	Vps13d
IP00420602.3	2	0	3	3	0	8	2	1.6	1.5	91.65%	1.7	86.60%	Zdhc17
IP00129304.7	0	2	4	0	2	8	2	1.6	2.0	100.00%	2.0	100.00%	Zw10
IP00310502.1	2	2	3	0	0	7	2	1.4	0.6	24.74%	1.7	173.21%	1810035L17Rik
IP00844754.2	3	4	0	0	0	7	0	1.4	2.1	89.21%	0.0	#DIV/0!	4833436C18Rik
IP00344583.2	0	7	0	0	0	7	0	1.4	4.0	173.21%	0.0	#DIV/0!	Abce1



IP100136274.2	2	0	0	3	2	0	0	0	7	2	1.4	1.5	91.65%	1.5	91.65%	191	Gapm1
IP100222750.7	2	3	2	0	2	0	0	0	7	2	1.4	0.6	24.74%	1.2	173.21%	192	Gapvd1
IP100319933.4	3	0	2	2	2	0	0	0	7	2	1.4	1.5	91.65%	1.2	86.60%	193	Gpsm1
IP100110827.1	3	0	0	4	2	0	0	0	7	0	1.4	1.7	173.21%	2.3	173.21%	194	Gsta4
IP100112138.1	0	0	3	2	2	0	0	0	7	2	1.4	1.7	173.21%	0.6	24.74%	195	H2-BI
IP100284925.3	4	0	3	0	0	0	0	0	7	0	1.4	2.1	89.21%	1.7	173.21%	196	Hip1r
IP100315325.1	0	0	0	7	0	0	0	0	7	0	1.4	0.0	#DIV/0!	4.0	173.21%	197	Hist1h1a
IP100471120.2	0	0	3	4	0	0	0	0	7	0	1.4	1.7	173.21%	2.1	89.21%	198	Hnrpr
IP100798610.1	2	0	3	2	2	0	0	0	7	2	1.4	1.5	91.65%	1.5	91.65%	199	Hnrph2
IP100848632.1	3	0	2	0	0	0	0	0	7	2	1.4	1.5	91.65%	1.2	86.60%	200	Itch
IP100330066.5	2	3	2	0	2	0	0	0	7	2	1.4	0.6	24.74%	1.2	173.21%	201	Kcnq2
IP100467435.2	2	3	2	0	0	0	0	0	7	2	1.4	0.6	24.74%	1.2	173.21%	202	Kit
IP100647997.1	3	0	2	0	2	0	0	0	7	2	1.4	1.5	91.65%	1.2	86.60%	203	Kpna3
IP100131674.3	0	0	2	3	2	0	0	0	7	2	1.4	1.2	173.21%	0.6	24.74%	204	Lasp1
IP100351252.4	2	0	3	2	2	0	0	0	7	2	1.4	1.5	91.65%	1.5	91.65%	205	Lnp
IP100284816.4	4	0	3	0	0	0	0	0	7	0	1.4	2.1	89.21%	1.7	173.21%	206	LOC100045493
IP100850771.1	2	3	2	0	0	0	0	0	7	2	1.4	0.6	24.74%	1.2	173.21%	207	LOC100047156
IP100222188.4	4	0	3	0	0	0	0	0	7	0	1.4	2.1	89.21%	1.7	173.21%	208	LOC100048369
IP100742415.1	0	0	0	2	0	0	0	5	7	0	1.4	0.0	#DIV/0!	2.5	107.85%	209	Ly6h
IP100469712.3	3	4	0	0	0	0	0	0	7	0	1.4	2.1	89.21%	0.0	#DIV/0!	210	Mest
IP100125662.2	3	4	0	0	0	0	0	0	7	0	1.4	2.1	89.21%	0.0	#DIV/0!	211	Mgea5
IP100466711.2	0	5	2	0	0	0	0	0	7	0	1.4	2.5	107.85%	1.2	173.21%	212	Mgst1
IP100223437.4	0	4	3	0	0	0	0	0	7	0	1.4	2.1	89.21%	1.7	173.21%	213	Mlistd2
IP100876478.1	0	0	3	4	0	0	0	0	7	0	1.4	1.7	173.21%	2.1	89.21%	214	Mobb
IP100340103.1	2	3	2	0	0	0	0	0	7	2	1.4	0.6	24.74%	1.2	173.21%	215	Mirp19
IP100881328.1	5	0	0	0	0	0	0	2	7	0	1.4	2.9	173.21%	1.2	173.21%	216	Mirp37
IP100117083.1	2	0	0	2	2	0	0	3	7	2	1.4	1.2	173.21%	1.5	91.65%	217	Mirps9
IP100113143.1	2	0	5	0	0	0	0	0	7	0	1.4	2.5	107.85%	2.9	173.21%	218	Mtmr2
IP100230355.1	2	0	3	2	0	0	0	0	7	2	1.4	1.5	91.65%	1.5	91.65%	219	Mycbp
IP100460127.2	0	0	0	7	0	0	0	0	7	0	1.4	0.0	#DIV/0!	4.0	173.21%	220	Mylpf
IP100229810.4	3	0	2	2	0	0	0	0	7	2	1.4	1.5	91.65%	1.2	86.60%	221	Nola2
IP100762140.1	0	2	2	0	0	0	0	3	7	2	1.4	1.2	86.60%	1.5	91.65%	222	Nup205
IP100471265.2	0	2	2	3	0	0	0	0	7	2	1.4	1.2	86.60%	1.5	91.65%	223	Nup54
IP100314989.2	2	0	3	0	0	0	0	2	7	2	1.4	1.5	91.65%	1.5	91.65%	224	Ocr1

IP100310627.3	3	0	2	0	0	2	0	0	2	0	0	2	1.4	1.5	91.65%	1.2	86.60%	Pacin3
IP100108143.1	2	0	2	0	0	3	0	0	2	0	0	2	1.4	1.2	86.60%	1.5	91.65%	Pak7
IP100227157.5	2	2	0	3	0	0	0	0	0	0	0	0	1.4	1.2	86.60%	1.7	173.21%	Pbxip1
IP100137172.1	3	0	2	2	0	0	0	0	0	0	0	0	1.4	1.5	91.65%	1.2	86.60%	Pepe
IP100649880.2	0	3	4	0	0	0	0	0	0	0	0	0	1.4	2.1	89.21%	2.3	173.21%	Pla2g7
IP100468204.1	0	3	0	4	0	0	0	0	0	0	0	0	1.4	1.7	173.21%	2.3	173.21%	Purb
IP100132089.2	0	3	2	2	0	0	0	0	0	0	0	0	1.4	1.5	91.65%	1.2	86.60%	Pycl
IP100120155.1	2	0	0	0	3	0	0	0	0	0	0	0	1.4	1.2	173.21%	1.5	91.65%	Ranbp9
IP100330167.4	2	0	0	0	3	0	0	0	0	0	0	0	1.4	1.2	173.21%	1.5	91.65%	Rnps1
IP100115591.1	4	0	3	0	0	0	0	0	0	0	0	0	1.4	2.1	89.21%	1.7	173.21%	Rpl11
IP100464151.3	0	0	7	0	0	0	0	0	0	0	0	0	1.4	4.0	173.21%	4.0	173.21%	Rpl23a
IP100228497.1	0	5	0	0	2	0	0	0	0	0	0	0	1.4	2.9	173.21%	1.2	173.21%	Rpl24
IP100457680.3	5	0	0	0	2	0	0	0	0	0	0	0	1.4	2.9	173.21%	1.2	173.21%	Rps24
IP100462005.1	2	3	2	0	0	0	0	0	0	0	0	0	1.4	0.6	24.74%	1.2	173.21%	Rps8
IP100853896.1	3	4	0	0	0	0	0	0	0	0	0	0	1.4	2.1	89.21%	0.0	#DIV/0!	Rsu1
IP100123744.1	0	4	3	0	0	0	0	0	0	0	0	0	1.4	2.1	89.21%	1.7	173.21%	Ryr1
IP100330866.1	2	0	0	0	3	0	0	0	0	0	0	0	1.4	1.2	173.21%	1.5	91.65%	Saps2
IP100172197.2	0	3	4	0	0	0	0	0	0	0	0	0	1.4	2.1	89.21%	2.3	173.21%	Sardh
IP100113563.2	3	0	2	2	0	0	0	0	0	0	0	0	1.4	1.5	91.65%	1.2	86.60%	Scyl2
IP100120674.2	0	2	3	0	0	0	0	0	0	0	0	0	1.4	1.5	91.65%	1.5	91.65%	Sgsm1
IP100153234.2	3	0	2	2	0	0	0	0	0	0	0	0	1.4	1.5	91.65%	1.2	86.60%	Sgta
IP100126011.1	5	0	2	0	0	0	0	0	0	0	0	0	1.4	2.5	107.85%	1.2	173.21%	Slc7a3
IP100762547.2	2	0	2	0	0	0	0	0	0	0	0	0	1.4	1.2	86.60%	1.5	91.65%	Slk
IP100755042.1	0	0	0	0	7	0	0	0	0	0	0	0	1.4	0.0	#DIV/0!	4.0	173.21%	Smarca5
IP100462453.4	3	2	2	0	0	0	0	0	0	0	0	0	1.4	0.6	24.74%	1.2	173.21%	Smpd1
IP100129489.1	0	0	4	3	0	0	0	0	0	0	0	0	1.4	2.3	173.21%	2.1	89.21%	Snx6
IP100462975.3	3	2	2	0	0	0	0	0	0	0	0	0	1.4	0.6	24.74%	1.2	173.21%	Spag9
IP100116285.2	3	0	0	0	2	0	0	0	0	0	0	0	1.4	1.7	173.21%	1.2	86.60%	Spq20
IP100387245.1	3	0	4	0	0	0	0	0	0	0	0	0	1.4	2.1	89.21%	2.3	173.21%	Spns1
IP100605141.2	3	2	0	0	0	0	0	0	0	0	0	0	1.4	1.5	91.65%	1.2	173.21%	Strn
IP100282957.4	5	0	0	0	0	0	0	0	0	0	0	0	1.4	2.9	173.21%	1.2	173.21%	Strn3
IP100858037.1	0	0	2	0	0	0	0	0	0	0	0	0	1.4	1.2	173.21%	2.5	107.85%	Syncrip
IP100119959.1	2	2	0	0	3	0	0	0	0	0	0	0	1.4	1.2	86.60%	1.7	173.21%	Syng3
IP100659001.1	3	2	0	0	2	0	0	0	0	0	0	0	1.4	1.5	91.65%	1.2	173.21%	Tmem132a











IP100127989.1	2	0	2	2	0	6	2	1.2	1.2	1.2	86.60%	1.2	86.60%	Safb
IP100762273.1	3	3	0	0	0	6	0	1.7	1.2	1.7	#DIV/0!	0.0	#DIV/0!	Sar1b
IP100458351.2	2	2	2	0	0	6	2	0.0	1.2	0.0	173.21%	1.2	173.21%	Scamp2
IP100377298.2	4	2	0	0	0	6	0	2.0	1.2	2.0	100.00%	0.0	#DIV/0!	Seg3
IP100225318.3	4	0	0	0	2	6	0	2.3	1.2	2.3	173.21%	1.2	173.21%	Scoc
IP100169524.2	3	3	0	0	0	6	0	1.7	1.2	1.7	86.60%	0.0	#DIV/0!	Sec23b
IP100515622.1	2	0	4	0	0	6	0	2.0	1.2	2.0	100.00%	2.3	173.21%	Sec24b
IP100119725.1	3	0	0	0	3	6	0	1.7	1.2	1.7	173.21%	1.7	173.21%	Sez6l
IP100128154.1	3	0	0	3	0	6	0	1.7	1.2	1.7	173.21%	1.7	173.21%	Sirt5
IP100466645.3	4	2	0	0	0	6	0	2.0	1.2	2.0	100.00%	0.0	#DIV/0!	Sic20a2
IP100648888.1	2	0	0	2	2	6	2	1.2	1.2	1.2	173.21%	1.2	86.60%	Sic7a14
IP100378224.2	2	4	0	0	0	6	0	2.0	1.2	2.0	100.00%	0.0	#DIV/0!	Sico1a4
IP100621272.3	2	2	2	0	0	6	2	0.0	1.2	0.0	0.00%	1.2	173.21%	Simap
IP100606142.3	2	0	0	4	0	6	0	1.2	1.2	1.2	173.21%	2.3	173.21%	Snx4
IP100117994.1	3	0	0	3	0	6	0	1.7	1.2	1.7	173.21%	1.7	173.21%	Srm
IP100109313.1	3	0	0	0	3	6	0	1.7	1.2	1.7	173.21%	1.7	173.21%	Sk39
IP100649324.2	2	0	2	0	2	6	2	1.2	1.2	1.2	86.60%	1.2	86.60%	Sult4a1
IP100378671.3	2	2	2	0	0	6	2	0.0	1.2	0.0	0.00%	1.2	173.21%	Surf1
IP100136915.2	0	0	0	0	6	6	0	0.0	1.2	0.0	#DIV/0!	3.5	173.21%	Tmem9b
IP100877236.1	0	2	0	4	0	6	0	1.2	1.2	1.2	173.21%	2.3	173.21%	Tra2a
IP100662189.2	2	2	0	2	0	6	2	1.2	1.2	1.2	86.60%	1.2	173.21%	Trappo4
IP100133700.2	6	0	0	0	0	6	0	3.5	1.2	3.5	173.21%	0.0	#DIV/0!	Try10
IP100473445.1	2	2	0	0	2	6	2	1.2	1.2	1.2	86.60%	1.2	173.21%	Ube3c
IP100849403.1	3	0	3	0	0	6	0	1.7	1.2	1.7	86.60%	1.7	173.21%	Vamp7
IP100122227.1	0	2	2	2	0	6	2	1.2	1.2	1.2	86.60%	1.2	86.60%	Vcpi1
IP100132942.1	0	4	2	0	0	6	0	2.0	1.2	2.0	100.00%	1.2	173.21%	Vps37c
IP100135345.3	0	4	0	2	0	6	0	2.3	1.2	2.3	173.21%	1.2	173.21%	Vtn
IP100122562.3	2	2	0	2	0	6	2	1.2	1.2	1.2	86.60%	1.2	173.21%	Wdr26
IP100111981.1	0	0	3	0	3	6	0	1.7	1.2	1.7	173.21%	1.7	86.60%	Wipi2
IP100652397.1	4	0	2	0	0	6	0	2.0	1.2	2.0	100.00%	1.2	173.21%	Wwp1
IP100420290.2	0	2	2	2	0	6	2	1.2	1.2	1.2	86.60%	1.2	86.60%	Zdhhc20
IP100749628.1	3	0	0	2	0	5	0	1.7	1.0	1.7	173.21%	1.2	173.21%	1110007L15RIK
IP100131535.1	0	0	3	0	2	5	0	1.7	1.0	1.7	173.21%	1.5	91.65%	1300012G16RIK
IP100128202.1	0	3	0	0	0	5	0	1.7	1.0	1.7	173.21%	1.2	173.21%	1810007P19RIK











IP100109932.1	0	2	0	3	0	0	5	0	1.0	1.5	91.65%	1.7	173.21%	Zyx
IP100555113.2	0	0	2	2	0	4	4	0	0.8	1.2	173.21%	1.2	86.60%	1110012J17RIK
IP100312501.3	4	0	0	0	0	4	4	0	0.8	2.3	173.21%	0.0	#DIV/0!	1110067D22RIK
IP100230415.5	0	2	0	2	0	4	4	0	0.8	1.2	86.60%	1.2	173.21%	1500032L24RIK
IP100222107.5	4	0	0	0	0	4	4	0	0.8	2.3	173.21%	0.0	#DIV/0!	1700009N14RIK
IP100421179.1	2	0	2	2	0	4	4	0	0.8	1.2	86.60%	1.2	173.21%	2310003L22RIK
IP100222549.6	0	0	2	2	0	4	4	0	0.8	1.2	173.21%	1.2	86.60%	2310066E14RIK
IP100122413.2	0	0	2	2	0	4	4	0	0.8	1.2	173.21%	1.2	86.60%	2310067B10RIK
IP100122451.1	2	0	0	0	2	4	4	0	0.8	1.2	173.21%	1.2	173.21%	2610301G19RIK
IP100471006.1	2	0	0	0	2	4	4	0	0.8	1.2	173.21%	1.2	173.21%	2610510H03RIK
IP100755700.2	2	2	0	0	0	4	4	0	0.8	1.2	86.60%	0.0	#DIV/0!	5830411G16RIK
IP100672748.4	0	2	0	0	2	4	4	0	0.8	1.2	173.21%	1.2	173.21%	6430517E21RIK
IP100458337.4	0	2	0	0	2	4	4	0	0.8	1.2	173.21%	1.2	173.21%	Aatk
IP100556893.3	2	0	2	2	0	4	4	0	0.8	1.2	86.60%	1.2	173.21%	Abca12
IP100108454.2	0	2	0	2	0	4	4	0	0.8	1.2	86.60%	1.2	173.21%	Abcb10
IP100458851.2	2	0	0	0	2	4	4	0	0.8	1.2	173.21%	1.2	173.21%	Abcb9
IP100122362.1	0	4	0	0	0	4	4	0	0.8	2.3	173.21%	0.0	#DIV/0!	Acot8
IP100831153.1	2	0	2	0	0	4	4	0	0.8	1.2	86.60%	1.2	173.21%	Accs2
IP100135561.3	0	0	4	0	0	4	4	0	0.8	2.3	173.21%	2.3	173.21%	Afg311
IP100408894.3	2	2	0	0	0	4	4	0	0.8	1.2	86.60%	0.0	#DIV/0!	Alg1
IP100406045.3	2	2	0	0	0	4	4	0	0.8	1.2	86.60%	0.0	#DIV/0!	Alg11
IP100754562.1	4	0	0	0	0	4	4	0	0.8	2.3	173.21%	0.0	#DIV/0!	Amfr
IP100226464.4	2	2	0	0	0	4	4	0	0.8	1.2	86.60%	0.0	#DIV/0!	Amt
IP100758066.2	2	0	2	0	2	4	4	0	0.8	1.2	173.21%	1.2	173.21%	Anln
IP100322431.9	2	2	0	0	0	4	4	0	0.8	1.2	86.60%	0.0	#DIV/0!	Ap3b1
IP100330155.4	2	2	0	0	0	4	4	0	0.8	1.2	86.60%	0.0	#DIV/0!	Ap1p1
IP100312468.5	2	0	2	2	0	4	4	0	0.8	1.2	86.60%	1.2	173.21%	Arhgap21
IP100117569.3	0	0	2	2	0	4	4	0	0.8	1.2	173.21%	1.2	86.60%	Arhgap26
IP100122831.5	2	0	0	0	2	4	4	0	0.8	1.2	173.21%	1.2	173.21%	Asphd1
IP100322562.5	0	0	2	2	0	4	4	0	0.8	1.2	173.21%	1.2	86.60%	Atxn10
IP100230124.5	0	2	0	2	0	4	4	0	0.8	1.2	86.60%	1.2	173.21%	Atxn21
IP100420245.1	0	2	0	2	0	4	4	0	0.8	1.2	86.60%	1.2	173.21%	AU022870
IP100408985.2	2	0	0	0	2	4	4	0	0.8	1.2	173.21%	1.2	173.21%	B230208H17RIK
IP100874641.1	2	0	2	0	0	4	4	0	0.8	1.2	173.21%	1.2	173.21%	Bcap29

IP00123296.1	2	0	0	0	0	0	0	0	0.8	1.2	173.21%	1.2	173.21%	Bcl2l13
IP00131502.3	2	0	2	0	0	0	0	0.8	1.2	86.60%	1.2	173.21%	Bpgm	
IP00132765.2	4	0	0	0	0	0	0	0.8	2.3	173.21%	0.0	#DIV/0!	C530043G21Rik	
IP00857151.1	0	0	0	4	0	0	0	0.8	0.0	#DIV/0!	2.3	173.21%	Cbx3	
IP00356888.4	0	2	0	0	2	0	0	0.8	1.2	173.21%	1.2	173.21%	Cdh15	
IP00118904.1	0	2	0	0	2	0	0	0.8	1.2	173.21%	1.2	173.21%	Cdk5rap3	
IP00230119.5	0	0	0	0	0	4	0	0.8	0.0	#DIV/0!	2.3	173.21%	Chmp5	
IP00119124.2	2	0	2	0	0	0	0	0.8	1.2	86.60%	1.2	173.21%	Clhd1	
IP0046200.3	2	0	2	0	0	0	0	0.8	1.2	86.60%	1.2	173.21%	Clec2d	
IP00308294.3	4	0	0	0	0	0	0	0.8	2.3	173.21%	0.0	#DIV/0!	Cnrip1	
IP00856927.1	0	0	2	0	2	0	0	0.8	1.2	173.21%	1.2	86.60%	Col6a1	
IP00323819.3	0	0	2	0	2	0	0	0.8	1.2	173.21%	1.2	86.60%	Coq6	
IP00130095.1	2	0	0	0	0	2	0	0.8	1.2	173.21%	1.2	173.21%	Coq9	
IP00648998.3	0	0	2	0	0	2	0	0.8	1.2	173.21%	1.2	86.60%	Cox18	
IP00653254.2	4	0	0	0	0	0	0	0.8	2.3	173.21%	0.0	#DIV/0!	Ctbp2	
IP00230283.5	2	0	0	2	0	0	0	0.8	1.2	86.60%	1.2	173.21%	Cyp2s1	
IP00221775.1	0	0	4	0	0	0	0	0.8	2.3	173.21%	2.3	173.21%	Dak	
IP00265219.4	0	0	0	0	0	0	0	0.8	0.0	#DIV/0!	2.3	173.21%	Ddx3y	
IP00170145.5	0	0	4	0	0	0	0	0.8	2.3	173.21%	2.3	173.21%	Ddx47	
IP00118739.3	2	0	0	0	0	0	0	0.8	1.2	86.60%	0.0	#DIV/0!	Dnahc8	
IP00459353.2	2	0	0	0	2	0	0	0.8	1.2	173.21%	1.2	173.21%	Dnajb1	
IP00165716.4	2	0	0	0	2	0	0	0.8	1.2	173.21%	1.2	173.21%	Dpp9	
IP00313824.2	2	0	2	0	0	0	0	0.8	1.2	86.60%	1.2	173.21%	Dsp	
IP00162932.1	0	0	4	0	0	0	0	0.8	2.3	173.21%	2.3	173.21%	EG384525	
IP00226416.3	2	0	0	0	0	2	0	0.8	1.2	173.21%	1.2	173.21%	EG434394	
IP00480341.2	2	0	0	0	0	2	0	0.8	1.2	173.21%	1.2	173.21%	EG436081	
IP00123131.3	0	2	0	0	0	0	0	0.8	1.2	86.60%	1.2	173.21%	EG639162	
IP00663587.1	4	0	0	0	0	0	0	0.8	2.3	173.21%	0.0	#DIV/0!	EG666274	
IP00173343.3	0	4	0	0	0	0	0	0.8	2.3	173.21%	0.0	#DIV/0!	Eif4g1	
IP00828628.2	0	0	2	0	0	0	0	0.8	1.2	173.21%	1.2	86.60%	Eifm2	
IP00469253.6	2	2	0	0	0	0	0	0.8	1.2	86.60%	0.0	#DIV/0!	Elp3	
IP00121149.1	0	0	2	0	0	2	0	0.8	1.2	173.21%	1.2	86.60%	ENSMUSG00000072758	
IP00464316.2	0	0	4	0	0	0	0	0.8	2.3	173.21%	2.3	173.21%	ENSMUSG00000078180	
IP00874334.1	0	4	0	0	0	0	0	0.8	2.3	173.21%	0.0	#DIV/0!	Epdr1	

















































PI00470138.4	0	0	0	2	0	0	0	0	0	0	0.4	1.2	173.21%	1.2	173.21%	173.21%	Pht3
PI00309419.2	0	0	0	0	2	0	0	0	0	0	0.4	0.0	#DIV/0!	1.2	173.21%	173.21%	Phip
PI00659860.2	0	0	0	0	2	0	0	0	0	0	0.4	0.0	#DIV/0!	1.2	173.21%	173.21%	Phka1
PI00313492.3	0	0	0	2	0	0	0	0	0	0	0.4	1.2	173.21%	1.2	173.21%	173.21%	Phldb1
PI00229726.2	2	0	0	0	0	0	0	0	0	0	0.4	1.2	173.21%	0.0	#DIV/0!	#DIV/0!	Pik3c2a
PI00330437.1	2	0	0	0	0	0	0	0	0	0	0.4	1.2	173.21%	0.0	#DIV/0!	#DIV/0!	Plekhh1
PI00113427.2	0	0	0	2	0	0	0	0	0	0	0.4	1.2	173.21%	1.2	173.21%	173.21%	Pnpla7
PI00458573.1	0	0	0	2	0	0	0	0	0	0	0.4	1.2	173.21%	1.2	173.21%	173.21%	Pogz
PI00315948.3	0	2	0	0	0	0	0	0	0	0	0.4	1.2	173.21%	0.0	#DIV/0!	#DIV/0!	Poldip3
PI00400411.3	0	0	0	0	2	0	0	0	0	0	0.4	1.2	173.21%	1.2	173.21%	173.21%	Polr2b
PI00124052.1	0	0	0	2	0	0	0	0	0	0	0.4	1.2	173.21%	1.2	173.21%	173.21%	Ppfbp1
PI00461857.2	0	2	0	0	0	0	0	0	0	0	0.4	1.2	173.21%	0.0	#DIV/0!	#DIV/0!	Ppp6c
PI00453815.3	2	0	0	0	0	0	0	0	0	0	0.4	1.2	173.21%	0.0	#DIV/0!	#DIV/0!	Prp
PI00323130.3	2	0	0	0	0	0	0	0	0	0	0.4	1.2	173.21%	0.0	#DIV/0!	#DIV/0!	Prkaa2
PI00471277.1	2	0	0	0	0	0	0	0	0	0	0.4	1.2	173.21%	0.0	#DIV/0!	#DIV/0!	Prkcd
PI00221932.1	0	0	0	2	0	0	0	0	0	0	0.4	1.2	173.21%	1.2	173.21%	173.21%	Prkg2
PI00114044.1	0	0	0	2	0	0	0	0	0	0	0.4	1.2	173.21%	1.2	173.21%	173.21%	Pripf3
PI00468346.1	2	0	0	0	0	0	0	0	0	0	0.4	1.2	173.21%	0.0	#DIV/0!	#DIV/0!	Pripf31
PI00224294.2	0	0	0	0	2	0	0	0	0	0	0.4	0.0	#DIV/0!	1.2	173.21%	173.21%	Prss2
PI00785267.1	0	0	0	0	0	0	0	0	0	0	0.4	0.0	#DIV/0!	1.2	173.21%	173.21%	Prune
PI00137439.1	0	0	0	0	0	0	0	2	0	0	0.4	0.0	#DIV/0!	1.2	173.21%	173.21%	Psemb7
PI00132644.4	0	0	0	0	0	0	0	2	0	0	0.4	0.0	#DIV/0!	1.2	173.21%	173.21%	Psme2
PI00816921.2	0	0	0	0	0	0	0	2	0	0	0.4	0.0	#DIV/0!	1.2	173.21%	173.21%	Psph
PI00119807.1	0	0	0	0	0	0	0	0	0	0	0.4	1.2	173.21%	1.2	173.21%	173.21%	Ptgds
PI00281011.7	0	0	0	2	0	0	0	0	0	0	0.4	1.2	173.21%	1.2	173.21%	173.21%	Ptprb
PI00118096.2	2	0	0	0	0	0	0	0	0	0	0.4	1.2	173.21%	0.0	#DIV/0!	#DIV/0!	Ptprt
PI00279494.6	0	0	0	2	0	0	0	0	0	0	0.4	1.2	173.21%	1.2	173.21%	173.21%	Puf60
PI00229893.1	0	0	0	2	0	0	0	0	0	0	0.4	1.2	173.21%	1.2	173.21%	173.21%	Pxn
PI00880411.1	2	0	0	0	0	0	0	0	0	0	0.4	1.2	173.21%	0.0	#DIV/0!	#DIV/0!	Rab11fp1
PI00118413.2	2	0	0	0	0	0	0	0	0	0	0.4	1.2	173.21%	0.0	#DIV/0!	#DIV/0!	Rab11fp2
PI00226952.4	0	0	0	0	0	0	0	2	0	0	0.4	0.0	#DIV/0!	1.2	173.21%	173.21%	Rab23
PI00114407.2	0	2	0	0	0	0	0	0	0	0	0.4	1.2	173.21%	0.0	#DIV/0!	#DIV/0!	Rab711
PI00221741.1	0	2	0	0	0	0	0	0	0	0	0.4	1.2	173.21%	0.0	#DIV/0!	#DIV/0!	Rab9
PI00380082.1	0	0	0	2	0	0	0	0	0	0	0.4	1.2	173.21%	1.2	173.21%	173.21%	Rabgap1













## Supplementary Table 2. Minimum Information About a Proteomics Experiment

### MIAPE v2.24 2008-05-28

#### 1. General Features – (a) Global Descriptors

Date Stamp	2009-03-18
Responsible Person	Mike Scott
Instrument Manufacturer, Model	Thermo Scientific, LTQ XL
Customizations	None

#### 1. General Features – (b) Control and Analysis Software

Software Name and Version	Xcalibur v2.0.7
Switching Criteria	30000 ions for full MS1
Isolation Width	2Da
Location of Parameters file	C:\Xcalibur\system\LTQ\msx\master.LTQ*

#### 2. Ion Source

Supply Type	Spray fed by liquid chromatography; flow at 250nL/min
Interface Manufacturer, Model, Catalog Number	Nanospray Source built by TSRI instrument shop
Sprayer type, coating, manufacturer, model, and catalog number	In-house 100um I.D. Column with 5um tip
Relevant voltages: tip, cone, acceleration	2.76kV spray voltage, 44V capillary voltage, -64V at gate lens
In-source dissociation?	Not Applicable (N/A)

#### 3. Post-source component

Final MS Stage Achieved	MS2
Gas Type and pressure for CID in	Helium gas, linear trap, 8.2 nBar
Collision Energy	35.0
Detector Type	Electron Multiplier
Detector Sensitivity	-1390V

#### 4. Spectrum and Peak List Generation and Annotation – (a) Spectrum Descriptions

Location and source of raw files including file name and type	C:\Xcalibur\data RAW files
MS level for spectra	MS2
Ion Mode	Positive
Precursor m/z and charge with the full mass spectrum for that peak	<i>in the raw data</i>

#### 4. Spectrum and Peak List Generation and Annotation – (b) Peak List Generation

Parameters triggering the generation of peak lists from raw data, where appropriate	N/A
Acquisition number from the RAW file	N/A
Smoothing	N/A
Background threshold	1000 counts
S/N estimation	N/A
% peak height for centroiding	N/A
Charge state calculation	calculated by SEQUEST in data processing

#### 4. Spectrum and Peak List Generation and Annotation – (c) Quantitation

Experimental Protocol	Label-free spectral counting using the software PatternLab for Proteomics. BMC Bioinformatics 2008, 9:316.
Number of combined samples and MS runs analyzed	6
Quantitation Approach	Spectral counting
Normalization Technique	Total Spectral Counts signal normalization
Location of quantitation data	TSRI MEM Torbett Lab

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