Accession	Name	Geomean	Stouffers P value
gi 40642994	Myosin essential light chain	1.3	0.0136
gi 405957584	Hypothetical protein CGI_10013948	1.3	0.0002
gi 405978198	Canopy-like protein 2	1.3	0.2984
gi 405976502	Uncharacterized protein c17orf98	1.3	0.0384
gi 405968273	Blastula protease 10	1.3	0.1437
gi 405962626	Serine-threonine kinase receptor-associated protein	1.3	0.0160
gi 405971647	Hypothetical protein CGI_10027007	1.3	0.0148
gi 405958005	Hypothetical protein CGI_10007019	1.3	
gi 405963883	Hypothetical protein CGI_10023805	1.3	0.0467
gi 405959794	Leucine-zipper-like transcriptional regulator 1	1.3	0.1776
gi 405972180	Tetratricopeptide repeat protein 25	1.3	0.0006
gi 405978588	Hypothetical protein CGI_10023566	1.3	0.0089
gi 405963560	Sarcoplasmic calcium-binding protein	1.3	0.0786
gi 405959327	Hypothetical protein CGI_10005346	1.3	0.0574
gi 405977511	Hypothetical protein CGI_10028134	1.3	0.2307
gi 405961165	26S proteasome non-ATPase regulatory subunit 11	1.3	
gi 405977606	Astacin	1.3	0.1926
gi 405955147	Plasminogen	1.3	0.0059
gi 405962751	RNA-binding protein 8A	1.3	0.1580
gi 405978866	Four and a half LIM domains protein 2	1.3	
gi 405977916	UPF0556 protein c19orf10-like protein	1.3	0.2907
gi 405950871	Hypothetical protein CGI_10010860	1.3	0.1289
gi 405960153	Neural/ectodermal development factor IMP-L2	1.3	0.0576
gi 405965635	Coiled-coil domain-containing protein 40	1.3	
gi 405970545	Hypothetical protein CGI_10022572	1.3	0.6108
gi 405970946	Fatty acid-binding protein, brain	1.3	0.2157
gi 405977975	Synapse-associated protein 1	1.3	0.0074
gi 353191	Myosin, regulatory la	1.3	0.3961
gi 405965909	Prohormone-4	1.3	0.0001
gi 405970837	Uncharacterized protein c7orf57-like protein	1.3	0.0444
gi 405957423	Transcription elongation regulator 1	1.3	0.0000
gi 405968105	Receptor-type tyrosine-protein phosphatase T	1.3	0.2063
gi 405955398	Proline-rich protein PRCC	1.3	
gi 405970386	Hypothetical protein CGI_10017976	1.3	0.0576
gi 405961983	Splicing factor 3B subunit 4	1.3	
gi 405970741	Translocon-associated protein subunit delta	1.3	
gi 405962196	ADAM family mig-17	1.3	0.2656
gi 22090632	Se-cadherin [Sexostrea echinata]	1.3	
gi 405970733	Lysosome-associated membrane glycoprotein 1	1.3	0.1723
gi 405971250	Glucosidase 2 subunit beta	1.3	0.0004
gi 405969368	Tetratricopeptide repeat protein 19	1.3	
gi 405975577	Fibrillin-1	1.3	0.0493
gi 405953094	Eukaryotic translation initiation factor 3 subunit J	1.3	0.0667
gi 405978332	Golgi membrane protein 1	1.3	
gi 405975553	Putative ferric-chelate reductase 1	1.3	0.3122
gi 405966927	Cytochrome P450 3A29	1.3	0.0290

Accession	Name	Geomean	Stouffers P value
gi 405966811	Intraflagellar transport protein 74-like protein	1.3	0.0057
gi 405969503	Hypothetical protein CGI_10015394	1.3	0.4036
gi 405960769	Lysyl oxidase-like protein 2	1.3	0.2132
gi 405978738	Hypothetical protein CGI_10022305	1.3	0.2138
gi 405977399	Metallothionein	1.3	0.1332
gi 405969789	Hypothetical protein CGI_10021722	1.3	
gi 405976897	LSM14-like protein A	1.3	0.0033
gi 405965308	Troponin T, skeletal muscle	1.3	0.0176
gi 405969739	Tankyrase-1	1.3	
gi 405953722	Calmodulin-like protein 12	1.3	0.0149
gi 405962468	Calmodulin-like protein 3	1.3	0.3286
gi 405958778	Hypothetical protein CGI_10021467	1.3	0.7819
gi 405966001	Non-specific lipid-transfer protein	1.3	0.0020
gi 405959472	AFG3-like protein 2	1.3	
gi 405968529	Hypothetical protein CGI_10017745	1.3	0.0053
gi 405978272	Calmodulin	1.3	0.0004
gi 405975550	Eukaryotic translation initiation factor 1A, X-chromosomal	1.3	0.3250
gi 405970197	Dysbindin	1.3	0.0997
gi 405973524	Thrombospondin-3b	1.3	
gi 405964593	Hypothetical protein CGI_10018739	1.4	0.0021
gi 405970776	Troponin C	1.4	0.0131
gi 405969229	NSFL1 cofactor p47	1.4	0.0041
gi 405968925	Hypothetical protein CGI_10004918	1.4	
gi 405959994	TP53-regulated inhibitor of apoptosis 1	1.4	0.0037
gi 405963180	Protein SET	1.4	0.0140
gi 405965328	Deoxyuridine 5'-triphosphate nucleotidohydrolase	1.4	0.1070
gi 405960921	Hypothetical protein CGI_10013223	1.4	
gi 405972179	Hypothetical protein CGI 10023112	1.4	
gi 405958084	Calmodulin	1.4	0.1367
gi 405954690	Hsc70-interacting protein	1.4	0.0000
gi 405972528	Hypothetical protein CGI_10020633	1.4	
gi 405971784	Hypothetical protein CGI_10018185	1.4	0.1715
gi 405955246	Hypothetical protein CGI_10002291	1.4	0.0901
gi 405959131	Uncharacterized protein c5orf48-like protein	1.4	0.0071
gi 405972975	Deleted in malignant brain tumors 1 protein	1.4	0.1003
gi 405952353	Calumenin-B	1.4	0.0001
gi 405977780	Calmodulin	1.4	
gi 405965382	Soma ferritin	1.4	0.0070
gi 405978177	Polyadenylate-binding protein 2	1.4	0.0147
gi 405970440	BAG family molecular chaperone regulator 1	1.4	
gi 405957935	Hypothetical protein CGI_10016657	1.4	
gi 405951401	Prefoldin subunit 1	1.4	
gi 405973521	Death-associated protein 1	1.4	0.0618
gi 405959947	Calmodulin	1.4	0.0571
gi 405952370	Hypothetical protein CGI_10006610	1.4	0.0000

Accession	Name	Geomean	Stouffers P value
gi 405969710	NADH dehydrogenase [ubiquinone] flavoprotein 3, mitochondrial	1.4	0.0745
gi 405960503	Peroxisomal biogenesis factor 19	1.4	0.0125
gi 405976121	WD repeat-containing protein c2orf44-like protein	1.4	0.1562
gi 405957059	Caprin-1	1.4	0.3680
gi 405973684	Calcium-regulated heat stable protein 1	1.4	0.5222
gi 405971204	E3 ubiquitin-protein ligase HECTD1	1.4	
gi 405978649	Laminin-like protein epi-1	1.4	0.0000
gi 405973965	Hypothetical protein CGI_10016182	1.4	0.1026
gi 148717307	Calreticulin	1.4	0.0000
gi 405965406	Aldehyde dehydrogenase	1.4	
gi 405959948	Hypothetical protein CGI_10013422	1.5	0.2817
gi 405957913	Calmodulin-like protein 12	1.5	0.0091
gi 405968717	Programmed cell death protein 6	1.5	0.0244
gi 405964391	Calmodulin	1.5	
gi 405967012	Hypothetical protein CGI_10026212	1.5	
gi 405978204	Tubulin-specific chaperone A	1.5	0.1713
gi 405952922	Calmodulin-5/6/7/8	1.5	0.0095
gi 405975575	Calmodulin	1.5	0.0057
gi 405952740	Latrophilin-3	1.5	0.3069
gi 405977513	High mobility group protein B3	1.5	0.0002
gi 405951638	Coiled-coil domain-containing protein 43	1.5	0.3364
gi 405968720	Hypothetical protein CGI_10023300	1.5	0.0001
gi 385843206	Ubiquitin/ribosomal protein S27 precursor	1.5	0.0090
gi 405963558	Sarcoplasmic calcium-binding protein	1.5	0.0000
gi 405975466	Hypothetical protein CGI 10010025	1.5	0.0011
gi 405977042	Elongation factor 1-beta	1.5	0.0000
gi 405976084	Tubulin polymerization-promoting protein family member 3	1.5	0.0007
gi 405973827	Cytochrome c	1.5	0.0001
gi 405951402	KH domain-containing, RNA-binding, signal transduction- associated protein 2	1.6	0.0001
gi 405974452	Adenylate kinase isoenzyme 6	1.6	
gi 405957087	Calumenin	1.6	0.0000
gi 405965135	Hypothetical protein CGI_10012662	1.6	0.0709
gi 405976946	Hypothetical protein CGI_10017526	1.6	0.0158
gi 405969780	Far upstream element-binding protein 3	1.6	0.0000
gi 405958088	Calmodulin	1.6	0.0006
gi 405955680	Nucleobindin-2	1.6	0.0156
gi 405972759	Hypothetical protein CGI 10018124	1.7	
gi 405975467	Hypothetical protein CGI_10010026	1.7	0.2697
gi 405960639	Deoxyuridine 5'-triphosphate nucleotidohydrolase	1.7	0.0428
gi 405946277	Hypothetical protein CGI_10000492	1.7	0.2095
gi 405970177	Eukaryotic translation initiation factor 4B	1.7	0.0000
gi 405968888	BET1-like protein	1.7	
gi 405975921	Clathrin light chain A	1.7	0.0020
gi 405958383	Hypothetical protein CGI 10007870	1.8	0.0035
gi 405969535	Hypothetical protein CGI_10016269	1.8	0.3226

Accession	Name	Geomean	Stouffers P value
gi 405957603	Dysferlin-interacting protein 1	1.8	
gi 405958091	Hypothetical protein CGI_10002445	1.9	
gi 405956668	LSM domain-containing protein 1	1.9	
gi 405978742	N-acetyltransferase 11	2.0	
gi 405969759	Vacuolar protein sorting-associated protein 13C	2.0	
gi 405978518	Plasminogen activator inhibitor 1 RNA-binding protein	2.2	0.0000
gi 405974686	Hypothetical protein CGI_10024481	6.2	0.3448
gi 405977783	Hypothetical protein CGI_10028016	0.4	
gi 405952065	Hypothetical protein CGI_10007261	0.5	
gi 405976102	Omega-crystallin	0.5	
gi 405960078	Hypothetical protein CGI_10010679	0.5	0.0092
gi 405960444	AP-3 complex subunit delta-1	0.5	
gi 405952183	Serine/threonine-protein kinase 3	0.6	
gi 405951710	Protein argonaute-2	0.6	
gi 405951433	Tubulin beta chain	0.6	
gi 405950782	Golgin subfamily A member 5	0.6	
gi 405963599	Endonuclease VIII-like 1	0.6	
gi 405969198	Tyrosine-protein kinase Yes, partial	0.6	
gi 405956722	Putative ATP-dependent RNA helicase DDX58	0.6	
gi 405954012	Thioredoxin	0.6	0.4928
gi 405957910	FMRFamide neuropeptide	0.6	0.3243
gi 405962092	Protein disabled	0.6	
gi 405969643	Hypothetical protein CGI_10012952	0.6	
gi 405956360	Multidrug resistance protein 1, partial	0.6	
gi 405957477	Putative thiopurine S-methyltransferase	0.6	
gi 405959119	60S ribosomal protein L3, partial	0.6	0.3909
gi 405969117	Dynein heavy chain 7, axonemal	0.6	0.1315
gi 405970293	Neuronal acetylcholine receptor subunit non-alpha-2	0.6	0.0649
gi 405973130	UDP-glucose:glycoprotein glucosyltransferase 1	0.6	
gi 405978806	Caspase-7	0.7	
gi 405971116	Hydroxyacylglutathione hydrolase, mitochondrial	0.7	
gi 405978371	Leucine-rich repeat-containing protein 51, partial	0.7	
gi 405966302	Heterogeneous nuclear ribonucleoprotein C	0.7	
gi 405965638	Tubulin alpha-1C chain	0.7	
gi 405975073	GMP synthase [glutamine-hydrolyzing]	0.7	
gi 405972926	Peroxiredoxin-4	0.7	
gi 405962986	Hypothetical protein CGI_10003439	0.7	
gi 405965431	Fatty acyl-CoA hydrolase precursor, medium chain	0.7	
gi 405964596	Hypothetical protein CGI_10018742	0.7	
gi 405972696	Hypothetical protein CGI_10011429	0.7	
gi 405970915	Serine/threonine-protein phosphatase 2A 55 kDa regulatory subunit B alpha isoform	0.7	
gi 2833326	Recname: Full=Actin, adductor muscle	0.7	0.2013
gi 405960911	Ninjurin-1	0.7	
gi 405961952	Importin subunit alpha-2	0.7	
gi 405965044	Neutral and basic amino acid transport protein RBAT	0.7	

Accession	Name	Geomean	Stouffers P value
gi 405971541	4-coumarateCoA ligase 1	0.7	
gi 405955166	Hypothetical protein CGI_10002360	0.7	
gi 405973425	Dynein intermediate chain 3, ciliary	0.7	
gi 405953996	Hypothetical protein CGI_10003795	0.7	0.1677
gi 405971953	Hypothetical protein CGI_10014226	0.7	
gi 405968884	Epithelial splicing regulatory protein 2	0.7	
gi 405964580	Actin, cytoplasmic	0.7	
gi 405966618	Carbonic anhydrase 2	0.7	0.0958
gi 405958743	Prostaglandin reductase 1	0.7	
gi 405951379	Amidase	0.7	
gi 405973433	Epoxide hydrolase 4	0.7	
gi 405975332	60S ribosomal protein L15	0.7	
gi 405950556	Copine-7	0.7	0.1801
gi 405966783	Myosin-11	0.7	
gi 405978774	Hypothetical protein CGI_10012934, partial	0.7	
gi 405975257	T-complex-associated testis-expressed protein 1	0.7	
gi 405972803	Radial spoke head protein 4-like protein A	0.7	0.0004
gi 405973865	Actin-1/3	0.7	0.0224
gi 405978703	Dynactin subunit 1	0.7	
gi 405963300	Serine/threonine-protein phosphatase 2A 56 kDa regulatory subunit delta isoform	0.7	
gi 405952433	Sodium- and chloride-dependent GABA transporter INE	0.7	0.1237
gi 405971621	Selenium-binding protein 1, partial	0.7	0.0166
gi 405974066	Eukaryotic translation initiation factor 4E-binding protein 1	0.7	0.1116
gi 405976463	Transketolase-like protein 2	0.7	0.0015
gi 405960423	Cadherin-23	0.7	
gi 405972486	Putative malate dehydrogenase 1B	0.7	
gi 405964311	Ankyrin repeat domain-containing protein 29, partial	0.7	
gi 405966284	Upstream activation factor subunit spp27	0.7	0.1182
gi 405969083	Hypothetical protein CGI_10018964	0.7	
gi 405963205	Heat shock 70 kDa protein 12A	0.8	
gi 405955982	Kyphoscoliosis peptidase	0.8	0.0000
gi 405964054	NADH-cytochrome b5 reductase 2	0.8	0.4340
gi 405963691	Cofilin	0.8	0.0263
gi 405971245	Eukaryotic translation initiation factor 3 subunit D	0.8	
gi 405965289	Adenine phosphoribosyltransferase	0.8	
gi 405975727	L-rhamnose-binding lectin CSL3, partial	0.8	0.3835
gi 405964567	Tubulin beta chain	0.8	0.0003
gi 405961313	Actin Cyl, cytoplasmic	0.8	0.1776
gi 405971057	Actin-related protein 2/3 complex subunit 3	0.8	0.0037
gi 405965394	Eukaryotic translation initiation factor 3 subunit B	0.8	
gi 405969278	Putative ATP-dependent RNA helicase an3	0.8	0.1980
gi 405969523	Catalase	0.8	0.1174
gi 405953522	Nucleolin	0.8	0.8253
gi 405964042	Coatomer subunit alpha	0.8	0.1012
gi 405963190	Alpha-aspartyl dipeptidase	0.8	

Accession	Name	Geomean	Stouffers P value
gi 405961262	Phosphoenolpyruvate carboxykinase, cytosolic [GTP]	0.8	0.1182
gi 405962165	Thioredoxin reductase 3	0.8	
gi 405972778	Protein phosphatase 1B	0.8	0.1687
gi 405951532	CAP-Gly domain-containing linker protein 1	0.8	
gi 405952292	Fibrillin-3	8.0	0.3843
gi 405968910	Gephyrin	0.8	
gi 405960133	Hypothetical protein CGI_10023385	0.8	
gi 405946057	Talin-2	0.8	
gi 405965007	DNA-directed RNA polymerase II subunit RPB1	0.8	
gi 405968439	Collagen alpha-1(XII) chain	0.8	0.4208
gi 405978645	Low-density lipoprotein receptor-related protein 6	0.8	
gi 405955753	Elongation factor 1-gamma	0.8	
gi 405967247	Sulfide:quinone oxidoreductase, mitochondrial	0.8	0.1003
gi 405952290	Armadillo repeat protein deleted in velo-cardio-facial syndrome-like protein	0.8	0.6182
gi 405975958	Heat shock 70 kDa protein 12A	0.8	0.1754
gi 405960652	60S ribosomal protein L17	0.8	
gi 405972862	Adenylate kinase isoenzyme 5	8.0	0.0441
gi 405972436	Cystatin-A2	0.8	0.7273
gi 405975234	Dual oxidase 2, partial	0.8	
gi 405959180	Protein SET	0.8	0.0132
gi 405963822	UPF0663 transmembrane protein c17orf28	0.8	0.0747
gi 405963573	Hypothetical protein CGI_10024254	0.8	
gi 405960428	PDZ and LIM domain protein 1 Epidermal growth factor receptor kinase substrate 8-like	0.8	0.0762
gi 405968887	protein 1		0.0702
gi 405971344	Mitochondrial carrier-like protein 2	8.0	
gi 405973648	Pericentriolar material 1 protein	8.0	0.3047
gi 405961193	Coatomer subunit beta', partial	8.0	
gi 405953995	Hypothetical protein CGI_10003794	8.0	0.0653
gi 405967835	Methylglutaconyl-CoA hydratase, mitochondrial	0.8	
gi 405953465	Tether containing UBX domain for GLUT4	0.8	
gi 405978602	Calcium-binding protein NCS-1	0.8	
gi 405971875	Hemicentin-1	0.8	
gi 405961385	Glutamine synthetase	0.8	0.0572
gi 405951942	Tubulin alpha-1C chain	0.8	0.5045
gi 405965290	Hypothetical protein CGI_10002265	0.8	
gi 405973146	Annexin A7	0.8	0.0002
gi 157154403	Sodium/potassium-transporting ATPase alpha chain	0.8	
gi 405953624	Hypothetical protein CGI_10004246	0.8	0.0=04
gi 405966819	Serine hydroxymethyltransferase, cytosolic	0.8	0.3721
gi 405973340	Actin	0.8	0.0208
gi 405962300	Frizzled-5	0.8	0.000
gi 405978690	Propionyl-CoA carboxylase beta chain, mitochondrial	0.8	0.0778
gi 405963030	Hypothetical protein CGI_10025157	0.8	0.0588
gi 405956092	DNA-directed RNA polymerases I, II, and III subunit RPABC2	0.8	0.2201
gi 405966314	Lysyl-tRNA synthetase	8.0	

Accession	Name	Geomean	Stouffers P value
gi 405965049	Nucleolar protein 58	0.8	0.4175
gi 405951327	Protein G7c	0.8	0.6167
gi 47076115	Mu class glutathione S-transferase	0.8	
gi 405977043	NADH-ubiquinone oxidoreductase 75 kDa subunit, mitochondrial	0.8	
gi 405952907	Hypothetical protein CGI_10005641	0.8	
gi 405959253	Myosin-VI	0.8	0.4115
gi 405974168	Titin	0.8	0.0203
gi 405954612	Fucolectin-5	0.8	
gi 405960204	Hypoxia up-regulated protein 1	0.8	
gi 405970532	Septin-2	0.8	
gi 405950592	Phosphoglucomutase-1	8.0	0.0982
gi 405960597	Phytanoyl-CoA dioxygenase domain-containing protein 1-like protein	0.8	0.3204
gi 405969341	Glycyl-tRNA synthetase	0.8	
gi 405975071	Coronin-1B	0.8	0.0008
gi 405960605	Uncharacterized protein c20orf26	0.8	
gi 405972553	Dual oxidase maturation factor 1	0.8	
gi 40643028	Ribosomal protein L19	0.8	0.2355
gi 405960660	Protein disulfide-isomerase A5	0.8	
gi 405967381	Cysteinyl-tRNA synthetase, cytoplasmic	0.8	
gi 405970695	Receptor-type tyrosine-protein phosphatase beta	0.8	
gi 405974068	Glucosaminefructose-6-phosphate aminotransferase [isomerizing] 1	0.8	0.0033
gi 405951651	2-amino-3-carboxymuconate-6-semialdehyde decarboxylase	0.8	
gi 405959219	Mitogen-activated protein kinase kinase kinase kinase 3, partial	0.8	
gi 405964691	Acetyl-coenzyme A synthetase	0.8	0.2508
gi 405959335	Serine/threonine-protein kinase Nek4	0.8	0.0682
gi 405971561	Zonadhesin	0.8	0.0077
gi 405966858	3-hydroxyacyl-CoA dehydrogenase type-2	0.8	
gi 405966303	Hypothetical protein CGI_10009847	0.8	0.1231
gi 405964597	Hypothetical protein CGI_10018743	0.8	
gi 405968446	Hypothetical protein CGI_10022012	8.0	
gi 405970058	Eukaryotic translation initiation factor 3 subunit A	0.8	0.0806
gi 405955211	Hypothetical protein CGI_10002332	0.8	0.4132
gi 405954502	Splicing factor, arginine/serine-rich 6	0.8	
gi 405960477	Laminin subunit beta-2	0.8	
gi 405966886	Hypothetical protein CGI_10025210	0.8	
gi 405978465	Delta-1-pyrroline-5-carboxylate dehydrogenase, mitochondrial	0.8	
gi 405956520	Putative aminopeptidase W07G4.4	0.8	0.0552
gi 405950078	Nucleoprotein TPR	0.8	
gi 405975672	Long-chain-fatty-acidCoA ligase 4	0.8	0.0875
gi 405976274	Hypothetical protein CGI_10021278	0.8	
gi 405953328	Protein kinase C delta type	0.8	
gi 405970941	Hypothetical protein CGI_10011199	0.8	
gi 405978868	Uncharacterized protein c1orf93-like protein	0.8	
gi 405976016	Villin-1	0.8	0.0779

Accession	Name	Geomean	Stouffers P value
gi 405953593	Double-stranded RNA-specific editase 1	0.8	
gi 405972492	Kinesin-related protein 1	0.8	
gi 405974500	Eukaryotic peptide chain release factor GTP-binding subunit ERF3B	0.8	0.2727
gi 405952027	Lethal(2) giant larvae-like protein 1	0.8	0.2618
gi 405967527	Nesprin-1	0.8	0.0121
gi 405975484	Protein lethal(2)essential for life	0.8	0.0154
gi 405951040	Eukaryotic translation initiation factor 3 subunit C	0.8	0.7591
gi 405960037	Serine/threonine-protein phosphatase 6 regulatory subunit 3	0.8	0.1371
gi 405964650	Extended synaptotagmin-2	0.8	0.2952
gi 405977935	Hypothetical protein CGI_10018270	0.8	0.3434
gi 405976336	Hypothetical protein CGI_10026535	0.8	0.5951
gi 71040623	Ribosomal protein S27	0.8	
gi 405966565	Putative glycogen [starch] synthase	0.8	
gi 405975901	Uncharacterized protein c3orf32	0.8	
gi 405973981	Calcium-dependent protein kinase C	0.8	
gi 405978279	AP-2 complex subunit sigma	0.8	
gi 405975834	Isocitrate dehydrogenase [NADP], mitochondrial	0.8	0.0000
gi 405950813	GPI inositol-deacylase	0.8	
gi 405953127	Allene oxide synthase-lipoxygenase protein, partial	0.8	0.0001
gi 405952800	Hypothetical protein CGI_10005804	0.8	
gi 405966125	Glutathione S-transferase 3	0.8	0.1176
gi 405962701	Malectin	0.8	
gi 405970446	Sodium- and chloride-dependent glycine transporter 2	0.8	
gi 405975785	Filamin-A	0.8	0.0415
gi 405969429	Plasma alpha-L-fucosidase	0.8	0.1322
gi 405964482	Hypothetical protein CGI_10003076	0.8	0.3253
gi 405966210	Collectin-12	0.8	0.5845
gi 405964165	Calcium/calmodulin-dependent protein kinase type II delta chain	0.8	0.2548
gi 405966818	Putative ATP-dependent RNA helicase DDX5	0.8	
gi 405971733	Formimidoyltransferase-cyclodeaminase	0.8	0.5047
gi 405945552	2,4-dienoyl-CoA reductase, mitochondrial	0.8	0.5527
gi 405955564	Cytoplasmic aconitate hydratase	0.8	0.2590
gi 405977889	Putative adenosylhomocysteinase 3	0.8	
gi 405978819	Leucine-rich repeat-containing protein 59	0.8	0.5226
gi 405959699	Aldo-keto reductase family 1 member B10	0.8	
gi 405965483	Dihydrolipoyl dehydrogenase, mitochondrial	0.8	0.3794
gi 405971538	Synaptophysin	0.8	0.3616
gi 405972424	Camp-dependent protein kinase type II regulatory subunit	0.8	0.3822
gi 405951265	Hypothetical protein CGI_10009262	0.8	
gi 405977348	Histone H5	0.8	0.3119
gi 405957950	SH3 domain-binding protein 5-like protein	0.8	
gi 405969389	DNA-directed RNA polymerases I, II, and III subunit RPABC3	0.8	
gi 405969881	Synaptobrevin	0.8	
gi 405974166	Titin	0.8	
gi 405953053	Echinoderm microtubule-associated protein-like 1	0.8	0.4557

Accession	Name	Geomean	Stouffers P value
gi 405960113	Ubiquitin carboxyl-terminal hydrolase 15	0.8	
gi 405978625	Metaxin-2	8.0	
gi 405961544	Plexin-A4	0.8	
gi 405978696	Alpha-aminoadipic semialdehyde dehydrogenase	0.8	0.0071
gi 405973025	Coatomer subunit delta	8.0	0.4564
gi 405964355	Calpain-A	0.8	0.0263
gi 405970056	Kyphoscoliosis peptidase	0.8	0.0112
gi 22758874	Ribosomal protein L	0.8	
gi 405970435	Thioredoxin domain-containing protein 5	0.8	0.6454
gi 405971145	Myosin heavy chain, non-muscle, partial	0.8	0.0000
gi 405975057	Myosin heavy chain, striated muscle	0.8	0.0024
gi 405960658	Plastin-1	0.8	0.6109
gi 405966381	Dynein heavy chain 6, axonemal	0.8	0.4499
gi 405977321	Hypothetical protein CGI_10021646	0.8	
gi 405965776	LISH domain-containing protein c16orf63-like protein	0.8	
gi 405956597	Hypothetical protein CGI_10000542	0.8	0.3625
gi 405970128	Nucleolar protein 56	0.8	
gi 405964721	EF-hand domain-containing protein 1	0.8	0.1079
gi 405975707	Hypothetical protein CGI_10015240	0.8	
gi 405974485	13 kDa ribonucleoprotein-associated protein	0.8	
gi 405977320	Neural cell adhesion molecule 1	0.8	0.0000
gi 157154401	IK cytokine	0.8	
gi 405972660	Nucleosome assembly protein 1-like 4	0.8	
gi 405960169	Cytoplasmic dynein 1 intermediate chain 2	0.8	
gi 405973904	Spermatogenesis-associated protein 6	0.8	
gi 405958791	Arginine kinase	0.8	
gi 405958081	Calmodulin	0.8	
gi 405960314	Hypothetical protein CGI_10005509	0.9	0.8697
gi 405969732	Laminin subunit alpha	0.9	0.0188
gi 405951570	15-hydroxyprostaglandin dehydrogenase [NAD+]	0.9	
gi 405960674	Hypothetical protein CGI_10010476	0.9	0.1742
gi 405967698	Protein disulfide-isomerase A3	0.9	0.0000
gi 405963261	Guanine nucleotide-binding protein subunit beta	0.9	0.4306
gi 405968440	Collagen alpha-6(VI) chain	0.9	0.0278
gi 405963184	Hypothetical protein CGI_10018076	0.9	0.7710
gi 405957246	Excitatory amino acid transporter 2	0.9	
gi 405976919	MORN repeat-containing protein 3	0.9	
gi 405957573	Hypothetical protein CGI_10013935	0.9	0.3525
gi 405966072	Putative adenylate kinase 7	0.9	0.0940
gi 405964761	Aconitate hydratase, mitochondrial	0.9	0.0029
gi 405961151	Xaa-Pro aminopeptidase 1	0.9	0.9001
gi 405972826	Collagen alpha-5(VI) chain	0.9	
gi 405957027	Heterogeneous nuclear ribonucleoprotein K	0.9	0.6010
gi 405973442	Nicotinamide riboside kinase 1	0.9	
gi 405965813	Williams-Beuren syndrome chromosomal region 27 protein	0.9	
gi 405961227	Serine/threonine-protein kinase phg2	0.9	0.4331