

Table S1, Roth et al. 2002

Men Before-ST	Women Before-ST	Before-ST Ratio	Acc#	Gene	Men After-ST	Women After-ST	After-ST Ratio
0.907	0.056	0.062	AA456636	RAN, member RAS oncogene family	1.025	0.492	0.480
0.832	0.055	0.066	H54289	Golgi vesicular membrane trafficking protein p18	0.849	0.480	0.566
3.463	0.259	0.075	N32201	osteomodulin	2.864	1.289	0.450
0.447	0.041	0.091	H70775	ESTs	0.455	0.246	0.541
1.235	0.127	0.103	R38619	fucose-1-phosphate guanylyltransferase	1.517	0.683	0.450
0.560	0.060	0.107	AA447793	H.sapiens mRNA for pur alpha extended 3'untranslated region	0.511	0.294	0.575
1.082	0.118	0.109	R92994	matrix metalloproteinase 12 (macrophage elastase)	1.120	0.656	0.585
1.881	0.206	0.109	AA598561	CD164 antigen, sialomucin	1.885	0.894	0.474
0.426	0.048	0.113	N53031	UDP glycosyltransferase 2 family, polypeptide B4	0.411	0.240	0.583
0.389	0.044	0.114	AA418524	phospholipase D2	0.327	0.180	0.550
0.691	0.083	0.121	W72437	general transcription factor IIIH, polypeptide 2 (44kD subunit)	0.649	0.373	0.575
0.574	0.070	0.121	H95088	poly (ADP-ribose) glycohydrolase	0.555	0.298	0.538
1.685	0.226	0.134	AA171449	biphenylhydrolase-like (serine hydrolase; breast epithelial mucin-associated antigen)	1.772	0.900	0.508
0.449	0.061	0.135	N81076	far upstream element (FUSE) binding protein 3	0.417	0.223	0.534
0.992	0.135	0.136	AA496149	3-hydroxy-3-methylglutaryl-Coenzyme A synthase 2 (mitochondrial)	1.051	0.569	0.541
0.691	0.096	0.140	AA416783	H-2K binding factor-2	0.718	0.322	0.448
0.571	0.080	0.140	H97488	N-ethylmaleimide-sensitive factor	0.538	0.225	0.418
1.116	0.158	0.141	H79779	histone deacetylase 3	1.054	0.455	0.432
1.008	0.145	0.144	H84871	Ste-20 related kinase	1.050	0.474	0.452
0.572	0.083	0.144	AA056013	EST	0.583	0.285	0.489
0.548	0.080	0.145	AA057233	S-antigen; retina and pineal gland (arrestin)	0.581	0.308	0.530
1.129	0.166	0.147	N48652	tumor protein p53-binding protein, 2	1.063	0.601	0.565
0.340	0.050	0.148	AA400474	ESTs, Highly similar to zona-pellucida-binding protein [H.sapiens]	0.466	0.200	0.428
1.005	0.151	0.150	R22179	PAX transcription activation domain interacting protein 1 like	1.053	0.591	0.561
1.723	0.265	0.154	W46972	solute carrier family 20 (phosphate transporter), member 1	1.934	0.915	0.473
1.245	0.192	0.155	AA063580	TATA box binding protein (TBP)-associated factor, RNA polymerase II, F, 55kD	1.227	0.661	0.539
0.490	0.076	0.155	AA453404	PPAR binding protein	0.457	0.250	0.547

Table S1, Roth et al. 2002

1.090	0.172	0.158	AA410207	kinesin 2 (60-70kD)	0.990	0.546	0.551
0.651	0.104	0.160	R52639	serine hydroxymethyltransferase 1 (soluble)	0.676	0.358	0.529
1.350	0.219	0.162	AA481562	aspartyl-tRNA synthetase	1.330	0.656	0.494
0.547	0.089	0.163	R62603	collagen, type VI, alpha 3	0.529	0.281	0.533
0.438	0.074	0.168	T70057	immunoglobulin J polypeptide, linker protein for immunoglobulin alpha and mu polypeptides	0.400	0.200	0.502
0.469	0.079	0.169	R51021	cytochrome P450, subfamily XXVIA, polypeptide 1	0.513	0.249	0.486
0.567	0.096	0.170	AA644092	non-metastatic cells 1, protein (NM23A) expressed in	0.676	0.288	0.426
0.678	0.118	0.173	AA448664	polymerase (DNA directed), epsilon 2	0.611	0.298	0.488
1.067	0.185	0.174	AA400234	nuclear receptor co-repressor 2	1.137	0.592	0.521
1.253	0.220	0.175	H58873	solute carrier family 2 (facilitated glucose transporter), member 1	1.183	0.610	0.516
0.580	0.107	0.184	AA447748	dihydrolipoamide dehydrogenase (E3 component of pyruvate dehydrogenase complex, 2-oxo-glutarate complex, branched chain keto acid dehydrogenase complex)	0.520	0.302	0.580
0.506	0.094	0.186	AA421269	EST, Highly similar to phosphatidylinositol 4-kinase 230 [H.sapiens]	0.522	0.254	0.487
0.383	0.073	0.190	AA127093	histone deacetylase 2	0.341	0.157	0.460
2.247	0.440	0.196	AA449361	ring finger protein 13	2.039	1.096	0.538
1.056	0.208	0.197	AA490996	interferon, gamma-inducible protein 16	1.198	0.673	0.562
1.054	0.208	0.197	AA610004	Homo sapiens putative oncogene protein mRNA, partial cds	1.015	0.574	0.566
0.635	0.126	0.199	AA699317	ring-box 1	0.709	0.336	0.474
0.476	0.095	0.200	AA504113	M-phase phosphoprotein 10 (U3 small nucleolar ribonucleoprotein)	0.459	0.269	0.585
0.315	0.063	0.201	AA432261	testis-specific ankyrin motif containing protein	0.357	0.173	0.483
1.011	0.207	0.204	W31899	zinc finger protein 165	0.968	0.493	0.509
1.021	0.211	0.206	N29901	dihydrolipoamide S-acetyltransferase (E2 component of pyruvate dehydrogenase complex)	1.105	0.626	0.566
1.071	0.225	0.211	R63702	cyclin T2	0.927	0.430	0.464

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0.600	0.126	0.211	H23075	hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), beta subunit	0.593	0.327	0.551
0.291	0.061	0.211	AA463411	chondroitin sulfate proteoglycan 6 (bamacan)	0.298	0.164	0.552
0.616	0.130	0.211	H87471	kynureninase (L-kynurenine hydrolase)	0.745	0.379	0.508
0.317	0.067	0.212	AA429895	ATP-binding cassette, sub-family C (CFTR/MRP), member 3	0.330	0.161	0.486
0.254	0.054	0.213	AA701929	bystin-like	0.274	0.161	0.588
0.373	0.080	0.214	H15677	protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), beta isoform	0.384	0.211	0.549
0.460	0.099	0.216	H05935	cytochrome P450, subfamily XXVIIA (steroid 27-hydroxylase, cerebrotendinous xanthomatosis), polypeptide 1	0.659	0.281	0.427
0.719	0.156	0.217	H68922	integrin, alpha 1	0.738	0.382	0.518
0.330	0.072	0.219	AA490159	glucose-6-phosphatase, transport (glucose-6-phosphate) protein 1	0.294	0.137	0.466
0.783	0.171	0.219	AA405989	death-associated protein 6	0.825	0.473	0.573
0.495	0.108	0.219	N31933	tyrosinase (oculocutaneous albinism IA)	0.473	0.252	0.533
1.197	0.263	0.220	W45688	caspase 6, apoptosis-related cysteine protease	1.361	0.690	0.507
0.396	0.087	0.220	R54807	sarcoglycan, beta (43kD dystrophin-associated glycoprotein)	0.467	0.265	0.568
0.345	0.076	0.221	AA122287	glycoprotein A repetitions predominant	0.334	0.186	0.556
1.305	0.296	0.227	R72097	pepsinogen 5, group I (pepsinogen A)	1.306	0.685	0.524
0.430	0.099	0.231	AA447986	heat shock 40kD protein 2	0.438	0.209	0.477
3.138	0.725	0.231	W86860	nuclear VCP-like	2.633	1.521	0.578
1.185	0.277	0.234	N36882	v-yes-1 Yamaguchi sarcoma viral oncogene homolog 1	1.312	0.727	0.554
0.310	0.073	0.236	N52533	EST	0.279	0.155	0.557
0.801	0.190	0.237	AA055163	calsequestrin 2, cardiac muscle zinc finger protein 184 (Kruppel-like)	0.983	0.478	0.487
0.522	0.124	0.238	AA455657	zinc finger protein 184 (Kruppel-like)	0.515	0.242	0.469
2.622	0.628	0.239	R48320	EphB1	3.200	1.554	0.486
0.382	0.092	0.242	R40059	nucleoporin 153kD	0.364	0.199	0.546
1.457	0.353	0.242	R54818	eukaryotic translation initiation factor 2B, subunit 5 (epsilon, 82kD)	1.688	0.966	0.573

Table S1, Roth et al. 2002

0.450	0.110	0.243	AA195959	cysteine and glycine-rich protein 3 (cardiac LIM protein)	0.396	0.184	0.464
1.002	0.244	0.244	AA150918	hydroxysteroid (11-beta) dehydrogenase 1	1.038	0.604	0.581
0.421	0.104	0.246	AA460265	ets variant gene 5 (ets-related molecule)	0.468	0.275	0.587
0.390	0.097	0.248	AA136710	glyoxalase I	0.367	0.209	0.571
0.421	0.104	0.248	AA504844	DKFZp434J1813 protein	0.409	0.234	0.573
2.344	0.582	0.248	N25204	leukemia associated gene 2	2.353	1.091	0.464
0.972	0.245	0.252	AA451792	protein phosphatase 3 (formerly 2B), catalytic subunit, alpha isoform (calcineurin A alpha)	1.149	0.611	0.531
0.775	0.196	0.252	R38433	phosphofructokinase, platelet	0.847	0.473	0.559
0.198	0.050	0.253	AA465386	ESTs	0.165	0.089	0.543
1.080	0.277	0.256	H49511	neurogranin (protein kinase C substrate, RC3)	1.119	0.650	0.581
0.276	0.071	0.257	T64878	peanut (Drosophila)-like 2	0.359	0.146	0.406
0.589	0.152	0.258	H08808	ESTs	0.602	0.350	0.582
0.513	0.135	0.263	R76437	thromboxane A synthase 1 (platelet, cytochrome P450, subfamily V)	0.494	0.281	0.568
0.857	0.226	0.264	AA453477	X-prolyl aminopeptidase (aminopeptidase P)-like	1.092	0.635	0.582
0.349	0.093	0.267	N50636	Homo sapiens guanine nucleotide exchange factor smgGDS (RAP1GDS1) mRNA, alternatively spliced, complete cds	0.363	0.155	0.426
1.947	0.526	0.270	H15431	polymerase (RNA) II (DNA directed) polypeptide D	2.160	1.094	0.507
2.419	0.660	0.273	H78385	huntingtin-interacting protein 2	2.235	0.987	0.442
0.516	0.142	0.274	AA679208	transforming growth factor beta-activated kinase-binding protein 1	0.501	0.246	0.492
1.362	0.376	0.276	AA431439	guanylate cyclase activator 1A (retina)	1.311	0.575	0.439
0.944	0.261	0.276	AA136533	transcription elongation factor B (SIII), polypeptide 1-like	0.947	0.517	0.546
1.014	0.281	0.277	W45148	acid phosphatase 1, soluble	1.026	0.582	0.567
2.348	0.654	0.279	AA496800	Human transposon-like element mRNA	2.217	1.294	0.583
0.274	0.076	0.279	AA401428	nucleoporin 214kD (CAIN)	0.344	0.202	0.588
0.627	0.176	0.280	AA460688	neuropeptide FF-amide peptide precursor	0.712	0.367	0.516
0.628	0.178	0.283	AA481555	thyroid hormone receptor coactivating protein	0.627	0.259	0.414
2.041	0.579	0.284	AA046430	lung type-I cell membrane-associated glycoprotein	2.720	1.426	0.524
0.438	0.125	0.286	AA490046	fibroblast growth factor (acidic) intracellular binding protein	0.516	0.303	0.587

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0.601	0.173	0.288	AA181300	proteasome (prosome, macropain) subunit, beta type, 8 (large multifunctional protease 7)	0.707	0.356	0.504
0.492	0.142	0.289	AA449459	sulfotransferase, estrogen-preferring	0.621	0.320	0.515
0.923	0.267	0.290	AA430035	reticulon 3	0.637	0.351	0.551
1.290	0.374	0.290	N50745	cortistatin	1.817	0.970	0.534
2.550	0.741	0.291	N94713	intersectin 2	2.759	1.416	0.513
0.349	0.102	0.292	AA001749	microtubule-associated protein, RP/EB family, member 1	0.374	0.208	0.556
0.749	0.223	0.297	AA680129	polymerase (DNA directed), epsilon	0.759	0.434	0.572
1.301	0.393	0.302	H05899	heterogeneous nuclear ribonucleoprotein C (C1/C2)	1.286	0.701	0.545
0.264	0.080	0.303	H13688	SMC (mouse) homolog, X chromosome	0.364	0.201	0.553
0.308	0.094	0.305	H23075	hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), beta subunit	0.347	0.171	0.492
0.635	0.195	0.306	AA419251	interferon induced transmembrane protein 1 (9-27)	0.579	0.297	0.512
0.886	0.275	0.310	AA401883	sialidase 1 (lysosomal sialidase)	0.723	0.317	0.439
0.357	0.111	0.311	T73558	deoxyribonuclease I-like 3	0.388	0.210	0.541
1.783	0.566	0.317	AA460251	NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 1 (6kD, KFYI)	1.465	0.794	0.542
0.843	0.270	0.320	AA463204	pleomorphic adenoma gene-like 1	0.758	0.412	0.544
0.307	0.101	0.328	AA520979	transmembrane 7 superfamily member 1 (upregulated in kidney)	0.297	0.150	0.506
0.400	0.131	0.328	AA425687	DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 1	0.375	0.205	0.547
0.145	0.048	0.329	H66484	nuclear body protein Sp140	0.135	0.058	0.431
0.767	0.256	0.334	AA403126	proteasome (prosome, macropain) 26S subunit, non-ATPase, 9	0.986	0.457	0.464
0.616	0.207	0.335	AA429882	similar to yeast BET3 (S. cerevisiae)	0.699	0.309	0.442
0.775	0.263	0.340	AA147594	tumor necrosis factor receptor superfamily, member 8	0.791	0.394	0.498
0.343	0.118	0.344	H26182	neuronal PAS domain protein 2	0.299	0.124	0.413
2.070	0.711	0.344	AA459401	kallikrein 10	2.268	1.272	0.561
0.186	0.064	0.344	AA708816	clathrin, heavy polypeptide-like 1	0.221	0.129	0.586
1.799	0.620	0.345	AA293192	putative nucleic acid binding protein RY-1	2.075	1.076	0.519

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0.651	0.225	0.345	AA521025	G protein pathway suppressor 1	0.564	0.284	0.504
0.246	0.085	0.345	N63988	ESTs	0.221	0.125	0.566
1.876	0.653	0.348	N94385	cartilage oligomeric matrix protein (pseudoachondroplasia, epiphyseal dysplasia 1, multiple)	2.123	1.232	0.580
1.751	0.616	0.352	N99154	protein kinase, Y-linked	1.347	0.722	0.536
0.861	0.307	0.357	AA429310	chromosome 2 open reading frame 3	0.927	0.502	0.541
0.262	0.094	0.359	AA504266	stromal antigen 2	0.279	0.164	0.586
1.644	0.593	0.360	AA088258	Human protein immuno-reactive with anti-PTH polyclonal antibodies mRNA, partial cds	1.899	1.115	0.587
1.004	0.365	0.364	AA630734	seryl-tRNA synthetase	1.083	0.635	0.586
1.725	0.628	0.364	N62245	CDC7 (cell division cycle 7, S. cerevisiae, homolog)-like 1	1.689	0.791	0.468
0.835	0.305	0.365	AA406040	maternal G10 transcript	0.885	0.465	0.526
0.412	0.150	0.365	AA427954	hook2 protein	0.478	0.242	0.507
0.474	0.175	0.369	AA133212	nuclear receptor coactivator 4	0.354	0.187	0.528
1.072	0.397	0.370	AA459909	6-pyruvoyl-tetrahydropterin synthase/dimerization cofactor of hepatocyte nuclear factor 1 alpha (TCF1)	1.160	0.666	0.574
3.442	1.277	0.371	AA150301	adrenal gland protein AD-004	3.038	1.700	0.560
0.250	0.093	0.373	H28952	ADP-ribosylation factor 4-like	0.288	0.163	0.567
0.870	0.325	0.373	AA454113	meprin A, alpha (PABA peptide hydrolase)	0.843	0.465	0.551
0.512	0.194	0.378	AA424344	uroporphyrinogen decarboxylase	0.579	0.249	0.430
1.051	0.399	0.379	AA469965	lymphocyte-specific protein tyrosine kinase	1.122	0.576	0.514
3.200	1.234	0.386	AA418564	ESTs, Moderately similar to I59372 cadherin 12 - human [H.sapiens]	3.369	1.639	0.487
0.538	0.210	0.390	AA427891	activated p21cdc42Hs kinase	0.563	0.321	0.570
0.299	0.118	0.394	AA436871	syntaxin 3A	0.293	0.171	0.584
2.691	1.060	0.394	AA136566	forkhead box M1	2.503	1.236	0.494
0.187	0.074	0.397	AA291484	cytochrome P450, subfamily IVB, polypeptide 1	0.207	0.107	0.515
0.237	0.096	0.406	AA448157	cytochrome P450, subfamily I (dioxin-inducible), polypeptide 1 (glaucoma 3, primary infantile)	0.260	0.147	0.566
3.422	1.390	0.406	AA450062	prostate differentiation factor	4.078	1.980	0.485
0.924	0.376	0.406	H08808	ESTs	0.972	0.521	0.535
0.242	0.099	0.409	N92901	fatty acid binding protein 4, adipocyte	0.527	0.216	0.409
0.301	0.123	0.411	AA625662	histone acetyltransferase 1	0.336	0.193	0.576
0.267	0.113	0.422	AA490044	solute carrier family 5 (inositol transporters), member 3	0.323	0.177	0.549
1.287	0.546	0.424	AA428196	POU domain, class 4, transcription factor 1	1.457	0.710	0.488

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0.657	0.279	0.425	AA701545	ribonuclease, RNase A family, k6	0.729	0.300	0.411
0.296	0.126	0.426	AA626867	KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 2	0.280	0.140	0.499
2.970	1.283	0.432	AA262504	eyes absent (Drosophila) homolog 3	2.998	1.739	0.580
0.915	0.402	0.439	AA446839	BCL2/adenovirus E1B 19kD-interacting protein 3	0.770	0.438	0.568
0.681	0.299	0.439	AA430178	RAN binding protein 2-like 1	0.630	0.313	0.497
0.415	0.183	0.440	AA446928	v-erb-b2 avian erythroblastic leukemia viral oncogene homolog 2 (neuro/glioblastoma derived oncogene homolog)	0.495	0.276	0.557
0.576	0.258	0.447	AA148536	nucleoporin 98kD	0.604	0.283	0.468
0.766	0.347	0.453	AA460728	voltage-dependent anion channel 3	0.660	0.384	0.581
1.532	0.703	0.459	AA447632	glycoprotein M6A	1.994	1.170	0.587
0.592	0.273	0.462	AA670380	diphtheria toxin resistance protein required for diphthamide biosynthesis (Saccharomyces)-like 1	0.505	0.263	0.521
0.572	0.266	0.464	W88884	TAR DNA binding protein	0.558	0.319	0.572
0.108	0.051	0.470	AA682386	oxidised low density lipoprotein (lectin-like) receptor 1	0.154	0.072	0.468
0.362	0.171	0.474	AA630017	transcription elongation factor B (SIII), polypeptide 2 (18kD, elongin B)	0.369	0.212	0.574
0.154	0.073	0.478	AA676802	sialic acid binding Ig-like lectin 5	0.137	0.079	0.577
0.343	0.166	0.485	AA476490	thyroid hormone receptor interactor 4	0.327	0.180	0.551
0.670	0.326	0.486	AA490538	zinc finger protein homologous to Zfp161 in mouse	0.616	0.341	0.554
0.461	0.227	0.493	AA476438	similar to rat HREV107	0.532	0.289	0.544
0.162	0.080	0.494	AA046067	ESTs	0.147	0.086	0.581
1.617	0.805	0.498	AA461506	testis-specific kinase 1	1.908	0.864	0.453
0.141	0.071	0.500	AA702663	myosin IXB	0.147	0.083	0.562
0.321	0.656	2.042	AA156793	similar to glucosamine-6-sulfatases	0.194	0.417	2.149
0.167	0.341	2.046	N64862	FYN-binding protein (FYB-120/130)	0.117	0.277	2.374
0.183	0.378	2.060	H12320	cAMP responsive element binding protein 1	0.136	0.260	1.913
0.575	1.190	2.070	AA191488	solute carrier family 31 (copper transporters), member 1	0.374	0.820	2.194
0.229	0.476	2.075	AA036881	chemokine (C-C motif) receptor 1	0.113	0.339	3.000
0.391	0.818	2.090	R45413	sarcoma amplified sequence	0.206	0.526	2.557
0.601	1.260	2.095	AA428778	ephrin-B1	0.399	0.937	2.344
0.577	1.216	2.107	H65066	visinin-like 1	0.411	0.810	1.972

Table S1, Roth et al. 2002

0.454	0.964	2.125	H99588	lymphoid nuclear protein related to AF4	0.286	0.716	2.502
0.244	0.518	2.128	R60019	membrane protein, palmitoylated 2 (MAGUK p55 subfamily member 2)	0.169	0.353	2.091
7.779	16.951	2.179	AA455272	ITBA1 gene	5.361	9.517	1.775
0.308	0.684	2.220	AA679177	butyrate-induced transcript 1	0.246	0.438	1.778
0.169	0.376	2.226	R72079	CD79B antigen (immunoglobulin-associated beta)	0.125	0.311	2.485
0.374	0.840	2.247	AA448400	plectin 1, intermediate filament binding protein, 500kD	0.298	0.561	1.887
0.185	0.419	2.261	AA011347	IGF-II mRNA-binding protein 3	0.151	0.286	1.893
0.326	0.749	2.294	H46425	purine-rich element binding protein A	0.254	0.470	1.850
0.200	0.470	2.346	R52085	growth differentiation factor 10	0.240	0.581	2.422
0.161	0.378	2.349	R51209	protein phosphatase 2A, regulatory subunit B' (PR 53)	0.249	0.478	1.919
0.341	0.806	2.363	AA868929	troponin T1, skeletal, slow	0.237	0.417	1.758
0.106	0.261	2.472	AA036975	amine oxidase, copper containing 3 (vascular adhesion protein 1)	0.076	0.206	2.712
0.285	0.706	2.477	AA683321	EST	0.205	0.697	3.405
0.170	0.427	2.510	N53449	RAD52 (S. cerevisiae) homolog	0.124	0.286	2.312
0.081	0.207	2.562	W61116	mitogen-activated protein kinase kinase kinase 14	0.070	0.207	2.948
0.166	0.430	2.597	AA683073	synaptotagmin 1	0.117	0.276	2.351
0.184	0.480	2.611	T39411	phosphatidylinositol-4-phosphate 5-kinase, type II, alpha	0.166	0.417	2.509
0.154	0.403	2.628	H05919	eukaryotic translation initiation factor 4A, isoform 2	0.160	0.391	2.444
0.093	0.252	2.712	N62761	fragile X mental retardation, autosomal homolog 1	0.055	0.114	2.059
0.624	1.898	3.041	T94169	mitogen-activated protein kinase 8	0.299	1.173	3.922
1.032	3.326	3.222	AA478436	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily d, member 2	0.792	2.177	2.749
0.512	1.654	3.230	AA455925	four and a half LIM domains 1	0.216	0.367	1.700
0.092	0.315	3.422	AA026557	spleen focus forming virus (SFFV) proviral integration oncogene spi1	0.077	0.271	3.537
0.173	0.677	3.904	T61948	FBJ murine osteosarcoma viral oncogene homolog B	0.135	0.276	2.050
2.940	14.259	4.850	AA076063	caldesmon 1	2.392	5.195	2.172
0.182	3.247	17.824	R68106	Fc fragment of IgG, low affinity IIb, receptor for (CD32)	0.111	0.240	2.173
0.103	2.502	24.253	AA443634	ubiquitin-conjugating enzyme E2G 2 (homologous to yeast UBC7)	0.084	0.209	2.490