AffyId	Gene Title	Gene Symbol	Emg1 1416393_at
1416393_at	EMG1 nucleol	Emg1	1.33E-07
1430129_a_a	COMM domai	Commd8	0.0002946
1442083_at;	RIKEN cDNA	1500011J06R	0.0002946
1416042_s_a	vacuolar prot	Vps4b	0.0003053
1417007_a_a	RIKEN cDNA	4732418C07F	0.0003053
1427978_at;	nuclear autoa	Nasp	0.0003053
1425338_at;	phospholipas	Plcb4	0.0003238
1428496_at;	SECIS bindin	Secisbp2	0.0004459
1416894_at	UDP-N-acetyl	Galnt3	0.0007326
1417588_at;	RIKEN cDNA	1500019G21F	0.0007326
1417740_at;	cell division c	Cdc37l1	0.0007326
1424318_at;	RIKEN cDNA	1110067D22F	0.000877
1430816_at;	RIKEN cDNA	3110003A22F	0.001136
1424412_at;	opioid growth	Ogfrl1	0.001188
1460585_x_a	phosphatidyls	Pisd	0.001579
1415723_at;	eukaryotic tra	Eif5	0.001671
1431192_at;	RIKEN cDNA	2010203O03F	0.001672
1450934_at	eukaryotic tra	Eif4a2	0.001757
1418330_at;	nuclear recep	Nr3c2	0.001807
1428843_at;	CCCTC-bindin	Ctcf	0.001807
1435991_at	membrane-as	Marf5	0.001807
1423684_at;	heterogeneou	Hnrpk	0.001982
1426870_at;	F-box only pr	Fbxo33	0.002049
1428847_a_a	microtubule-a	Macf1	0.002111
1449018_at	profilin 1	Pfn1	0.002386
1423949_at;	ubiquitin-con	Ube2z	0.002545
1416911_a_a	RIKEN cDNA	6330407G11F	0.002613
1417907_at;	trinucleotide	Tnrc9	0.002888
1436600_at	ubiquitin-con	Ube2l3 /// Rp	0.002888
1416461_at;	attractin	Atrn	0.003053
1416689_at	asp (abnorma	Aspm	0.003053
1416896_at	guanine nucle	Gnb1l	0.003053
1417064_at	fucosyltransfe	Fut4	0.003053
1417202_s_a	homeo box g	Hesx1	0.003053
1418412_at	leukocyte cell	Lect1	0.003053

1418447_at	protocadherin	Pcdh10	0.003053
1418512_at;:	progesterone	Pgr	0.003053
1418905_at;:	prolactin-like	Prlpe	0.003053
1418957_at	pre-mRNA pr	Prpf8	0.003053
1419316_s_a	ribosomal pro	Rps6ka1	0.003053
1419343_at	retinoschisis :	Rs1h	0.003053
1419502_at	chemokine (C	Ccl3	0.003053
1419561_at	src homology	Stac	0.003053
1420379_at;:	tumor protein	Tpd52l1	0.003053
1420604_at	proteasome (	Psmd3	0.003053
1420780_at	tuftelin 1	Tuft1	0.003053
1420918_at;:	ubiquitin-acti	Ube1c	0.003053
1421084_at;:	chemokine (C	Xcr1	0.003053
1421166_at;:	nucleotide bir	Nubp1	0.003053
1421311_at;:	golgi autoanti	Golga5	0.003053
1421444_at;:	solute carrier	Slco1a1	0.003053
1422294_at	tropomodulin	Tmod4	0.003053
1422453_at	GPI-anchored	Gpiap1	0.003053
1422814_at;:	neuronal gua	Ngef	0.003053
1424532_at;:	serine/threon	Stk3	0.003053
1432047_at;:	solute carrier	Slc15a1	0.003053
1441447_at;:	achaete-scute	Ascl3	0.003053
1441873_at;:	jagunal homo	Jagn1	0.003053
1448479_at	GH3 domain c	Ghdc	0.003053
1448978_at	DEP domain c	Depdc2	0.003053
1449969_at	serum/glucoc	Sgk3	0.003053
1450834_at;:	nucleolar prot	Nol4	0.003053
1455219_at;:	YLP motif con	Ylpm1	0.003053
1460258_at	RIKEN cDNA :	1110030E23F	0.003053
1424015_at	Rab6 interact	Rab6ip1	0.003168
1418579_at;:	centrin 2	Cetn2	0.003236
1417166_at;:	PC4 and SFR	Psip1	0.003419
1419757_at;:	phosphatidyli	Pitpnm2	0.003419
1423734_at;:	RAS-related C	Rac1	0.003555
1433585_at;:	transportin 1	Tnpo1	0.004005
1428297_at;:	mitogen activ	Map4k2	0.004041

1434355_at;	RIKEN cDNA 2	2900019G14F	0.004041
1435101_at;	zinc finger pr	Zfp617	0.004041
1438194_at;	Der1-like don	Derl2	0.004041
1416179_a_a	radixin	Rdx	0.00408
1417054_a_a	RIKEN cDNA (	0610009D07F	0.004219
1415856_at;	paired-like ho	Phox2a	0.004274
1416258_at	crystallin, bet	Cryba1	0.004274
1417047_at;	prominin 2	Prom2	0.004274
1417459_at	S100 calcium	S100a8	0.004274
1417682_a_a	secretogranin	Scg2	0.004274
1418158_at;	sarcoglycan, a	Sgca	0.004274
1418361_at;	silver	Si	0.004274
1418466_at;	serine (or cys	Serpina1a	0.004274
1418620_at	transformatio	Trp63	0.004274
1418753_at	nudix (nucleo	Nudt8	0.004274
1419003_at;	Crn, crooked	Crnkl1	0.004274
1419389_at;	ring finger pr	Rnf141	0.004274
1419394_s_a	growth arrest	Gas8	0.004274
1419565_a_a	solute carrier	Slc35b3	0.004274
1419690_at;	aryl hydrocar	Arnt	0.004274
1419733_at	Arginine-tRNA	Ate1	0.004274
1419996_s_a	carbonic anhy	Car6	0.004274
1420036_at;	catenin (cadh	Ctnna2	0.004274
1420106_at;	demilune cell	Dcpp	0.004274
1420553_x_a	defensin beta	Defb2	0.004274
1420754_at;	embigin	Emb	0.004274
1420849_at;	glutamine fru	Gfpt2	0.004274
1421001_a_a	granzyme M (	Gzmm	0.004274
1421020_at;	caspase recru	Card14	0.004274
1421248_at	glutamate rec	Grid2ip	0.004274
1421815_at;	noggin	Nog	0.004274
1422300_at	surfactant ass	Sftpa1	0.004274
1422334_a_a	thymidine kin	Tk1	0.004274
1422523_at	protease, seri	Prss2	0.004274
1422654_at	transcription	Ttf1	0.004274
1422709_a_a	zona pellucida	Zp2	0.004274

1433970_at	blood vessel (	Bves	0.004274
1448723_at	phosphodiest	Pde10a	0.004274
1448895_a_a	synapsin III	Syn3	0.004274
1448937_at	prolactin-like	Prlpj	0.004274
1449016_at	stromal antig	Stag3	0.004274
1449501_a_a	retinol dehyd	Rdh7	0.004274
1449830_at	WD repeat do	Wdr46	0.004274
1450111_a_a	RAN binding p	Ranbp17	0.004274
1450142_a_a	RIKEN cDNA 2	2610002M06I	0.004274
1450310_at	GLI pathogen	Glipr1l2	0.004274
1450554_at	ependymin re	Epdr2	0.004274
1450671_at	DNA Segment	D9Mgi7	0.004274
1450708_at	zinc finger pr	Zfx	0.004274
1460229_at	bolA-like 3 (E	Bola3	0.004274
1422768_at;	synaptotagmi	Syncrip	0.004283
1440445_at	Pax6 opposite	Pax6os1	0.0044
1433519_at;	nuclear casei	Nucks1	0.004614
1449590_a_a	muscle and m	Mras	0.004859
1448622_at	LSM4 homolo	Lsm4	0.004913
1446636_at;	tensin 3	Tns3	0.00512
1425378_at;	hect (homolo	Herc1	0.005258
1436035_at	RIKEN cDNA 3	3830431G21I	0.005258
1428645_at;	guanine nucle	Gnai3	0.005595
1416767_a_a	RIKEN cDNA 1	1110003E01I	0.005648
1431050_at;	ribosomal pro	Rps6ka5	0.005769
1421249_at;	similar to solu	LOC674308	0.005983
1416654_at;	solute carrier	Slc31a2	0.00604
1416319_at;	adenosine kin	Adk	0.006105
1418465_at	neutrophil cyt	Ncf4	0.006105
1419974_at;	tumor necrosi	Tnfrsf21	0.006105
1422740_at;	sterol carrier	Scp2	0.006105
1416781_at;	PRA1 domain	Praf2	0.006223
1418621_at;	RAB2, membe	Rab2	0.006239
1428114_at;	solute carrier	Slc14a1	0.006275
1430265_at;	amyloid beta	Appbp2	0.006444
1426089_a_a	cDNA sequen	BC003331	0.007082

1420899_at;	RAB18, meml	Rab18	0.007231
1419917_s_a	tyrosine 3-mo	Ywhaq	0.007407
1420828_s_a	transmembra	Tmed7	0.007407
1425669_at;	MOB1, Mps O	Mobkl2b	0.007885
1428202_at;	RIKEN cDNA	1810037C20F	0.007885
1415968_a_a	kidney andro	Kap	0.007937
1420336_at;	myeloid/lymp	Mllt10	0.007937
1420869_at;	Pbx/knotted 1	Pknox1	0.007937
1421233_at;	ADAMTS-like	Adamtsl5	0.007937
1421433_at;	zinc finger ho	Zfhx4	0.007937
1433521_at;	ankyrin repea	Ankrd13c	0.007937
1415695_at;	proteasome (	Psma1	0.007939
1422597_at;	matrix metall	Mmp15	0.007939
1425354_a_a	angiogenic fa	Aggf1	0.007985
1424632_a_a	G protein-cou	Gpr103	0.008079
1433597_at;	RIKEN cDNA 9	9430010O03F	0.008079
1457048_at	REV3-like, ca	Rev3l	0.008079
1416679_at;	ATP-binding c	Abcd3	0.00814
1415686_at;	RAB14, meml	Rab14	0.008463
1422167_at;	sema domain	Sema5a	0.008761
1421835_at;	microtubule-a	Mtap7	0.009033
1433412_at;	RIKEN cDNA 5	5730446C15F	0.009097
1417250_at;	ring finger pr	Rnf12	0.00915
1416268_at;	E26 avian leu	Ets2	0.00922
1419385_a_a	ubiquilin 1	Ubqln1	0.009341
1417419_at;	cyclin D1	Ccnd1	0.009843
1435149_at;	phospholipase	Plcg1	0.009843
1428717_at;	secernin 1	Scrn1	0.009967
1415986_at;	chloride chan	Clcn4-2	0.01008
1417385_at;	aminopeptida	Npepps	0.01008
1421260_a_a	spermidine sy	Srm	0.01021
1426736_at;	G1 to S phase	Gspt1	0.01035
1417765_a_a	popeye doma	Popdc3	0.01062
1423856_at	amylase 1, sa	Amy1	0.01062
1426958_at;	ribosomal pro	Rps9 /// LOC	0.01097
1435436_at	Endothelial PA	Epas1	0.01112

1420391_at;	solute carrier	Slc35a2	0.01148
1422677_at;	diacylglycerol	Dgat2	0.01148
1422827_x_a	par-3 (partiti	Pard3	0.01148
1422781_at;	toll-like recep	Tlr3	0.01159
1428544_at;	RIKEN cDNA (	0610007L01R	0.01161
1417733_at;	solute carrier	Slc25a37	0.01188
1417750_a_a	YY1 associate	Yaf2	0.01188
1422675_at;	SWI/SNF rela	Smarce1	0.01188
1423399_a_a	ring finger pr	Rnf146	0.01188
1427430_at;	expressed se	AI848100	0.01188
1435293_at;	RIKEN cDNA 2	2900022I03R	0.01202
1430583_at;	RIKEN cDNA (	C130076O07I	0.01211
1417872_at;	four and a ha	Fhl1	0.01222
1416188_at;	Phosphatidyls	Ptdss1	0.01235
1418478_at	aldehyde deh	Aldh1a7	0.01235
1418601_at	LIM domain o	Lmo1	0.01235
1449739_at	GM2 ganglios	Gm2a	0.01235
1450959_at	RIKEN cDNA I	D930014E17F	0.01235
1450966_at	carnitine O-o	Crot	0.01245
1439251_at	Transcribed l	---	0.01253
1420477_at;	nucleosome a	Nap1l1	0.01291
1422296_at;	RIKEN cDNA !	5730470L24R	0.01291
1416381_a_a	peroxiredoxin	Prdx5	0.01294
1424185_a_a	mortality fact	Morf4l1 /// LC	0.01296
1442319_at;	ubiquitin spe	Usp4	0.01343
1416005_at	asparagine-li	Alg14	0.01363
1417149_at	protease (pro	Psmc1	0.01363
1417348_at	transmembra	Tmub1	0.01363
1417542_at;	RIKEN cDNA 2	2310039H08F	0.01363
1417720_at;	NTF2-related	Nxt1	0.01363
1417737_at;	zinc finger pr	Zfp524	0.01363
1419114_at;	RIKEN cDNA (	0610007P06F	0.01363
1419351_a_a	open reading	ORF61	0.01363
1420979_at;	polymerase (I	Polr2j	0.01363
1421754_at	mitochondrial	Mrps31	0.01363
1422488_at	cDNA sequen	AY036118	0.01363

1427109_at	procollagen-p	P4ha2	0.01363
1430577_at;:	p21 (CDKN1A	Pak1	0.01363
1448840_at	ribosomal pro	Rps6ka2	0.01363
1448915_at	centrosomal p	Cep27	0.01363
1448956_at	START domai	Stard10	0.01418
1434129_s_a	lipoma HMGI(	Lhfpl2	0.01465
1447477_at;:	RIKEN cDNA 3	3110006E14F	0.01503
1417712_at;:	eukaryotic tra	Eif2s2	0.01514
1424863_a_a	homeodomain	Hipk2	0.01519
1416573_at;:	matrix metall	Mmp24	0.01527
1417678_at;:	enolase 2, ga	Eno2	0.01527
1418829_a_a	protein O-fuc	Pofut2	0.01527
1422528_a_a	zinc finger pr	Zfp36l1	0.01537
1421416_at;:	mitogen activ	Map2k7	0.01553
1426331_a_a	FERM domain	Frmd4b	0.01562
1417182_at;:	epoxide hydr	Ephx2	0.01568
1448499_a_a	DnaJ (Hsp40)	Dnaja2	0.01568
1434083_a_a	ELMO domain	Elmod1	0.01578
1416244_a_a	tetratricopept	Ttc19	0.01601
1427114_at;:	cellular nucle	Cnbp	0.01601
1437354_at;:	RIKEN cDNA (	C230091D08I	0.01601
1416186_at;:	proline-rich n	Pnrc2	0.01648
1424293_s_a	transmembra	Tmem55a	0.01686
1426743_at	Dip3 beta	Dip3b	0.01717
1418060_a_a	mitogen activ	Mapk7	0.01739
1420844_at;:	zinc finger pr	Zfp180	0.01739
1428170_at	ubiquilin 2	Ubqln2	0.01739
1443175_at	RIKEN cDNA A	A830010M09I	0.01739
1415775_at;:	retinoblastom	Rbbp7	0.01762
1435151_a_a	ribosomal pro	Rps3	0.01767
1431995_at;:	anthrax toxin	Antxr1	0.01789
1426554_a_a	phosphoglyce	Pgam1 /// LO	0.0179
1419568_at;:	mitogen activ	Mapk1	0.01815
1423994_at;:	succinate deh	Sdha	0.01824
1426688_at;:	kinesin family	Kif1b	0.01824
1454671_at	insulin induce	Insig1	0.01868

1423332_at;	syndecan bin	Sdcbp	0.01882
1431369_at;	development	Ddef2	0.01883
1417536_at;	zinc finger, m	Zmat2	0.0194
1419585_at;	retinitis pigm	Rp2h	0.0194
1416028_a_a	hematologica	Hn1	0.01965
1420365_a_a	heterogeneou	Hnrpa2b1	0.01977
1434549_at;	---	---	0.01978
1420472_at;	myotrophin	Mtpn	0.01991
1419291_x_a	growth arrest	Gas5	0.02001
1415916_a_a	telomeric rep	Terf2ip	0.02018
1419381_at;	methylenetet	Mthfd1	0.02018
1428277_at	ADAMTS-like	Adamtsl1	0.02018
1430313_at;	methionine a	Mat2b	0.02018
1448196_at	OTU domain	Otud6b	0.02018
1418526_at;	FUS interactir	Fusip1	0.02175
1428236_at;	zinc finger pr	Zfp192	0.02183
1430334_at;	acyl-Coenzym	Acbd5	0.02183
1423112_at;	ubiquitin-con	Ube2d3	0.02264
1432391_at;	coiled-coil do	Ccdc21	0.02287
1435475_at;	lectin, manno	Lman2l	0.02292
1417723_at;	ubiquitin-con	Ube2j1	0.02303
1423154_at;	tetratricopept	Ttc14	0.02327
1426544_a_a	cDNA sequen	BC005537	0.02327
1430592_at;	transmembra	Tmem106b	0.02327
1422664_at;	RAB10, meml	Rab10	0.02343
1429537_at;	RIKEN cDNA !	5730406M06I	0.02357
1415920_at	cell adhesion	Cdon	0.02395
1421452_at;	HESB like dor	Hbld2	0.02395
1423651_at;	cleavage stim	Cstf2t	0.02395
1424873_at;	ring finger pr	Rnf2	0.02395
1420694_a_a	dachshund 1	Dach1	0.02422
1424365_at	RIKEN cDNA	1810037I17R	0.02436
1427278_at;	restin-like 2	Rsnl2	0.02472
1424215_at;	FUN14 domai	Fundc1	0.02487
1435828_at;	avian muscul	Maf	0.02504
1420131_s_a	pituitary tum	Pttg1ip	0.02543

1417367_at;	protein phosp	Ppp2ca	0.02597
1430988_at;	RIKEN cDNA 2	2810407C02F	0.02611
1423503_at;	bolA-like 2 (E	Bola2	0.02625
1434543_a_a	junction adhe	Jam3	0.02625
1434558_at;	WD repeat do	Wdr47	0.02625
1436333_a_a	synaptojanin	Synj1	0.02661
1422879_at;	synaptophysii	Sypl	0.02681
1434082_at;	---	---	0.02741
1428817_at;	patched hom	Ptch1	0.02762
1428853_at;	protocadherin	Pcdh9 /// LOC	0.02762
1429861_at;	MON2 homolo	Mon2	0.02762
1428140_at;	3-oxoacid Co/	Oxct1	0.02766
1428835_at	myosin, heav	Myh14 /// LO	0.02771
1427408_a_a	thyroid horm	Thrap3	0.02772
1419736_a_a	eukaryotic tra	Eif1ay	0.02821
1433816_at;	mitochondrial	Mcart1	0.02851
1423517_at;	chaperonin su	Cct6a	0.02934
1416185_a_a	alcohol dehyd	Adh5	0.03039
1416531_at;	calcium chan	Cacna1s	0.03175
1417050_at;	cellular retino	Crabp1	0.03175
1418650_at;	glutathione S	Gsto1	0.03175
1420442_at	ubiquitin-like	Ubl3	0.03175
1420679_a_a	ankyrin repea	Ankrd57	0.03175
1421642_a_a	androgen-ind	Aig1	0.03175
1423461_a_a	C1q and tum	C1qtnf4	0.03175
1429974_at;	spermatogen	Spata6	0.03175
1431670_at	NADH dehydr	Ndufc2	0.03175
1448326_a_a	cysteinyl leuk	Cysltr2	0.03175
1454736_at	T-box18	Tbx18	0.03175
1424118_a_a	spindle pole b	Spbc25	0.03209
1428365_a_a	protease, seri	Prss15	0.03209
1431254_at;	kelch repeat a	Kbtbd11 /// L	0.03263
1426945_at;	RAN binding p	Ranbp5	0.0327
1428078_at;	peptidylprolyl	Ppil6	0.0327
1453310_at	RIKEN cDNA (	0610013E23F	0.0327
1416610_a_a	chloride chan	Clcn3	0.0337

1416201_at;	tubulin, beta	Tubb5	0.03376
1416256_a_a	v-crk sarcom	Crk	0.03376
1418323_at;	dual specificit	Dusp12	0.03428
1420636_a_a	Rho GTPase a	Arhgap17	0.03428
1426623_a_a	feminization	Fem1b	0.03428
1428997_at	PHD finger pr	Phf23	0.03438
1416555_at	phosphodiest	Pde7a	0.03538
1423313_at;	etoposide ind	Ei24	0.03538
1427876_at;	RIKEN cDNA 2	2610312B22F	0.03588
1434301_at	RIKEN cDNA I	D330050I23R	0.036
1451665_a_a	RIKEN cDNA 2	2610014I16R	0.03623
1455135_at	adaptor-relat	Ap4s1	0.03623
1420522_at;	coiled-coil do	Ccdc50	0.03625
1433724_at;	DNA segment	D15Ertd621e	0.03631
1416105_at;	solute carrier	Slc9a6	0.03638
1433975_at	cyclin-depenc	Cdk10	0.03638
1435008_at;	nicotinamide	Nnt	0.03638
1415697_at;	RIKEN cDNA I	E430034L04R	0.03648
1423297_at;	adducin 3 (ga	Add3	0.03691
1423681_at;	RIKEN cDNA 1	1300018I05R	0.03787
1435783_at	RIKEN cDNA I	B230112C05F	0.038
1444511_at;	solute carrier	Slc41a1	0.03816
1422045_a_a	protein tyrosi	Ptpn12	0.03833
1423240_at;	Rous sarcom	Src	0.03881
1434250_at;	p21 (CDKN1A	Pak2	0.03881
1415796_at	DAZ associate	Dazap2	0.03923
1431989_at;	glutamate rec	Grin2c	0.03968
1449245_at	RIKEN cDNA 4	4930433E13F	0.03968
1421871_at;	SH3-binding (	Sh3bgrl	0.04045
1425241_a_a	WD repeat an	Wsb1	0.04117
1416884_at;	chromobox h	Cbx3 /// LOC	0.04194
1417637_a_a	high mobility	Hmg20b	0.04194
1451075_s_a	CTD (carboxy	Ctdsp2	0.04195
1421866_at;	RIKEN cDNA (	C730029A08F	0.04197
1434860_at	NMDA recept	Narg3	0.04197
1436168_at	nuclear recep	Nr3c1	0.04197

1437632_at;	Adult male ce	---	0.04197
1420610_at;	protein kinas	Prkacb	0.04203
1423091_a_a	glycoprotein r	Gpm6b	0.04211
1415858_at;	capping prote	Capza2	0.04213
1423057_at;	eukaryotic tra	Eif3s8	0.04213
1447776_x_a	RAB6, membe	Rab6	0.04213
1421117_at;	ethanolamine	Etnk1	0.0423
1430996_at;	dystonin	Dst	0.0423
1427456_at;	WD repeat an	Wdfy3	0.04244
1434418_at;	longevity assu	Lass6	0.04255
1423187_at;	gamma-amin	Gabarapl2	0.04269
1423451_at	progesterone	Pgrmc1	0.04324
1417343_at	FXYD domain	Fxyd6	0.04329
1420953_at;	adducin 1 (al	Add1	0.04392
1434702_at;	RIKEN cDNA 2	2600009E05F	0.04464
1431385_a_a	membrane-bo	Mbtps1	0.04476
1418937_at;	deiodinase, io	Dio2	0.04506
1433552_a_a	polymerase (I	Polr2b	0.04642
1426401_at;	protein phosp	Ppp3ca	0.04647
1417349_at;	copine III	Cpne3	0.04667
1428514_at;	pallidin	Pldn	0.04667
1420958_at;	---	---	0.04682
1431404_at;	transmembra	Tmem56	0.04728
1441850_x_a	transcobalam	Tcn2	0.04736
1415708_at;	taurine upreg	Tug1	0.04798
1417690_at;	chordin-like 1	Chrdl1	0.04838
1421295_at;	vacuolar prot	Vps11	0.04838
1422937_at;	protein kinas	Prkag1	0.04838
1423103_at	frizzled homo	Fzd5	0.04838
1428109_at;	regulatory fac	Rfx5	0.04838
1425114_at;	retinoblastom	Rbbp6	0.04859
1427004_at	F-box only pr	Fbxo2	0.04859
1425382_a_a	aquaporin 4	Aqp4	0.049
1419255_at;	spectrin beta	Spnb2	0.05036
1425567_a_a	annexin A5	Anxa5	0.05038
1424822_at;	RIKEN cDNA 9	9630044O09F	0.05043

1421908_a_a	transcription	Tcf12	0.05045
1424033_at;	splicing factor	Sfrs7	0.05097
1415946_at;	phosphatidyli	Pigq	0.05106
1434300_at;	RIKEN cDNA 2	2610101N10F	0.05134
1426531_at;	zinc finger, M	Zmynd11	0.05144
1416967_at	SRY-box cont	Sox2	0.05185
1416556_at;	tetraspanin 3	Tspan31	0.05245
1416037_a_a	chaperonin su	Cct2	0.05272
1417846_at;	RIKEN cDNA 2	2210419D22F	0.05282
1425037_at;	CDC14 cell di	Cdc14a	0.05282
1426208_x_a	pleiomorphic	Plagl1	0.05282
1428500_at	Unc-51 like ki	Ulk2	0.05282
1436913_at;	FYVE, RhoGEF	Fgd4	0.05282
1417999_at;	integral mem	Itm2b	0.05345
1415870_at;	calumenin	Calu	0.05541
1418349_at;	sodium chanr	Scn7a	0.05607
1427495_at;	RIKEN cDNA A	A230083H22F	0.05607
1439274_at;	heparin-bindi	Hbegf	0.05607
1420115_at;	exosome com	Exosc8	0.05684
1415700_a_a	signal sequen	Ssr3	0.05698
1415701_x_a	ribosomal pro	Rpl23	0.05709
1418124_at;	transmembra	Tmem85	0.05709
1435106_at;	RIKEN cDNA 3	3732412D22F	0.05751
1416081_at;	MAD homolog	Smad1	0.05773
1426414_a_a	EF hand dom	Efhd1	0.05777
1426776_at;	protein phosp	Ppp4r2 /// LC	0.05777
1433850_at;	ring finger pr	Rnf7	0.05777
1448507_at	Wiskott-Aldri	Wasl	0.05777
1418066_at;	cofilin 2, mus	Cfl2	0.05792
1423564_a_a	phosphoribos	Paics	0.05799
1425742_a_a	TSC22 domai	Tsc22d1	0.05822
1433752_s_a	RIKEN cDNA I	D030016E14F	0.05849
1423680_at;	fatty acid des	Fads1	0.05853
1433706_a_a	protein tyrosi	Ptplad1	0.05856
1422764_at;	microtubule-a	Mapre1	0.05863
1415692_s_a	NADH dehydr	Ndufb10	0.05897

1423641_s_a	calnexin	Canx	0.05897
1428322_a_a	CCR4-NOT tra	Cnot7	0.05897
1423846_x_a	tubulin, alpha	Tuba2	0.05979
1416323_at;	potassium ch	Kctd20	0.06014
1416509_at;	transmembra	Tm9sf3	0.06035
1426442_at;	glycoprotein r	Gpm6a	0.06041
1416941_s_a	Williams-Beu	Wbscr1	0.06046
1419734_at;	actin, beta, c	Actb	0.06141
1426994_at	PH domain ar	Phlpp	0.06157
1417267_s_a	FK506 bindin	Fkbp11	0.06165
1423662_at;	ATPase, H+ t	Atp6ap2	0.06166
1433985_at;	abl-interactor	Abi2	0.06206
1424636_at;	RIKEN cDNA 2	2610204L23R	0.06287
1426668_at	UDP-N-acetyl	Galnt13	0.06298
1439899_at;	solute carrier	Slc30a9	0.06298
1423100_at	FBJ osteosarc	Fos	0.06341
1416211_a_a	pleiotrophin	Ptn	0.06393
1426124_a_a	CDC-like kina	Clk1	0.06408
1433561_at;	centaurin, be	Centb2	0.06444
1421828_at;	ATPase, Na+/	Atp1b3	0.06453
1423126_at	karyopherin (	Kpna3	0.06453
1421887_a_a	amyloid beta	Aplp2	0.06475
1424342_at;	zinc finger an	Zbtb4	0.06558
1429722_at;	forty-two-thre	Fyttd1	0.06558
1415984_at	acyl-Coenzym	Acadm	0.06576
1435895_at;	RIKEN cDNA I	D930023J12R	0.06576
1427048_at;	smoothened h	Smo	0.06604
1429107_at;	RIKEN cDNA 1	1110059H15R	0.06671
1439159_at;	---	---	0.06779
1416144_a_a	DEAH (Asp-G	Dhx15	0.068
1432143_a_a	high mobility	Hbp1	0.06813
1416315_at;	abhydrolase c	Abhd4	0.06876
1424950_at;	SRY-box cont	Sox9	0.06886
1427310_at;	fetal Alzheime	Falz	0.06918
1423052_at;	ADP-ribosylat	Arf4	0.06983
1418285_at;	IMP1 inner m	Immp1l	0.07005

1435743_at	kelch-like 23	Klhl23	0.07005
1438409_at;	ephrin B1	Efnb1	0.07005
1445473_at;	centrosomal p	Cep63	0.07005
1451217_a_a	5-azacytidine	Azi2 /// Diras	0.07005
1416723_at;	transcription	Tcf4	0.0704
1416476_a_a	ubiquitin-con	Ube2d2	0.07096
1433490_s_a	erythrocyte p	Epb4.1l2	0.07124
1416922_a_a	BCL2/adenovi	Bnip3l	0.07128
1419574_at;	zinc finger pr	Zfp292	0.07228
1433754_at;	muscleblind-l	Mbnl2	0.07299
1428163_at	SAR1 gene ho	Sar1b	0.07329
1420058_s_a	contactin 1	Cntn1	0.07469
1426029_a_a	RIKEN cDNA 2	2410166I05R	0.07469
1449563_at;	nuclear factor	Nfat5	0.07469
1426997_at;	thyroid hormo	Thra	0.07474
1421933_at;	chromobox ho	Cbx5	0.07523
1452953_at	family with se	Fam18b	0.07551
1416027_at	programmed	Pdcd6	0.07612
1431766_x_a	ribosomal pro	Rps2	0.07637
1435740_at	Retinoic acid	Rai17	0.07763
1419616_at;	bone morpho	Bmpr2	0.07834
1415866_at;	RIKEN cDNA (	6030458C11F	0.07863
1426018_a_a	Rho GTPase a	Arhgap26	0.07863
1428013_at;	unc-45 homo	Unc45a	0.07863
1430957_at;	SRY-box cont	Sox6	0.07863
1417661_at	RAD52 motif	Rdm1	0.07935
1429040_at;	RIKEN cDNA 2	2610005L07R	0.07968
1426340_at;	solute carrier	Slc1a3	0.07997
1416498_at	peptidylprolyl	Ppic	0.08027
1423220_at;	eukaryotic tra	Eif4e /// LOC	0.08047
1435593_x_a	ribosomal pro	Rps7	0.08085
1416056_a_a	NADH dehydr	Ndufb11	0.08108
1419640_at;	purine rich el	Purb	0.08115
1437060_at	olfactomedin	Olfm4	0.0812
1419449_a_a	guanine nucle	Gnai2	0.08207
1425411_at;	ADP-ribosylat	Arl4a	0.08236

1444086_at	RIKEN cDNA I	E030049G20F	0.08354
1427468_at;	protein phosp	Ppp3cb	0.0836
1418663_at;	multiple PDZ	Mpdz	0.08395
1422017_s_a	RIKEN cDNA 4	4833439L19F	0.08395
1427129_a_a	heterogeneou	Hnrpr	0.08402
1423834_s_a	golgi associat	Gga1	0.08443
1425810_a_a	cysteine and (	Csrp1	0.08495
1438984_x_a	proteasome (	Psmb4	0.08511
1416170_at;	TNF receptor-	Trap1	0.08548
1450880_at	mitochondrial	Mrpl16	0.08582
1417794_at;	zinc finger, M	Zmym3	0.08593
1424635_at	eukaryotic tra	Eef1a1	0.08621
1421927_at;	RIKEN cDNA I	E130218I03R	0.08669
1435318_at	transducin (b	Tbl2	0.08669
1422504_at;	glycine recepl	Glrb	0.08671
1428386_at;	acyl-CoA synt	Acsl3	0.08678
1415778_at;	mortality fact	Morf4l2	0.08704
1423599_a_a	phosducin-lik	Pdcl	0.08749
1430123_a_a	aldo-keto red	Akr1a4	0.08778
1428011_a_a	Erbb2 interac	Erbb2ip	0.0879
1415879_a_a	ribosomal pro	Rplp2	0.08796
1426698_a_a	heterogeneou	Hnrpm	0.09075
1419371_s_a	golgi SNAP re	Gosr2	0.09081
1452596_at	polymerase (I	Polr2k	0.09109
1423200_at;	glutamate-am	Glul	0.09274
1425142_a_a	heterogeneou	Hnrpd	0.09274
1426235_a_a	nuclear recep	Ncor1	0.09274
1434108_at	F-box protein	Fbxo11	0.09274
1417512_at;	glucosamine-	Gnpda1 /// L(	0.09341
1417912_at;	sorting nexin	Snx10	0.09341
1418427_at;	platelet-activ	Pafah1b2	0.09341
1418827_at;	Lipin 2	Lpin2	0.09341
1421151_a_a	Eph receptor	Epha2	0.09341
1422791_at;	ecotropic vira	Evi5	0.09341
1431055_a_a	kinesin family	Kif5b	0.09341
1446316_at;	transmembra	Tmem93	0.09341

1448163_at;	three prime h	Thex1	0.09341
1416288_at;	DnaJ (Hsp40)	Dnaja1	0.09357
1424657_at;	TAO kinase 1	Taok1	0.09393
1440120_at;	Guanine nucl	Gnb2l1	0.09401
1460196_at	carbonyl redu	Cbr1	0.09403
1421624_a_a	enabled hom	Enah	0.09488
1418010_a_a	SH3-domain (	Sh3glb1	0.09558
1433751_at;	solute carrier	Slc39a10	0.09562
1448418_s_a	WD repeat do	Wdr23	0.09573
1417230_at;	Ral GEF with	Ralgps2	0.09599
1434235_at;	solute carrier	Slc20a2	0.09599
1453638_at;	RIKEN cDNA 9	9030420J04R	0.09599
1423812_s_a	expressed se	AW146242	0.09601
1423545_a_a	zinc finger pr	Zfp207	0.09657
1425554_a_a	ring finger pr	Rnf6	0.0966
1427898_at;	CDC16 cell di	Cdc16	0.0966
1430543_at;	RIKEN cDNA 1	1500005P14F	0.09725
1420614_at;	dynein light c	Dynlt3	0.09726
1417427_at;	RIKEN cDNA 4	4933428P19F	0.09744
1426100_a_a	Eph receptor	Ephb1	0.09744
1428273_at;	thymidine kin	Tk2	0.09744
1432593_at;	protein kinas	Prkg1	0.09744
1444232_at;	RIKEN cDNA 1	1500026D16F	0.09744
1455188_at	abhydrolase c	Abhd13	0.09744
1422694_at;	tweety homol	Ttyh1	0.09746
1438992_x_a	activating tra	Atf4	0.09749
1433770_at;	dihydropyrim	Dpysl2	0.09751
1425628_a_a	similar to Ger	LOC669007	0.09779
1420174_s_a	---	---	0.0985
1426645_at;	heat shock pr	Hsp90aa1	0.0985
1420592_a_a	acidic (leucin	Anp32e	0.09892
1419088_at;	tissue inhibit	Timp3	0.09915
1418716_at;	mitochondrial	Mrps25	0.09918
1415981_at;	HERPUD fami	Herpud2	0.1012
1423177_a_a	abl-interactor	Abi1	0.1024
1419247_at;	regulator of G	Rgs2	0.1027

1423083_at	RAB33B, mem	Rab33b		0.1028
1428782_a_a	ubiquinol-cyt	Uqcrc1		0.1029
1428286_at;	---	---		0.1039
1422455_s_a	heterogeneou	Hnrpu		0.104
1423050_s_a	RIKEN cDNA 2	2310057D15F		0.104
1423705_at;	N-ethylmaleir	Nsf		0.104
1432164_a_a	glycine cleava	Gcsh /// LOC		0.1042
1426404_a_a	ring finger pr	Rnf11		0.1065
1417952_at;	PHD finger pr	Phf17		0.1068
1421509_at;	cytochrome P	Cyp2j6		0.1068
1426752_at;	VPS10 domai	Sorcs1		0.1068
1425162_at;	RAR-related c	Rorb		0.1071
1427773_a_a	Rab acceptor	Rabac1		0.1073
1425550_a_a	protein kinase	Prkar1a		0.1077
1415840_at;	ELOVL family	Elovl5		0.1083
1418297_at;	dihydropyrim	Dpysl4		0.1084
1448559_at	flotillin 1	Flot1		0.1084
1423472_at;	septin 2		40424	0.1085
1427060_at	mitogen activ	Mapk3		0.1092
1451596_a_a	sphingosine k	Sphk1		0.1107
1417663_a_a	N-myc downs	Ndrg3		0.111
1433442_at	Kelch-like 9 (	Klhl9		0.1131
1436319_at;	sulfatase 1	Sulf1		0.1133
1423723_s_a	TAR DNA binc	Tardbp		0.1151
1417060_at;	solute carrier	Slc7a2		0.1157
1422648_at;	protein phosp	Ppp1r11		0.1157
1450280_a_a	retinal pigme	Rrh		0.1158
1425516_at;	O-linked N-ac	Ogt		0.1165
1421148_a_a	Tial1 cytotoxi	Tial1		0.1168
1416112_at;	cytochrome c	Cox8a		0.117
1440905_at;	Heparan sulfa	Hs2st1		0.1173
1423839_a_a	basic transcri	Btf3		0.1176
1428573_at;	chimerin (chi	Chn2		0.1177
1424979_at;	anterior phar	Aph1a		0.118
1423394_at;	prenylcystein	Pcyox1		0.1191
1428619_at;	RIKEN cDNA 2	2310005N03F		0.1192

1451695_a_a	glutathione p	Gpx4	0.1193
1418374_at	discs, large h	Dlg5	0.1205
1424198_at;	FXYD domain	Fxyd3	0.1205
1415990_at	janus kinase	Jakmip1	0.1217
1430597_at;	voltage-depe	Vdac2	0.1217
1416161_at;	RAD21 homol	Rad21	0.1244
1421955_a_a	neural precur	Nedd4	0.1249
1422321_a_a	splicing facto	Sf1	0.1249
1422226_at;	---	---	0.1264
1431805_a_a	rhophilin, Rh	Rhpn2	0.1266
1430526_a_a	SWI/SNF rela	Smarca2	0.127
1451572_a_a	RIKEN cDNA	5230400G24	0.1272
1423632_at;	G protein-cou	Gpr146	0.1278
1437626_at;	zinc finger pr	Zfp36l2	0.1284
1423791_at;	IK cytokine	Ik	0.1294
1423215_at;	signal peptid	Spcs2	0.1299
1416720_at;	splicing facto	Sfrs6	0.1302
1425914_a_a	transient rece	Trpm3	0.1302
1439026_at;	armadillo rep	Armcx1	0.1302
1416855_at;	growth arrest	Gas1	0.1305
1428804_at;	eukaryotic tra	Eif5a	0.1307
1437859_x_a	microfibrillar-	Mfap3l	0.1307
1424309_a_a	molybdenum	Mocs2	0.131
1421198_at;	integrin alpha	Itgav	0.1314
1423640_at	synaptoporin	Synpr	0.1314
1425732_a_a	Max interacti	Mxi1	0.1314
1428470_at	exocyst comp	Exoc2	0.1314
1443651_at;	Polyhomeotic	Phc3	0.1314
1415894_at;	ectonucleotid	Enpp2	0.1321
1429841_at	heterogeneou	Hnrpa3	0.1323
1452712_at;	multiple EGF-	Megf10	0.1323
1435987_x_a	RIKEN cDNA	1110059G02	0.1324
1428231_at;	cleavage and	Cpsf6	0.1325
1417923_at;	p21 (CDKN1A	Pak3	0.1328
1449145_a_a	caveolin, cav	Cav1	0.1339
1418436_at	syntaxin 7	Stx7	0.134

1423895_a_a	CUG triplet re	Cugbp2	0.1343
1420821_at;	sphingosine-1	Sgpp1	0.1346
1447720_x_a	protein kinas	Prkaca	0.1353
1418117_at;	NADH dehydr	Ndufs4	0.1358
1434735_at;	hepatic leuke	Hlf	0.1358
1428256_at;	RIKEN cDNA 2	2310047H23	0.1365
1417516_at;	DNA-damage	Ddit3	0.1379
1417718_at	eukaryotic tra	Eif3s4	0.1388
1420811_a_a	catenin (cadh	Ctnnb1	0.1388
1438081_at;	mutated in co	Mcc	0.1391
1445208_at;	exonuclease 3	Exdl2	0.1391
1451336_at	lectin, galacto	Lgals4	0.1397
1417693_a_a	Cd99 antigen	Cd99l2	0.1405
1423965_at;	growth factor	Gab1	0.1405
1425457_a_a	RIKEN cDNA 4	4930402H24	0.1412
1429694_at;	growth factor	Grb10	0.1412
1424311_at	RIKEN cDNA 0	0710008K08	0.142
1434335_at;	expressed se	AI317237	0.142
1456878_at	expressed se	AI646023	0.1424
1421162_a_a	nuclear factor	Nfia	0.1434
1418505_at;	nudix (nucleo	Nudt4	0.1438
1419454_x_a	protein inhibi	Pias2	0.1443
1417769_at;	---	---	0.1447
1419822_at;	Eukaryotic tra	Eif3s6	0.1456
1439094_at;	phosphoserin	Psat1	0.1468
1451064_a_a	clathrin, heav	Cltc	0.1468
1419601_at	potassium inv	Kcnj10	0.1474
1423911_at;	protein phosp	Ppp2r5a	0.1475
1435865_at;	histone 3, H2	Hist3h2a	0.1484
1418814_s_a	NADH dehydr	Ndufa12	0.1487
1415706_at;	coatomer pro	Copa	0.1493
1421860_at;	calsyntenin 1	Clstn1	0.1518
1426497_at;	jumonji, AT ri	Jarid1c	0.1526
1416551_at;	ATPase, Ca++-	Atp2a2	0.1527
1417802_at;	UPF3 regulato	Upf3b	0.1538
1420140_at;	bin3, bicoid-in	Bcdin3	0.1538

1434842_s_a	RIKEN cDNA :	1110032A04F		0.1538
1438758_at;	ADP-ribosylat	Arl1		0.1538
1447089_at;	---	---		0.1538
1427117_at;	myotubularin	Mtmr3		0.154
1431337_a_a	RIKEN cDNA :	1810055E12F		0.154
1431424_at;	RIKEN cDNA :	2810055G20F		0.154
1455286_at;	BTB (POZ) do	Btbd1		0.1559
1423120_at;	insulin degra	Ide		0.157
1452942_at;	transmembra	Tmem65		0.157
1416749_at;	HtrA serine p	Htra1		0.1576
1423745_at;	RIKEN cDNA :	1110031B06F		0.1576
1416945_at;	cathepsin O	Ctso		0.1577
1433501_at;	prostate tum	Ptov1		0.1577
1417830_at;	---	---		0.1583
1444021_at;	structural ma	Smc1a		0.1583
1423106_at;	ubiquitin-con	Ube2b		0.1587
1423345_at;	degenerative	Degs1		0.1587
1425667_at;	acyl-CoA thio	Acot11		0.1592
1421413_a_a	PDZ and LIM	Pdlim5 /// LO		0.1593
1432055_at;	SAM and SH3	Sash1		0.1594
1419963_at;	DEP domain c	Depdc6		0.1601
1418253_a_a	heat shock pr	Hspa4l		0.1607
1416934_at;	X-linked myo	Mtm1		0.162
1421120_at;	myosin VI	Myo6		0.1632
1437995_x_a	septin 7		40429	0.1638
1423782_at;	MOB1, Mps O	Mobk1b		0.1641
1431285_at;	mahogunin, r	Mgrn1		0.1645
1416496_at	Morf4 family	Mrfap1		0.1652
1421966_at;	neural cell ad	Ncam1		0.1657
1419408_at	sine oculis-re	Six6		0.1661
1424312_at;	adiponectin r	Adipor1		0.1661
1423223_a_a	peroxiredoxin	Prdx6		0.1677
1416589_at;	monocyte to	Mmd2		0.1681
1424534_at;	secreted acid	Sparc		0.1681
1415734_at;	disabled hom	Dab1		0.1689
1416998_at;	RAB7, memb	Rab7		0.1689

1417011_at;	5'-nucleotidas	Nt5c3	0.1689
1417932_at	RIKEN cDNA 2	2900052N01F	0.1689
1418667_at;	syndecan 2	Sdc2	0.1689
1421100_a_a	RRS1 ribosom	Rrs1	0.1689
1423591_at;	RIKEN cDNA 2	2410002O22F	0.1689
1431306_at;	FGFR1 oncoge	Fgfr1op2	0.1689
1435223_at	interleukin 18	Il18	0.1689
1436231_at;	SPFH domain	Spfh2	0.1689
1424768_at;	caldesmon 1	Cald1	0.169
1416818_at;	parvin, alpha	Parva	0.1706
1420982_at;	RIKEN cDNA 4	4930570C03F	0.1707
1450409_a_a	RNA binding r	Rbm39	0.1707
1423773_at;	GC-rich prom	Gpbp1	0.1724
1416024_x_a	chaperonin su	Cct3	0.1735
1415807_s_a	splicing factor	Sfrs2	0.1739
1418563_at;	Serpine1 mRI	Serbp1	0.1739
1423692_at	NADH dehydr	Ndufa8	0.1739
1423654_a_a	ring finger pr	Rnf4	0.1742
1434866_x_a	carnitine paln	Cpt1a	0.1749
1427371_at	ATP-binding c	Abca8a	0.1769
1434021_at;	RIKEN cDNA (	C230096C10F	0.1772
1415740_at	protease (pro	Psmc5	0.1773
1421139_a_a	zinc finger pr	Zfp386	0.1774
1420858_at;	protein kinase	Pkia	0.178
1435770_at;	thioredoxin d	Txndc13	0.1792
1436798_at;	ribosomal pro	Rpl9 /// LOC4	0.1801
1424562_a_a	solute carrier	Slc25a4	0.1812
1423133_at	RIKEN cDNA (	0610040D20F	0.1817
1432466_a_a	apolipoprotein	Apoe	0.1821
1433516_a_a	myeloma ove	Myeov2	0.1824
1416036_at;	FK506 bindin	Fkbp1a	0.1826
1434063_at;	zinc finger pr	Zfp664	0.1826
1420878_a_a	tyrosine 3-mo	Ywhab	0.1828
1418694_at;	potassium ch	Kcmf1	0.1838
1424827_a_a	casein kinase	Csnk1a1	0.1845
1430511_at;	RIKEN cDNA 1	1500037O19F	0.1871

1433676_at;	WNK lysine d	Wnk1	0.1873
1416427_at;	cyclin I	Ccni	0.1881
1423295_at;	transmembra	Tm9sf2	0.1883
1417142_at;	kinesin family	Kif2a	0.1884
1426326_at;	RIKEN cDNA 4	4932442K08F	0.1884
1432072_at;	zinc finger pr	Zfp91	0.1884
1420621_a_a	amyloid beta	App	0.1888
1420920_a_a	ADP-ribosylat	Arf1	0.1907
1426983_at;	formin bindin	Fnbp1	0.1923
1435240_at;	bromodomain	Baz2b	0.193
1416250_at;	B-cell translo	Btg2	0.1938
1428431_at;	RIKEN cDNA 2	2310047A01F	0.1942
1417494_a_a	ceruloplasmin	Cp	0.1962
1417001_a_a	DNA segment	D4Wsu53e	0.1966
1416659_at;	eukaryotic tra	Eif3s10	0.1971
1423961_at;	WD repeat do	Wdr26	0.1987
1416208_at;	ubiquitin spec	Usp14	0.1991
1449526_a_a	glycerophospl	Gdpd3	0.1991
1417431_a_a	SLIT-ROBO R	Srgap3	0.2008
1424588_at;	sphingosine k	Sphk2	0.2008
1417152_at;	BTB (POZ) do	Btbd14a	0.2009
1423795_at;	splicing factor	Sfpq	0.2009
1423879_at;	RIKEN cDNA I	D030056L22F	0.2009
1427009_at;	CGG triplet re	Cggbp1	0.2009
1454641_at;	laminin, alpha	Lama5 /// LO	0.2009
1423349_at;	suppressor of	Socs5	0.2031
1416465_a_a	vesicle-associ	Vapa	0.2061
1422946_a_a	DNA methyltr	Dnmt1	0.2063
1434604_at;	eukaryotic tra	Eif5b	0.2063
1455106_a_a	creatine kinas	Ckb	0.2079
1424108_at;	leucine-rich r	Lgr4 /// LOC6	0.2087
1433891_at	glyoxalase 1	Glo1	0.2087
1434515_at;	nuclear recep	Ncoa1	0.2104
1419628_at	C. elegans ce	Chx10	0.2107
1422449_s_a	polymerase (I	Pole4	0.2109
1423371_at;	reticulocalbin	Rcn2	0.2109

1416985_at;	signal-regulat	Sirpa	0.2113
1419650_at;	zinc finger RN	Zfr	0.2116
1434156_at;	---	---	0.2118
1437396_at;	cAMP respons	Creb3l2	0.2125
1428314_at;	PEST proteoly	Pcnp	0.2126
1429314_at;	synaptotagmi	Syt11	0.2135
1421215_a_a	sarcolemma a	Slmap	0.2142
1418102_at	thioredoxin d	Txndc11	0.2143
1421282_at;	general trans	Gtf2e1	0.2143
1428124_at;	hairy and enh	Hes1	0.2143
1448782_at;	bone morpho	Bmp5	0.2143
1420910_at;	phosphatidic	Ppap2c	0.216
1436631_at;	RIKEN cDNA 2	2010010M04I	0.2164
1417428_at	guanine nucle	Gng3	0.2169
1422615_at;	mitogen-activ	Map4k4	0.2179
1420142_s_a	ATP synthase	Atp5g3	0.2196
1454661_at	proliferation-a	Pa2g4	0.2196
1420123_at;	T-cell leukem	Tcta	0.2201
1434319_at;	malate dehyd	Mdh1	0.2205
1421789_s_a	ADP-ribosylat	Arf3	0.2209
1452142_at;	solute carrier	Slc6a1	0.2225
1448122_at	t-complex pro	Tcp1	0.2236
1425576_at;	S-adenosylho	Ahcyl1	0.2245
1420833_at;	vesicle-associ	Vamp2	0.2257
1423408_a_a	methionine a	Metap2	0.2289
1423610_at;	RIKEN cDNA 2	2500003M10I	0.2289
1426053_a_a	xenotropic an	Xpr1	0.2302
1427270_a_a	BSD domain c	Bsdc1	0.2315
1416719_a_a	ribosomal pro	Rps10 /// LO	0.2317
1425965_at;	ubiquitin C	Ubc	0.2321
1423615_at;	zinc finger pr	Zfp364	0.2326
1442767_s_a	ubiquitin-acti	Ube1x	0.2343
1436421_s_a	actin related	Arpc5l	0.2387
1432270_a_a	chromatin mo	Chmp5	0.2428
1420042_at;	THO complex	Thoc1	0.2433
1451366_at;	COP9 (constit	Cops6	0.2462

1427020_at	scavenger rec	Scara3	0.247
1436179_a_a	DnaJ (Hsp40)	Dnajc5	0.247
1416517_at;:	patatin-like p	Pnpla6	0.2475
1417053_at;:	prohibitin ///	Phb /// LOC6	0.2477
1416290_a_a	proteasome (	Psmc4	0.2479
1426974_at;:	RIKEN cDNA 4	4632413K17F	0.2483
1439320_at;:	cDNA sequen	BC030477	0.2489
1442376_at;:	actin-binding	Ablim1	0.2492
1418501_a_a	oxidation resi	Oxr1	0.2514
1435030_at;:	UPF2 regulato	Upf2	0.252
1457181_at;:	DNA segment	D15Wsu75e	0.2546
1434249_s_a	---	---	0.2551
1416280_at;:	RIKEN cDNA A	A730098D12F	0.2566
1426827_at;:	ubiquitin-like	Uble1b	0.2566
1434394_at;:	Bcl3 binding p	B3bp	0.2573
1425199_a_a	erythrocyte p	Epb4.1l5	0.2577
1419937_at	cDNA sequen	BC008163	0.2583
1425327_at;:	expressed se	AA536749	0.2583
1444327_at;:	E3 ubiquitin p	Edd1	0.2583
1426812_a_a	RIKEN cDNA 9	9130404D14F	0.259
1416240_at	proteasome (	Psmb7	0.2599
1434215_at;:	RIKEN cDNA I	B230308N11F	0.2603
1427134_at;:	splicing facto	Sfrs12	0.2622
1445857_at;:	cDNA sequen	BC026585	0.2632
1416851_at;:	Suppression c	St13	0.2643
1424050_s_a	fibroblast gro	Fgfr1	0.2664
1435442_at	WD repeats a	Wdsof1	0.2664
1434348_at	fasciculation a	Fez2	0.2682
1416440_at;:	BCL2-associa	Bclaf1	0.2688
1428844_a_a	CD164 antige	Cd164	0.2688
1423707_at	RIKEN cDNA 3	3830406C13F	0.2695
1434550_at;:	transmembra	Tmem50b	0.2695
1431777_a_a	high mobility	Hmgn3	0.2702
1415758_at;:	furry homolo	Fryl	0.2723
1433577_at;:	RIKEN cDNA A	A730017C20F	0.274
1416343_a_a	lysosomal me	Lamp2	0.2766

1415728_at;	poly(A) bindi	Pabpn1	0.2791
1445597_s_a	HRAS like su	Hrasls3	0.283
1428192_at;	kelch repeat	Kbtbd7	0.2844
1421241_at;	nerve growth	Ngfr	0.2847
1416064_a_a	heat shock 7(	Hspa5	0.2851
1430449_at;	RIKEN cDNA (	C330002I19R	0.2868
1419251_at;	epidermal gr	Eps15	0.2872
1428252_at;	chromatin m	Chmp2b	0.2872
1426369_at;	male sterility	Mlstd2	0.2882
1418718_at;	chemokine (C	Cxcl16	0.2883
1417115_at;	mitogen activ	Map3k12	0.2885
1428368_at;	Rho GTPase a	Arhgap21	0.2893
1429362_a_a	splicing facto	Sf3b2	0.2914
1454754_a_a	angio-associa	Aamp	0.2928
1434002_at;	checkpoint su	Ches1	0.2932
1426258_at;	sortilin-relate	Sorl1	0.2936
1420620_a_a	dystroglycan	Dag1	0.295
1423872_a_a	ring finger pr	Rnf13	0.295
1429454_at;	GTPase activa	Gapvd1	0.295
1441689_at;	nucleoporin 1	Nup153	0.2958
1415979_x_a	ribosomal pro	Rpl7	0.2963
1428221_at	kelch domain	Klhdc8b	0.2978
1425068_a_a	Component o	Cog7	0.2999
1443486_at	testis express	Tex264	0.2999
1422073_a_a	cadherin EGF	Celsr2	0.301
1423374_at	nuclear recep	Ncoa6	0.3019
1422971_at;	GCN5 genera	Gcn5l2	0.3038
1426713_s_a	glutamyl-prol	Eprs /// LOC6	0.3042
1434892_x_a	retinoblastom	Rbbp4	0.3043
1436810_x_a	RIKEN cDNA 2	2900010M23I	0.3043
1421381_a_a	procollagen, t	Col9a1	0.3046
1424039_at	RIKEN cDNA 2	2700060E02F	0.3055
1435413_x_a	transmembra	Tmem66	0.3055
1448752_at	carbonic anh	Car2	0.3072
1420612_s_a	protein tyrosi	Ptp4a2	0.3112
1424204_at;	mitochondrial	Mrpl13	0.3124

1434633_at;	CREB binding	Crebbp	0.3139
1417083_at	Sec61 beta su	Sec61b	0.314
1415716_a_a	ribosomal pro	Rps27	0.3142
1418174_at;	D site albumi	Dbp	0.3143
1450623_at	guanine nucle	Gnb2	0.3181
1420020_at;	RIKEN cDNA 2	2310004N24F	0.3184
1427895_at	suppressor of	Suz12	0.3184
1426968_a_a	retinol dehydı	Rdh10	0.3189
1416762_at;	S100 calcium	S100a10	0.3232
1422555_s_a	guanine nucle	Gna13	0.3235
1422064_a_a	yippee-like 3	Ypel3	0.3277
1426624_a_a	zinc finger an	Zbtb20	0.3277
1434337_at	Transcribed lo	---	0.3282
1434676_at;	myotubularin	Mtmr9	0.3284
1422429_at;	ring finger pr	Rnf14	0.3301
1452625_at	potassium ch	Kctd2	0.3301
1416829_at	ATP synthase	Atp5b	0.3304
1416951_a_a	ATPase, H+ tı	Atp6v1d	0.3308
1452239_at	gene trap RO	Gt(ROSA)26S	0.3313
1428579_at	---	---	0.3317
1431235_at;	formin-like 2	Fmnl2	0.3317
1435986_x_a	succinate deh	Sdhc	0.3318
1445024_at;	START domai	Stard7	0.3338
1452196_a_a	NCK-associat	Nckap1	0.3348
1426384_a_a	tyrosine 3-mo	Ywhae	0.3384
1429298_at;	dimethylargin	Ddah1	0.341
1423047_at;	toll interactin	Tollip	0.3447
1428125_at;	RIKEN cDNA 4	4921506J03R	0.349
1424175_at;	thyrotroph en	Tef	0.35
1417274_at	small nuclear	Snrpa	0.3501
1436316_at	RIKEN cDNA 9	9430029L20R	0.3517
1454668_at	zinc finger, U	Zubr1	0.3544
1443339_at;	RIKEN cDNA 2	2310056P07F	0.3547
1423972_at	electron trans	Etfa	0.3552
1434931_at;	neogenin	Neo1	0.3553
1419653_a_a	DEAD (Asp-G	Ddx5	0.3576

1452580_a_a	mitochondrial	Mrpl21	0.3579
1428272_at	eukaryotic tra	Eif1b	0.359
1419548_at;:	LY6/PLAUR do	Lypd6	0.3635
1454969_at;:	karyopherin (	Kpna1	0.3635
1448589_at	NADH dehydr	Ndufb5	0.3642
1430716_at;:	selenoprotein	Sepp1	0.3643
1452141_a_a	RIKEN cDNA !	5330421F07F	0.3643
1416842_at	glutathione S	Gstm5	0.3673
1418822_a_a	Notch gene h	Notch2	0.3679
1451889_at;:	ADP-ribosylat	Arf6	0.3679
1425282_at;:	IBR domain c	Ibrdc2	0.368
1423974_at;:	nuclear mitot	Numa1	0.373
1416614_at;:	aspartate-bet	Asph	0.3748
1420959_at;:	CREBBP/EP30	Cri1	0.3748
1420485_at;:	nucleolar prot	Nol7	0.375
1416791_a_a	nuclear RNA e	Nxf1	0.3755
1423384_s_a	testis express	Tex261	0.3766
1423032_at;:	ribosomal pro	Rpl39	0.3782
1426448_at;:	praja1, RING-	Pja1	0.38
1416219_at	ribosomal pro	Rpl19	0.384
1440151_s_a	endothelial di	Edf1	0.3862
1419039_at;:	cytochrome P	Cyp2d22	0.387
1416417_a_a	NADH dehydr	Ndufb7	0.3872
1417365_a_a	calmodulin 1	Calm1	0.3873
1424085_at	NADH dehydr	Ndufa4	0.3882
1415766_at;:	protein kinase	Prkcn	0.3889
1419999_at;:	SEC22 vesicle	Sec22b	0.3889
1420567_at;:	immunoglobu	Igbp1	0.3889
1417594_at	G kinase anch	Gkap1	0.3905
1436192_at;:	ADP-ribosylat	Arfgef2 /// LC	0.3942
1427177_at;:	FYVE and coil	Fyco1	0.3943
1422432_at;:	diazepam bin	Dbi	0.395
1417963_at;:	phospholipid	Pltp	0.3956
1451172_at	RIKEN cDNA	1200015A19F	0.3973
1416577_a_a	ring-box 1	Rbx1	0.3984
1436689_a_a	aldehyde deh	Aldh9a1	0.4

1417326_a_a	anaphase pro	Anapc11	0.4002
1421152_a_a	guanine nucle	Gnaq	0.4002
1428938_at;	guanine nucle	Gnao1	0.4002
1454161_s_a	RIKEN cDNA (	0610007P14F	0.4007
1425326_at;	ATP citrate ly	Acly	0.4024
1421116_a_a	reticulon 4	Rtn4	0.4025
1428707_at;	parathymosin	Ptms	0.4027
1418115_s_a	RIKEN cDNA :	1110004F10F	0.4032
1423335_at	interferon alp	Ifrg15	0.4032
1451310_a_a	cathepsin L	Ctsl	0.4036
1423898_a_a	thyroid hormo	Trip12	0.4037
AFFX-GapdhM	similar to gly	LOC14433	0.4041
1415867_at;	myristoylated	Marcks	0.4055
1415971_at;	chaperonin su	Cct4	0.4055
1415736_at;	prefoldin 5	Pfdn5	0.4065
1423272_at;	ribosomal pro	Rpl27a	0.4065
1437729_at	polymerase (I	Polg	0.4065
1415752_at	cDNA sequen	BC031181	0.4127
1418341_at;	RAB4A, meml	Rab4a	0.4171
1416102_at;	RAB5C, meml	Rab5c	0.4178
1424684_at	tyrosine 3-mo	Ywhaz	0.4178
1416416_x_a	glutathione S	Gstm1	0.4179
1420945_at;	tankyrase, TR	Tnks2	0.4179
1428388_at;	alpha thalass	Atrx	0.4179
1424308_at	solute carrier	Slc24a3	0.4185
1422692_at;	SUB1 homolo	Sub1	0.4189
1455955_s_a	sorting nexin	Snx17	0.4194
1425621_at;	tripartite mot	Trim35	0.4239
1428381_a_a	RIKEN cDNA 2	2700038C09F	0.4239
1422009_at;	ATPase, Na+/	Atp1b2	0.4242
1415911_at;	imprinted anc	Impact	0.4244
1423330_at;	endosulfine a	Ensa	0.4252
1424415_s_a	spondin 1, (f-	Spon1	0.4263
1417483_at;	nuclear factor	Nfkbiz	0.4284
1436959_x_a	nasal embryo	Nelf	0.4289
1416565_at;	cytochrome c	Cox6b1	0.4305

1421102_a_a	vesicle-associ	Vamp3	0.4323
1440455_at;	expressed se	AI848599	0.4336
1426722_at;	solute carrier	Slc38a2	0.4363
1429798_s_a	CKLF-like MA	Cmtm5	0.4368
1428631_a_a	ubiquinol cyt	Uqcrc2	0.4372
1416810_at;	male enhance	Mea1	0.4376
1444028_s_a	dedicator of c	Dock9 /// LO	0.4378
1448447_at	vacuolar prot	Vps28	0.4379
1417766_at;	RIKEN cDNA 2	2400003N08	0.4397
1427896_at;	cytochrome b	Cyb5b	0.4397
1436221_at;	DNA segment	D1Ertd471e	0.4403
1448346_at;	cofilin 1, non-	Cfl1	0.4421
1424886_at;	protein tyrosi	Ptprd	0.443
1416066_at	CD9 antigen	Cd9	0.4439
1416394_at	Bcl2-associat	Bag1	0.4445
1428145_at;	acetyl-Coenzy	Acaa2	0.4451
1424707_at;	transmembra	Tmed10	0.4503
1426419_at;	RNA binding r	Rbm26	0.4504
1428407_at	heterogeneou	Hnrpa0	0.4505
1423007_a_a	glial cell line	Gfra2	0.4525
1423689_a_a	CD44 antigen	Cd44	0.4525
1423760_at;	G-protein sig	Gpsm1	0.4525
1451971_at	cullin 4A	Cul4a	0.4525
1426139_a_a	chemokine (C	Ccrl1	0.4527
1433593_at;	yippee-like 5	Ypel5	0.4599
1426788_a_a	structure spe	Ssrp1	0.4624
1424049_at	leucine rich r	Lrrc42	0.4644
1418529_at;	O-sialoglycop	Osgep	0.465
1416546_a_a	G protein-cou	Gpr37	0.4661
1450875_at	ribosomal pro	Rpl6	0.4661
1416300_a_a	solute carrier	Slc25a3	0.4676
1425097_a_a	zinc finger pr	Zfp106	0.4683
1416032_at;	transmembra	Tmem109	0.4687
1418576_at;	Yip1 domain f	Yipf5	0.4728
1416508_at;	mediator of R	Med28	0.4735
1418348_a_a	activating sig	Ascc2	0.4735

1423798_a_a	eukaryotic tra	Eif1	0.474
1415998_at;	voltage-depe	Vdac1	0.4754
1417965_at;	pleckstrin hor	Plekha1	0.4764
1415970_at;	cytochrome c	Cox6c	0.4789
1426565_at;	insulin-like gr	Igf1r	0.4794
1416009_at	tetraspanin 3	Tspan3	0.4816
1423686_a_a	RIKEN cDNA	3300001P08F	0.4828
1424802_a_a	proline rich 1	Prr13	0.4828
1423362_at;	sortilin 1	Sort1	0.4853
1419933_at;	p53 and DNA	Pdrg1	0.4874
1423433_at;	TROVE domai	Trove2	0.4874
1425183_a_a	ribosomal pro	Rpl4	0.4876
1419428_a_a	glucosidase, a	Gaa	0.4882
1426758_s_a	GTL2, imprint	Gtl2	0.4882
1420093_s_a	heterogeneou	Hnrpdl	0.4886
1430978_at;	ribosomal pro	Rps25	0.4895
1415863_at;	eukaryotic tra	Eif4g2	0.4903
1425364_a_a	solute carrier	Slc3a2	0.4945
1456675_at;	expressed se	AA536749	0.4963
1416467_at;	DEAD/H (Asp	Ddx3x	0.4969
1421945_a_a	brix domain c	Bxdc1	0.4972
1438118_x_a	vimentin	Vim	0.4981
1416901_at;	Niemann Pick	Npc2	0.5009
1415881_at;	metaxin 1	Mtx1	0.501
1418521_a_a	growth hormo	Ghitm	0.501
1417440_at;	AT rich intera	Arid1a /// LO	0.5019
1421814_at;	moesin	Msn	0.5025
1417710_at;	methyltransfe	Mettl9	0.5026
1428739_at;	RIKEN cDNA	2310040A07F	0.5051
1431081_a_a	phospholipid	Plscr3	0.5051
1415676_a_a	proteasome (	Psmb5	0.5056
1435203_at;	mannosidase	Man2a2	0.5069
1420575_at	metallothione	Mt3	0.5088
1453568_at	RIKEN cDNA	2310032F03F	0.5088
1455316_x_a	Solute carrier	Slc15a2 /// K	0.5089
1437773_x_a	DEAD (Asp-G	Ddx17	0.5099

1416478_a_a	malate dehyd	Mdh2	0.5121
1416669_s_a	nascent polyp	Naca	0.5127
1428332_at;	RIKEN cDNA	1500004A08F	0.5128
1435897_at	ribosomal pro	Rpl32	0.5161
1424017_a_a	histidine triad	Hint1	0.5165
1416364_at;	heat shock pr	Hsp90ab1	0.5181
1416514_a_a	fascin homolo	Fscn1	0.5193
1415922_s_a	MARCKS-like	Marcksl1	0.5197
1421196_at;	protein tyrosi	Ptpn11	0.5205
1422490_at;	BCL2/adenovi	Bnip2	0.5205
1423251_at;	LUC7-like 2 (S	Luc7l2	0.5206
1415797_at;	discoidin dom	Ddr1	0.5227
1421557_x_a	thioredoxin 2	Txn2	0.5238
1425791_at;	paraoxonase	Pon2	0.5246
1416906_at;	anaphase-pro	Anapc5	0.5251
1421081_a_a	barrier to aut	Banf1	0.526
1427347_s_a	tubulin, beta	Tubb2a /// Tu	0.5262
1418497_at;	fibroblast gro	Fgf13	0.5267
1428512_at	basic helix-lo	Bhlhb9	0.5267
1420293_at;	plexin C1	Plxnc1	0.5299
1423213_at;	acyltransferas	Aytl2	0.5299
1425941_a_a	protein-L-isoa	Pcmtd1	0.5299
1429429_s_a	endothelial di	Edg3	0.5299
1437173_at;	cDNA sequen	BC025462	0.5299
1417612_at;	immediate ea	Ier5	0.5313
1415724_a_a	cell division c	Cdc42	0.536
1422605_at	protein phosp	Ppp1r1a	0.5391
1416511_a_a	CDC42 effecto	Cdc42ep4	0.5435
1423818_a_a	ADP-ribosylat	Arl6ip1	0.5438
1436292_a_a	ornithine dec	Oaz1	0.5456
1423676_at;	ATP synthase	Atp5h	0.5461
1415675_at	dolichol-phos	Dpm2	0.5473
1416065_a_a	ankyrin repea	Ankrd10	0.5499
1416285_at;	NADH dehydr	Ndufc1	0.5503
1433679_at	far upstream	Fubp3	0.5524
1436315_at;	MYST histone	Myst3	0.5524

1415849_s_a	stathmin 1	Stmn1	0.5534
1424440_at;	mitochondrial	Mrps6	0.5534
1437005_a_a	ribosomal pro	Rpl18	0.5554
1451974_at	oxysterol bin	Osbpl2	0.5576
1419489_at;	expressed se	AW049604	0.5626
1426599_a_a	solute carrier	Slc2a1	0.5626
1424053_a_a	transcription	Tcf25	0.563
1448822_at	proteasome (	Psmb6	0.5638
1436995_a_a	gene model 1	Gm106	0.5696
1456335_at;	ribosomal pro	Rpl26	0.5696
1434987_at;	aldehyde deh	Aldh2	0.5709
1427131_s_a	leucine rich re	Lrrc58	0.5712
1420887_a_a	asparaginyl-t	Nars	0.575
1428666_at;	Bcl2-like 1	Bcl2l1	0.575
1431018_at;	RIKEN cDNA	1810013L24R	0.5773
1415760_s_a	ATX1 (antioxi	Atox1	0.5786
1416630_at	inhibitor of DI	Id3	0.5788
1417864_at;	phosphoglyce	Pgk1 /// LOC4	0.5788
1455006_at	RIKEN cDNA 2	2310016M24I	0.585
1427100_at;	meteorin, glia	Metrn	0.5859
1415838_at;	peroxiredoxin	Prdx1	0.5877
1416000_a_a	signal-induce	Sipa1l1	0.5877
1434222_at;	serine incorpo	Serinc1	0.5877
1427269_at;	splicing facto	Sfrs11	0.5888
1420037_at;	ATP synthase	Atp5a1	0.5916
1416148_at;	lysosomal-ass	Laptm4b	0.5923
1434198_at;	ATPase inhibi	Atpif1	0.5923
1415908_at;	testis-specific	Tspyl1	0.5929
1416937_at	gamma-amin	Gabarap	0.5935
1448388_a_a	RIKEN cDNA	1110002B05F	0.5977
1433504_at	brain glycoge	Pygb	0.598
1449622_s_a	ATPase, H+ t	Atp6ap1	0.5997
1421239_at;	interleukin 6	Il6st	0.6016
1419497_at;	cyclin-depenc	Cdkn1b	0.609
1427103_at	pleckstrin hor	Plekhq1	0.6142
1422521_at;	dynactin 1	Dctn1	0.6145

1420841_at;	protein tyrosi	Ptprf	0.6148
1423493_a_a	nuclear factor	Nfix	0.6152
1420814_at;	guanosine dip	Gdi2	0.6164
1415819_a_a	protein phosp	Ppp2r1a	0.6166
1417502_at;	tetraspanin 7	Tspan7	0.6178
1426393_a_a	stromal cell d	Sdf4	0.6238
1418885_a_a	isocitrate deh	Idh3b	0.6247
1424115_at;	protein phosp	Ppp5c	0.6272
1423567_a_a	proteasome (	Psma7	0.6297
1423188_a_a	RIKEN cDNA (	6720456B07F	0.6316
1418101_a_a	reticulon 3	Rtn3	0.6374
1426484_at;	UBX domain (	Ubxd2	0.6393
1423233_at;	CCAAT/enhan	Cebpd	0.6399
1419223_a_a	dynamin 1-lik	Dnm1l	0.6427
1428008_at;	dystrobrevin a	Dtna	0.6427
1418840_at	eukaryotic tra	Eif3s12	0.6428
1452674_a_a	programmed	Pdcd4 /// LOC	0.6428
1424544_at	nuclear recep	Nrbp2	0.6448
1431328_at	protein phosp	Ppp1cb	0.6456
1433480_at	RIKEN cDNA 2	2900010J23R	0.6462
1416788_a_a	isocitrate deh	Idh3g	0.6466
1416307_at;	adaptor-relat	Ap1m1	0.6471
1430713_s_a	NADH dehydr	Ndufa13	0.649
1417734_at	A kinase (PRK	Akap8l	0.6493
1433781_a_a	claudin 12	Cldn12	0.6502
1437195_x_a	mitogen activ	Mapk10	0.6502
1415771_at;	nucleolin	Ncl	0.6525
1426794_at;	protein tyrosi	Ptprs	0.6527
1423909_at;	RIKEN cDNA (	0610011I04R	0.6533
1421928_at;	Eph receptor	Epha4	0.6534
1417312_at;	dickkopf hom	Dkk3	0.6549
1427021_s_a	ferritin heavy	Fth1	0.6552
1428025_s_a	phosphatidyli	Pitpnc1	0.6604
1436586_x_a	ribosomal pro	Rps14	0.6628
1415918_a_a	triosephospha	Tpi1	0.6636
1432635_a_a	translocation	Tloc1	0.6645

1416321_s_a	proline argini	Prelp	0.6646
1435496_at;	RIKEN cDNA !	5730469M10I	0.6677
1416925_at;	karyopherin (	Kpnb1	0.6699
1426195_a_a	cystatin C	Cst3	0.6714
1418119_at;	RNA binding r	Rbm8a	0.6716
1416452_at	ornithine ami	Oat	0.6728
1426400_a_a	calpain, small	Capns1	0.6746
1417423_at;	glutamate rec	Grina	0.6751
1452102_at;	coatomer pro	Copb2	0.6844
1428768_at;	PAN3 polyA s	Pan3	0.6849
1417753_at;	polycystic kid	Pkd2	0.6885
1421991_a_a	insulin-like gr	Igfbp4	0.6885
1454652_at	zinc finger, R	Zranb2	0.6889
1417311_at	cysteine rich	Crip2	0.691
1421938_at;	serine hydrola	Serhl	0.6939
1421313_s_a	cortactin	Cttn	0.6942
1416183_a_a	lactate dehyd	Ldhb	0.6969
1415678_at;	protein phosp	Ppm1a	0.7039
1427880_at;	ubiquinol-cyt	Uqcrq	0.7055
1416470_a_a	ribophorin I	Rpn1	0.7066
1416089_at	ribosomal pro	Rps15	0.7075
1452133_at;	ubiquinol-cyt	Uqcrh	0.7085
1433723_s_a	small EDRK-r	Serf2	0.7088
1417376_a_a	immunoglobu	Igsf4a	0.7099
1416035_at;	branched cha	Bcat1	0.7116
1430111_a_a	---	---	0.7116
1430539_at;	RIKEN cDNA	1810057P16F	0.7119
1422945_a_a	kinesin family	Kif5c	0.719
1415751_at;	cytochrome c	Cox7a2	0.7209
1416970_a_a	Ribosomal pr	Rpl18a	0.7209
1428152_a_a	heterochroma	Hp1bp3	0.7209
1421216_a_a	transmembra	Tmem58	0.721
1438426_at	iduronate 2-s	Ids	0.721
1417258_at	chaperonin su	Cct5	0.7215
1415820_x_a	non-POU-dom	Nono	0.7237
1424736_at	eukaryotic tra	Eef2	0.7249

1420129_s_a	DNA segment	D10Wsu52e	0.7321
1423896_a_a	ring finger pr	Rnf187	0.7321
1454916_s_a	ADP-ribosylat	Arfip1	0.7321
1448729_a_a	septin 4	40426	0.7324
1452285_a_a	eukaryotic tra	Eif3s5	0.7328
1435745_at;	RIKEN cDNA !	5031439G07F	0.7344
1456628_x_a	ribosomal pro	Rps24	0.7344
1435003_at;	phosphatidyli	Pik4ca	0.7347
1423833_a_a	brain protein	Brp44	0.7368
1425461_at;	F-box and WI	Fbxw11	0.7386
1435304_at;	superoxide di	Sod1	0.7425
1447810_x_a	pleckstrin hor	Plekhb2	0.7447
1416883_at;	quininoid dih	Qdpr	0.7531
1423664_at;	cleft lip and p	Clptm1	0.7531
1417027_at;	tripartite mot	Trim2	0.7565
1422128_at;	ribosomal pro	Rpl14	0.7662
1423069_at	activity-depe	Adnp	0.7678
1429951_at;	single-strand	Ssbp2	0.7681
1420622_a_a	heat shock pr	Hspa8 /// LO	0.7685
1419441_at	acidic riboson	Arbp	0.7712
1423906_at;	heat shock fa	Hsbp1	0.7726
1429452_x_a	RIKEN cDNA 4	4933439C20F	0.7727
1418234_s_a	breast carcino	Bcas2	0.7789
1428258_at	RIKEN cDNA 2	2010107E04F	0.7809
1416050_a_a	scavenger rec	Scarb1	0.7816
1425014_at;	nuclear recep	Nr2c2	0.7836
1416877_a_a	mitochondrial	Mrpl51	0.786
1422976_x_a	NADH dehydr	Ndufa7	0.7884
1415907_at;	cyclin D3	Ccnd3	0.7899
1416276_a_a	ribosomal pro	Rps4x	0.7913
1416858_a_a	FK506 bindin	Fkbp3	0.7947
1418567_a_a	signal recogn	Srp14	0.796
1418365_at;	cathepsin H	Ctsh	0.7967
1418561_at;	splicing factor	Sf3b1	0.7977
1427853_a_a	heat shock pr	Hspb1	0.7996
1415755_a_a	plastin 3 (T-is	Pls3	0.802

1415928_a_a	microtubule-a	Map1lc3b	0.802
1416383_a_a	pyruvate carb	Pcx	0.802
1423725_at	ubiquitin-con	Ube2v1 /// LC	0.802
1423759_a_a	phosphomanr	Pmm1	0.8045
1424167_a_a	transmembra	Tmco1	0.8045
1420924_at;	tissue inhibitc	Timp2	0.8068
1418002_at	HIG1 domain	Higd2a	0.8087
1427832_at;	tubulin alpha,	Tuba-rs1	0.8114
1421772_a_a	cytochrome c	Cox7a2l	0.8151
1427913_at;	RWD domain	Rwdd1	0.8159
1423653_at;	ATPase, Na+/	Atp1a1	0.8186
1431340_a_a	RIKEN cDNA 2	2310002J21R	0.8198
1416468_at	aldehyde deh	Aldh1a1	0.82
1416332_at	cold inducible	Cirbp	0.8287
1455222_a_a	upstream bin	Ubp1	0.8327
1418188_a_a	Metastasis as	Malat1	0.8339
1422451_at	mitochondrial	Mrps21	0.8354
1415872_at;	heterogeneou	Hnrph1	0.8371
1440265_at;	Jun proto-onc	Jund1	0.8385
1416406_at;	phosphoprote	Pea15	0.8415
1424917_a_a	WD repeat do	Wipi1	0.8418
1416054_at	ribosomal pro	Rps5	0.8423
1415888_at;	hepatoma-de	Hdgf	0.8454
1434274_at;	pam, highwir	Phr1	0.8476
1431997_at	RIKEN cDNA 3	3000002C10F	0.8541
1416001_a_a	coactosin-like	Cotl1	0.8555
1417600_at;	solute carrier	Slc15a2	0.856
1428257_s_a	dynein light c	Dynlrb1	0.8597
1451168_a_a	Rho GDP diss	Arhgdia	0.8598
1424714_at;	aldolase 3, C	Aldoc	0.8705
1434914_at;	RAB6B, mem	Rab6b	0.8711
1427173_a_a	mitochondrial	Mrps33	0.8726
1421110_at;	Metadherin	Mtdh	0.8747
1451294_s_a	small nuclear	Snrpe	0.8773
1416150_a_a	splicing factor	Sfrs3	0.8778
1437027_x_a	ribonucleic ac	Rnps1	0.8808

1416921_x_a	aldolase 1, A	Aldoa	0.8844
1416415_a_a	H2A histone f	H2afz /// LOC	0.8882
1422579_at;	heat shock pr	Hspe1 /// LOC	0.8933
1426742_at;	ATP synthase	Atp5f1	0.8998
1450149_a_a	protein phosp	Ppp1cc /// LC	0.906
1417432_a_a	guanine nucle	Gnb1	0.9067
1427183_at	epidermal grc	Efemp1	0.9097
1454813_at	coiled-coil do	Ccdc72	0.9131
1438633_x_a	LIM and SH3	Lasp1	0.9226
1423716_s_a	ATP synthase	Atp5d	0.9248
1427058_at;	eukaryotic tra	Eif4a1	0.9263
1424625_a_a	DENN/MADD	Dennd1a	0.928
1424874_a_a	polypyrimidin	Ptbp1	0.9332
1415759_a_a	hepatitis B vi	Hbxip	0.9343
1417608_a_a	ribosomal pro	Rpl13a	0.935
1426710_at;	phosphohistic	Phpt1	0.935
1427903_at	calmodulin 3	Calm3	0.935
1422031_a_a	zinc finger, A	Zfand6	0.9375
1420951_a_a	Son cell prolif	Son	0.9401
1416269_at;	ATP synthase	Atp5j2	0.9431
1423254_x_a	ribosomal pro	Rps27l	0.9438
1420038_at;	VATPase, H+	Atp6v1e1	0.9501
1416849_at;	zinc finger, C	Zcd1	0.9516
1429209_at;	procollagen, t	Col23a1	0.9553
1418209_a_a	profilin 2	Pfn2	0.9575
1424038_a_a	RIKEN cDNA 2	2310044H10F	0.9632
1456373_x_a	ribosomal pro	Rps20	0.9652
1416521_at;	selenoprotein	Sepw1	0.9734
1417417_a_a	cytochrome c	Cox6a1	0.9734
1428608_at	myosin light c	Mylc2b	0.9734
1436803_a_a	NADH dehydr	Ndufb9	0.9734
1437510_x_a	ribosomal pro	Rps17	0.9787
1417490_at;	cathepsin B	Ctsb	0.9818
1421374_a_a	FXYD domain	Fxyd1	0.9846
1416522_a_a	gene rich clus	Grcc10	0.9902
1425591_a_a	chromatin mc	Chmp2a	0.9976

1415869_a_a	S100 calcium	S100a16	1
1415950_a_a	forkhead box	Foxo1	1
1415991_a_a	DNA segment	D12Ertd553e	1
1416096_at;:	crystallin, alp	Cryab	1
1416239_at;:	paired box ge	Pax6	1
1416455_a_a	solute carrier	Slc25a36	1
1416526_a_a	RIKEN cDNA (	6330577E15F	1
1416769_s_a	phosphatidyle	Pebp1	1
1416834_x_a	transmembra	Tmepai	1
1416981_at;:	centrin 3	Cetn3	1
1417087_at;:	serine/arginir	Srpk2	1
1417125_at	receptor acce	Reep5	1
1417134_at;:	golgi apparatı	Glg1	1
1417239_at	NADH dehydr	Ndufb2	1
1417285_a_a	ribosomal pro	Rps21	1
1418626_a_a	translocase o	Timm17b	1
1419271_at;:	ataxin 10	Atxn10	1
1419398_a_a	argininosucci	Ass1	1
1419656_at;:	ATP synthase	Atp5k	1
1419665_a_a	host cell fact	Hcfc1r1	1
1422525_at;:	single-strand	Ssbp3	1
1422576_at;:	clusterin	Clu	1
1422623_x_a	basigin	Bsg	1
1422705_at;:	nuclear prote	Nupr1	1
1424000_a_a	ribosomal pro	Rpl37	1
1424313_a_a	NADH dehydr	Ndufs6	1
1424355_a_a	Ribosomal pr	Rpl41	1
1424364_a_a	ATPase, H+ t	Atp6v0b	1
1425194_a_a	transcriptiona	Sin3b	1
1425560_a_a	Parkinson dis	Park7	1
1425940_a_a	transcription	Tceb2	1
1428182_at	S-adenosylho	Ahcy	1
1428263_a_a	anaphase pro	Anapc13	1
1428405_at;:	RIKEN cDNA :	1110020P15F	1
1430288_x_a	kelch domain	Klhdc3	1
1431045_at;:	tripartite mot	Trim28	1

1433603_at	phosphoribos	Prpsap1	1
1438656_x_a	expressed se	AI413782	1
1448642_at	RIKEN cDNA	1500032L24R	1
1452584_at	ribosomal pro	Rps11	1
1452586_at	poly(rC) bindi	Pcbp1	1
1453729_a_a	NADH dehydr	Ndufs7	1
1456616_a_a	NADH dehydr	Ndufa5	1