

Tissue-conserved, beta cell-abundant transcripts

symbol	cluster	probe set	gene (Rat)	Accession	av FC	Beta score	probe set	Accession	av FC	Beta score	probe set	Accession	av FC	Beta score
AACS	B	1368126_at	acetoacetyl-CoA synthetase	NM_023104	2.1	4	218434_s_at	NM_023928	1.2	10	1451116_at	BC026817	4.837	17
ACBD3	C	1372552_at	acyl-Coenzyme A binding domain containing 3	AI102273	2.05	6	202324_s_at	NM_022735	2.28	14	1456316_a_at	BI965035	2.798	17
ACBD3	C	1372552_at	acyl-Coenzyme A binding domain containing 3	AI102273	2.05	6	202323_s_at	AI636775	1.7	14	1452137_at	BB704602	2.283	13
ACBD3	C	1372552_at	acyl-Coenzyme A binding domain containing 3	AI102273	2.05	6	202323_s_at	AI636775	1.7	14	1426635_at	BB704602	1.803	13
ACBD3	C	1372552_at	acyl-Coenzyme A binding domain containing 3	AI102273	2.05	6	202323_s_at	AI636775	1.7	14	1428668_at	BI790311	1.591	15
ACLY	C	1367854_at	ATP citrate lyase	NM_016987	2.91	2	201127_s_at	AI971281	2.43	14	1446315_at	BB261367	6.284	17
ACLY	C	1367854_at	ATP citrate lyase	NM_016987	2.91	2	201128_s_at	NM_001096	3.79	14	1438389_x_at	BB424434	2.938	17
ACLY	C	1367854_at	ATP citrate lyase	NM_016987	2.91	2	210337_s_at	U18197	4.94	14	1439445_x_at	AV347837	1.746	13
ACLY	C	1367854_at	ATP citrate lyase	NM_016987	2.91	2	201128_s_at	NM_001096	3.79	14	1451666_at	BI456232	2.52	17
ACLY	C	1367854_at	ATP citrate lyase	NM_016987	2.91	2	201128_s_at	NM_001096	3.79	14	1439459_x_at	BB276877	1.902	17
ADRA2A		1387708_at	adrenergic receptor, alpha 2a	U79031	2.64	6	209869_at	AF284095	1.22	8	1423022_at	NM_007417	2.889	17
AGA	C	1372421_at	aspartylglucosaminidase (predicted)	AA943766	2.72	6	204332_s_at	M64073	1.53	12	1434665_at	AV052058	3.899	17
AGA	C	1372421_at	aspartylglucosaminidase (predicted)	AA943766	2.72	6	204333_s_at	NM_000027	1.67	14	1434665_at	AV052058	3.899	17
AGA	C	1372421_at	aspartylglucosaminidase (predicted)	AA943766	2.72	6	216064_s_at	W27131	3.12	14	1434665_at	AV052058	3.899	17
AMPD2	C	1372782_a_at	similar to adenosine monophosphate deaminase 2 (isoform L)	BE111787	1.88	4	212360_at	AI916249	1.76	10	1438128_at	BB427907	8.87	17
AMPD2	C	1372782_a_at	similar to adenosine monophosphate deaminase 2 (isoform L)	BE111787	1.88	4	212360_at	AI916249	1.76	10	1438941_x_at	AV330806	2.884	15
AMPD2	C	1372782_a_at	similar to adenosine monophosphate deaminase 2 (isoform L)	BE111787	1.88	4	212360_at	AI916249	1.76	10	1426757_at	AK004759	2.263	17
AP3B2	A	1376844_at	adaptor-related protein complex 3, beta 2 subunit (predicted)	BI298352	1.44	2	205678_at	NM_004644	1.2	8	1451837_at	BC024722	2.044	17
AP3B2	A	1376844_at	adaptor-related protein complex 3, beta 2 subunit (predicted)	BI298352	1.44	2	205678_at	NM_004644	1.2	8	1426927_at	AB030202	1.991	15
AP3B2	A	1376844_at	adaptor-related protein complex 3, beta 2 subunit (predicted)	BI298352	1.44	2	205678_at	NM_004644	1.2	8	1445224_at	BI134892	1.858	13
AP3D1		1375444_at	adaptor-related protein complex 3, delta subunit (predicted)	BF409829	1.6	6	210974_s_at	AF130042	1.21	10	1417964_at	NM_007460	1.579	15
AP3D1		1367549_a_at	adaptor-related protein complex 3, delta subunit (predicted)	BI282757	1.37	0	208710_s_at	AI424923	1.2	8	1417964_at	NM_007460	1.579	15
APLP1	A	1371572_at	amyloid beta (A4) precursor protein	BM986220	1.35	-2	209462_at	U48437	6.71	8	1435857_s_at	AI848048	3.641	15
APLP1	A	1371571_at	amyloid beta (A4) precursor protein	BM986220	1.4	0	209462_at	U48437	6.71	8	1416134_at	NM_007467	1.909	15
ARCN1	C	1398896_at	archain	AA892567	1.61	6	201176_s_at	NM_001655	1.75	14	1423743_at	BC017124	2.415	15
ARCN1	C	1398896_at	archain	AA892567	1.61	6	201176_s_at	NM_001655	1.75	14	1451089_a_at	BC017124	1.806	17
ARCN1	C	1398896_at	archain	AA892567	1.61	6	201176_s_at	NM_001655	1.75	14	1436062_at	BB757123	1.763	15
ARF4	C	1398758_at	ADP-ribosylation factor 4	NM_024151	2.02	6	201096_s_at	AL537042	1.94	10	1434074_x_at	BB754834	3.116	17
ARF4	C	1398758_at	ADP-ribosylation factor 4	NM_024151	2.02	6	201097_s_at	NM_001660	1.69	12	1434074_x_at	BB754834	3.116	17
ARFIP1	C	1369026_at	ADP-ribosylation factor interacting protein 1	NM_021763	2.2	6	218230_at	AL044651	1.86	14	1454916_s_at	AV087417	1.733	15
ARFIP1	C	1369026_at	ADP-ribosylation factor interacting protein 1	NM_021763	2.2	6	214483_s_at	AF124489	1.5	12	1454916_s_at	AV087417	1.733	15
ARHGAP1	C	1393113_at	Rho GTPase activating protein 1 (predicted)	AA957578	1.88	4	202117_at	BG468434	1.51	10	1458118_at	AV233159	2.512	17
CD164	C	1386886_at	CD164 antigen	NM_031812	2.35	6	208654_s_at	BF669455	2.26	14	1416440_at	NM_016898	1.794	17
ARHGEF9		1377216_at	Cdc42 guanine nucleotide exchange factor (GEF) 9	BI295997	1.77	0	203263_s_at	AI625739	1.49	10	1436577_at	BB380620	3.924	17
ARHGEF9		1377216_at	Cdc42 guanine nucleotide exchange factor (GEF) 9	BI295997	1.77	0	203263_s_at	AI625739	1.49	10	1436258_at	BE651574	3.665	15
ARMCX2		1373156_at	armadillo repeat containing, X-linked 2	AI385330	4.91	4	203404_at	NM_014782	2.05	14	1456739_x_at	BB392869	3.011	15
ARMCX2		1373156_at	armadillo repeat containing, X-linked 2	AI385330	4.91	4	203404_at	NM_014782	2.05	14	1437849_x_at	BB443695	2.746	15
ARMCX2		1373156_at	armadillo repeat containing, X-linked 2	AI385330	4.91	4	203404_at	NM_014782	2.05	14	1415780_a_at	NM_026139	1.953	13
ARMCX3	A	1373958_at	armadillo repeat containing, X-linked 3 (similar to ALEX3 protein)	AA943149	1.82	6	217858_s_at	NM_016607	1.29	10	1424373_at	AK004598	3.482	17
ARMET	C	1372352_at	arginine-rich, mutated in early stage tumors (predicted)	AI170666	3.46	6	202655_at	NM_006010	1.89	14	1428112_at	AK014338	1.999	17
ASAH1	C	1369077_at	N-acylsphingosine amidohydrolase 1	NM_053407	2.61	6	210980_s_at	U47674	1.61	8	1416735_at	NM_019734	2.675	17
ASAH1	C	1374770_at	N-acylsphingosine amidohydrolase 1	AA800062	2.22	6	213702_x_at	AI934569	1.56	10	1416735_at	NM_019734	2.675	17
ATF2	C	1369472_a_at	activating transcription factor 2	NM_031018	1.98	6	212984_at	BE786164	2.13	12	1452116_s_at	BM119623	1.691	15
ATF3	B	1369268_at	activating transcription factor 3	NM_012912	7.46	4	202672_s_at	NM_001674	9.16	10	1449363_at	BC019946	11.24	17

ATF4	C	1367624_at	activating transcription factor 4	NM_024403	2.54	6	200779_at	NM_001675	1.27	8	1448135_at	M94087	1.798	15
ATP2A2	C	1369065_a_at	ATPase, Ca++ transporting, cardiac muscle, slow twitch 2	NM_017290	2.4	4	209186_at	M23114	1.71	8	1416551_at	NM_009722	2.461	13
ATP2A2	C	1370426_a_at	ATPase, Ca++ transporting, cardiac muscle, slow twitch 2	J04024	2.56	6	209186_at	M23114	1.71	8	1427250_at	AA245637	2.08	17
ATP2A3	C	1368477_at	ATPase, Ca++ transporting, ubiquitous	NM_012914	10.1	6	207522_s_at	NM_005173	5.44	14	1450124_a_at	NM_016745	2.63	13
ATP2A3	C	1368477_at	ATPase, Ca++ transporting, ubiquitous	NM_012914	10.1	6	213036_x_at	Y15724	3.91	14	1450124_a_at	NM_016745	2.63	13
ATP2A3	C	1368477_at	ATPase, Ca++ transporting, ubiquitous	NM_012914	10.1	6	207521_s_at	AF068220	2.88	14	1450124_a_at	NM_016745	2.63	13
ATP2A3	C	1371165_a_at	ATPase, Ca++ transporting, ubiquitous	AF458230	6.39	6	213042_s_at	AA877910	3.35	14	1450124_a_at	NM_016745	2.63	13
ATP6V1B2	A	1387664_at	ATPase, H+ transporting, V1 subunit B, isoform 2	NM_057213	2.46	2	201089_at	NM_001693	2.47	10	1446465_at	BB500880	1.77	17
ATP6V1B2	A	1371402_at	ATPase, H+ transporting, V1 subunit B, isoform 2	BM390600	2.16	2	201089_at	NM_001693	2.47	10	1446465_at	BB500880	1.77	17
ATP6V1C1	A	1374396_at	ATPase, H+ transporting, V1 subunit C, isoform 1 (predicted)	BI293306	2.02	4	202872_at	AW024925	1.77	10	1419546_at	AA987147	1.631	15
BACE1	A	1369256_at	beta-site APP cleaving enzyme	NM_019204	1.46	6	217904_s_at	NM_012104	2.21	10	1421825_at	AF200346	4.681	17
BACE1	A	1369256_at	beta-site APP cleaving enzyme	NM_019204	1.46	6	217904_s_at	NM_012104	2.21	10	1455826_a_at	BB114336	3.043	17
BEXL1	A	1373458_at	Brain expressed X-linked 4	BI289546	6.38	4	218332_at	NM_018476	7.43	12	1428209_at	AK010400	4.119	17
BEXL1	A	1373458_at	Brain expressed X-linked 4	BI289546	6.38	4	215440_s_at	AL523320	3.27	12	1428209_at	AK010400	4.119	17
BICD1	C	1373902_at	Bicaudal D homolog 1 (Drosophila) (predicted)	AI230054	1.83	6	214806_at	U90030	1.4	8	1437548_at	AV367839	6.374	17
BICD1	C	1373902_at	Bicaudal D homolog 1 (Drosophila) (predicted)	AI230054	1.83	6	214806_at	U90030	1.4	8	1438701_at	BB130665	2.79	15
BICD1	C	1373902_at	Bicaudal D homolog 1 (Drosophila) (predicted)	AI230054	1.83	6	214806_at	U90030	1.4	8	1442298_at	AI426427	2.117	17
BICD1	C	1373902_at	Bicaudal D homolog 1 (Drosophila) (predicted)	AI230054	1.83	6	214806_at	U90030	1.4	8	1451684_a_at	BC016192	1.803	15
BTG2	B	1386995_at	B-cell translocation gene 2, anti-proliferative	BI288701	4.46	4	201235_s_at	BG339064	4.38	14	1448272_at	NM_007570	2.977	13
BTG2	B	1386994_at	B-cell translocation gene 2, anti-proliferative	BI288701	2.22	4	201236_s_at	NM_006763	2.96	14	1416250_at	NM_007570	5.667	17
CACNA1A	A	1367821_a_at	calcium channel, voltage-dependent, P/Q type, alpha 1A subunit	BF400631	1.44	6	214933_at	AA769818	2.29	8	1459996_at	AI848293	3.153	17
CACNA1A	A	1367821_a_at	calcium channel, voltage-dependent, P/Q type, alpha 1A subunit	BF400631	1.44	6	214933_at	AA769818	2.29	8	1450510_a_at	AB066608	1.911	15
CACNA1D	A	1370640_a_at	calcium channel, voltage-dependent, L type, alpha 1D subunit	AF370009	1.58	6	210108_at	BE550599	5.53	14	1427974_s_at	BB048682	3.773	17
CACNA1D	A	1370640_a_at	calcium channel, voltage-dependent, L type, alpha 1D subunit	AF370009	1.58	6	207998_s_at	NM_000720	1.56	14	1427974_s_at	BB048682	3.773	17
CACNB2	A	1388023_a_at	calcium channel, voltage-dependent, beta 2 subunit	AF394941	1.45	4	213714_at	AI040163	1.7	8	1444693_at	BG069383	4.709	17
CACNB2	A	1370178_at	calcium channel, voltage-dependent, beta 2 subunit	BG373973	3.03	6	213714_at	AI040163	1.7	8	1456401_at	BB078175	2.511	13
CADM1		1370594_at	immunoglobulin superfamily, member 1	AF322217	1.35	4	207695_s_at	NM_001555	1.22	10	1447702_x_at	AV152162	5.647	15
CADM1		1370410_at	immunoglobulin superfamily, member 1	AF322216	1.74	2	207695_s_at	NM_001555	1.22	10	1438589_at	BB023061	1.902	17
CADPS	A	1368523_at	Ca<2+-dependent activator protein for secretion	NM_013219	4.97	2	204814_at	NM_003716	6.22	14	1445586_at	BB283411	6.856	17
CADPS	A	1368523_at	Ca<2+-dependent activator protein for secretion	NM_013219	4.97	2	204814_at	NM_003716	6.22	14	1448955_s_at	NM_012061	3.46	13
CADPS	A	1368523_at	Ca<2+-dependent activator protein for secretion	NM_013219	4.97	2	204814_at	NM_003716	6.22	14	1458298_at	AI462674	3.21	13
CAMKK2		1368753_at	calcium/calmodulin-dependent protein kinase kinase 2, beta	NM_031338	1.43	6	207359_at	NM_006549	1.65	14	1424476_at	BI157430	3.018	15
CANX	C	1371687_at	calnexin	AA848326	2.29	6	208853_s_at	L18887	2.51	14	1428935_at	AI988026	3.552	17
CANX	C	1371686_at	calnexin	AA848326	1.9	6	208852_s_at	AI761759	2.18	14	1428935_at	AI988026	3.552	17
CANX	C	1388442_at	calnexin	AA848326	2.19	6	200068_s_at	M94859	1.32	10	1428935_at	AI988026	3.552	17
CCDC6		1372134_at	coiled-coil-helix-coiled-coil-helix domain containing 6 (predicted)	AW918650	1.55	6	204716_at	NM_005436	1.62	12	1423779_at	BC011331	1.989	15
CCPG1	B	1375027_at	Cell cycle progression 1 (predicted)	AI407365	1.9	6	222156_x_at	AK022459	3.38	14	1424420_at	BC006717	2.791	17
CCPG1	B	1375027_at	Cell cycle progression 1 (predicted)	AI407365	1.9	6	221511_x_at	AB033080	2.15	14	1420127_s_at	C85100	3.918	17
CCPG1	B	1375027_at	Cell cycle progression 1 (predicted)	AI407365	1.9	6	221156_x_at	NM_004748	1.81	14	1420127_s_at	C85100	3.918	17
CD164		1386886_at	CD164 antigen	NM_031812	2.35	6	208405_s_at	NM_006016	1.62	10	1416440_at	NM_016898	1.794	17
CDK2AP1		1375326_at	CDK2 (cyclin-dependent kinase 2)-associated protein 1 (predicted)	AI602458	2.37	6	201938_at	NM_004642	1.51	10	1458451_at	BE985383	3.259	17
CHGA	B	1387235_at	chromogranin A	NM_021655	14.3	4	204697_s_at	NM_001275	18.2	14	1418149_at	NM_007693	25.11	17
CHM		1369227_at	choroideremia	NM_017067	2.4	6	207099_s_at	NM_000390	1.5	14	1448712_at	NM_018818	2.281	17
CHM		1373802_at	Choroideremia	BF392902	2.15	6	207099_s_at	NM_000390	1.5	14	1448712_at	NM_018818	2.281	17
CHST3		1369512_at	carbohydrate (chondroitin 6/keratan) sulfotransferase 3	NM_053408	1.51	6	208252_s_at	NM_004273	1.2	10	1460322_at	NM_016803	2.43	17
CKAP4		1375654_at	cytoskeleton-associated protein 4 (predicted)	BI278813	1.41	6	200999_s_at	NM_006825	1.85	8	1426755_at	BB312117	2.106	13
CLCN3	B	1392453_at	chloride channel 3	AI763523	1.74	0	201735_s_at	NM_001829	2.89	14	1416610_a_at	NM_007711	1.967	13
CLCN3	B	1380547_at	chloride channel 3	BI288519	1.82	0	201734_at	AI760629	2.43	14	1416610_a_at	NM_007711	1.967	13
CLCN3	B	1380547_at	chloride channel 3	BI288519	1.82	0	201732_s_at	AF029346	1.98	14	1416610_a_at	NM_007711	1.967	13
CLCN3	B	1380547_at	chloride channel 3	BI288519	1.82	0	201733_at	AA902971	1.77	10	1416610_a_at	NM_007711	1.967	13

CNOT2	1376067_at	CCR4-NOT transcription complex, subunit 2	AA800637	1.5	4	222181_at	BG105204	1.3	14	1447212_at	BB165759	2.197	17
COPA	1388377_at	coatamer protein complex subunit alpha (predicted)	AW142717	1.91	6	214336_s_at	AI621079	1.33	8	1437274_at	BM204808	3.998	17
COPA	1388377_at	coatamer protein complex subunit alpha (predicted)	AW142717	1.91	6	214336_s_at	AI621079	1.33	8	1415706_at	BQ175820	1.619	17
COPB2	C 1367454_at	coatamer protein complex, subunit beta 2 (beta prime)	NM_021765	1.88	6	201098_at	NM_004766	1.85	14	1456175_a_at	AV109541	1.69	17
COPG	1372464_at	coatamer protein complex, subunit gamma	BE109050	2.1	6	217749_at	NM_016128	1.4	12	1416017_at	BC024686	2.018	15
COPG	1371810_at	Coatamer protein complex, subunit gamma	BF284093	1.39	4	217749_at	NM_016128	1.4	12	1415670_at	BC024686	2.364	15
COX15	C 1377444_at	COX15 homolog, cytochrome c oxidase assembly protein (yeast)	BE120680	1.77	6	221550_at	BC002382	1.15	14	1426693_x_at	BC021498	2.671	17
COX15	C 1377444_at	COX15 homolog, cytochrome c oxidase assembly protein (yeast)	BE120680	1.77	6	221550_at	BC002382	1.15	14	1452146_a_at	BC021498	1.564	15
CPE	B 1386921_at	carboxypeptidase E	NM_013128	12.3	6	201116_s_at	AI922855	7.13	14	1415949_at	BC010197	3.248	15
CPE	B 1386921_at	carboxypeptidase E	NM_013128	12.3	6	201117_s_at	NM_001873	6.85	14	1415949_at	BC010197	3.248	15
CPEB1	1392996_at	cytoplasmic polyadenylation element binding protein 1 (predicted)	BG668435	2.18	6	219578_s_at	NM_030594	1.27	10	1417960_at	NM_007755	8.554	17
CPLX2	A 1368584_a_at	complexin 2	NM_053878	1.63	2	206368_at	NM_006650	2.59	14	1421477_at	NM_009946	3.797	15
CPLX2	A 1368584_a_at	complexin 2	NM_053878	1.63	2	206368_at	NM_006650	2.59	14	1455672_s_at	BE946238	2.065	15
CRMP1	A 1387899_at	collapsin response mediator protein 1	U52102	3.51	0	202517_at	NM_001313	1.81	8	1448289_at	AB006714	2.205	13
CRYZ	1374628_at	crystallin, zeta (predicted)	AI232098	1.38	4	202950_at	NM_001889	3.03	10	1430546_at	AK010433	1.905	15
CTNNA1	C 1371921_at	catenin (cadherin-associated protein), alpha 1, 102kDa	BM986272	1.88	6	200765_x_at	NM_001903	1.6	8	1443662_at	BE989574	3.473	17
CTNNA1	C 1371921_at	catenin (cadherin-associated protein), alpha 1, 102kDa	BM986272	1.88	6	200764_s_at	AI826881	1.54	8	1443662_at	BE989574	3.473	17
CYB561	1398383_at	cytochrome b-561 (predicted)	BI282164	1.5	4	209163_at	AL514271	2.34	14	1417507_at	BC006732	2.049	11
CYB561	1398383_at	cytochrome b-561 (predicted)	BI282164	1.5	4	209164_s_at	BC002976	1.23	12	1417507_at	BC006732	2.049	11
DAAM1	1389318_at	Dishevelled associated activator of morphogenesis 1 (predicted)	BF282986	1.36	4	216060_s_at	AK021890	1.97	12	1431035_at	AW988556	2.658	17
DACH1	B 1384907_at	similar to dachshund homolog 1 (Drosophila)	not on RG230A array			205471_s_at	AW772082	3.46	14	1433743_at	BG075820	3.216	17
DAD1	C 1367465_at	defender against cell death 1	AI013627	2.1	6	200046_at	NM_001344	1.61	14	1454860_x_at	BI966630	3.403	17
DAD1	C 1367465_at	defender against cell death 1	AI013627	2.1	6	200046_at	NM_001344	1.61	14	1418528_a_at	NM_010015	1.975	17
DCX	A 1374966_at	doublecortin	BE109057	12	6	204850_s_at	NM_000555	1.69	12	1418139_at	BB418548	29.54	17
DCX	A 1374966_at	doublecortin	BE109057	12	6	204850_s_at	NM_000555	1.69	12	1418141_at	BB418548	9.299	17
DCX	A 1374966_at	doublecortin	BE109057	12	6	204851_s_at	AF040254	2.46	12	1448974_at	BB418548	2.602	15
DDIT3	A 1369590_a_at	DNA-damage inducible transcript 3	NM_024134	9.57	6	209383_at	BC003637	1.95	14	1417516_at	NM_007837	3.35	17
DDIT3	A 1376178_at	DNA-damage inducible transcript 3	BF403703	7.01	4	209383_at	BC003637	1.95	14	1443897_at	BB200603	2.104	15
DDOST	C 1372247_at	dolichyl-di-phosphooligosaccharide-protein glycotransferase (predictec	BF282907	2.63	6	208674_x_at	BC002594	1.91	12	1416493_at	NM_007838	2.059	17
DDOST	C 1372247_at	dolichyl-di-phosphooligosaccharide-protein glycotransferase (predictec	BF282907	2.63	6	208675_s_at	D29643	1.84	14	1416493_at	NM_007838	2.059	17
DDR1	B 1370216_at	discoidin domain receptor family, member 1	L26525	1.76	2	210749_x_at	L11315	4.37	14	1459990_at	BF462661	7.157	17
DDR1	B 1370216_at	discoidin domain receptor family, member 1	L26525	1.76	2	207169_x_at	NM_001954	5.07	14	1439382_x_at	BB378700	2.187	15
DDR1	B 1370216_at	discoidin domain receptor family, member 1	L26525	1.76	2	208779_x_at	L20817	4.73	14	1439382_x_at	BB378700	2.187	15
DDX50	1390688_at	DEAD (Asp-Glu-Ala-Asp) box polypeptide 50 (predicted)	AA799576	1.55	2	221699_s_at	AF334103	1.71	12	1439082_at	AA518255	3.026	17
DDX50	1390688_at	DEAD (Asp-Glu-Ala-Asp) box polypeptide 50 (predicted)	AA799576	1.55	2	221699_s_at	AF334103	1.71	12	1417875_at	NM_053183	2.205	17
DDX6	1398945_at	DEAD (Asp-Glu-Ala-Asp/His) box polypeptide 6 (mapped)	AA851439	1.48	4	204909_at	NM_004397	1.27	14	1424598_at	BC021452	6.716	17
DDX6	1398945_at	DEAD (Asp-Glu-Ala-Asp/His) box polypeptide 6 (mapped)	AA851439	1.48	4	204909_at	NM_004397	1.27	14	1447789_x_at	BB150520	3.517	17
DLG5	1377121_at	Discs, large homolog 5 (Drosophila) (predicted)	AW520410	2.01	6	201681_s_at	AB011155	1.58	12	1431187_s_at	BF140264	2.872	13
DNAJB9	B 1387116_at	DnaJ (Hsp40) homolog, subfamily B, member 9	NM_012699	3.87	6	202843_at	NM_012328	3.68	14	1417191_at	NM_013760	1.903	13
DNAJB9	B 1387116_at	DnaJ (Hsp40) homolog, subfamily B, member 9	NM_012699	3.87	6	202842_s_at	AL080081	2.83	14	1417191_at	NM_013760	1.903	13
DNAJC1	1376079_at	DnaJ (Hsp40) homolog, subfamily C, member 1 (predicted)	BI295806	1.83	4	218409_s_at	NM_022365	1.59	12	1459791_at	AV129500	2.56	17
DNM1L	C 1369220_at	dynamamin 1-like	NM_053655	1.36	2	203105_s_at	NM_012062	1.4	8	1428008_at	BC027538	5.16	17
DNM1L	C 1388022_a_at	dynamamin 1-like	AF132727	1.43	2	203105_s_at	NM_012062	1.4	8	1428087_at	BM249101	1.591	15
DPH4	C 1390168_a_at	DPH4, JJJ3 homolog (S. cerevisiae) (zinc finger, CSL-type containing	BI289564	2.92	6	213853_at	AL050199	1.68	12	1451389_at	AY028460	10.74	17
DPH4	C 1390168_a_at	DPH4, JJJ3 homolog (S. cerevisiae) (zinc finger, CSL-type containing	BI289564	2.92	6	213853_at	AL050199	1.68	12	1430718_s_at	AV313469	7.305	17
DST	1375357_at	dystonia 1	BE112237	1.35	4	220154_at	NM_020388	1.36	10	1450119_at	NM_010081	2.144	13
DSTN	C 1371375_at	destrin	BI285599	2.79	6	201021_s_at	BF697964	1.58	10	1439535_at	BB204161	2.742	17
DSTN	C 1375881_at	destrin	AI170442	2.41	6	201021_s_at	BF697964	1.58	10	1439535_at	BB204161	2.742	17
DYNC2L1	C 1387782_at	dynein light chain-2	NM_080697	1.65	6	203762_s_at	NM_016008	1.33	10	1428446_at	AK008822	2.038	17
DYNLT1	1388903_at	T-complex associated-testis-expressed 1-like (Protein 91/23)	AI179335	1.51	4	201999_s_at	NM_006519	1.66	8	1420614_at	NM_025975	1.846	15

DYNLT1		1388903_at	T-complex associated-testis-expressed 1-like (Protein 91/23)	AI179335	1.51	4	201999_s_at	NM_006519	1.66	8	1449928_at	NM_025975	1.767	15
EBAG9	C	1391510_at	estrogen receptor-binding fragment-associated gene 9 (predicted)	BF398331	1.7	6	204274_at	AA812215	1.86	14	1422669_at	AY009091	1.607	13
EBAG9	C	1391510_at	estrogen receptor-binding fragment-associated gene 9 (predicted)	BF398331	1.7	6	204278_s_at	NM_004215	1.27	8	1422669_at	AY009091	1.607	13
ECHDC1	C	1374585_at	enoyl Coenzyme A hydratase domain containing 1	AW915152	1.7	4	219974_x_at	NM_018479	1.2	8	1425001_at	BC023104	1.523	17
EFCBP2	A	1368443_at	EF hand calcium binding protein 2	NM_133415	1.73	6	215005_at	AV723666	1.29	10	1418881_at	NM_054095	5.643	17
EGR4	B	1387442_at	early growth response 4	NM_019137	7.92	2	207767_s_at	NM_001965	2.18	14	1449977_at	NM_020596	4.169	17
EGR4	B	1375784_at	Early growth response 4	BF412806	1.37	6	207768_at	NM_001965	34.1	14	1449977_at	NM_020596	4.169	17
EIF3S10	C	1389968_at	eukaryotic translation initiation factor 3, subunit 10 (theta) (predicted)	BG376953	1.38	4	200596_s_at	BE614908	1.27	14	1416661_at	AW701127	8.263	17
EIF3S10	C	1389968_at	eukaryotic translation initiation factor 3, subunit 10 (theta) (predicted)	BG376953	1.38	4	200595_s_at	NM_003750	1.65	12	1416660_at	AW701127	1.907	13
ELAVL4	A	1389975_at	ELAV (embryonic lethal, abnormal vision, Drosophila)-like 4 (Hu antigen)	BE116949	2.23	4	206051_at	NM_021952	3.69	10	1428741_at	AK013588	6.031	15
ELAVL4	A	1389975_at	ELAV (embryonic lethal, abnormal vision, Drosophila)-like 4 (Hu antigen)	BE116949	2.23	4	206051_at	NM_021952	3.69	10	1450258_a_at	NM_010488	3.83	15
ELAVL4	A	1389975_at	ELAV (embryonic lethal, abnormal vision, Drosophila)-like 4 (Hu antigen)	BE116949	2.23	4	206051_at	NM_021952	3.69	10	1452894_at	AK013588	3.563	15
ELAVL4	A	1389975_at	ELAV (embryonic lethal, abnormal vision, Drosophila)-like 4 (Hu antigen)	BE116949	2.23	4	206051_at	NM_021952	3.69	10	1457399_at	BB211758	2.708	17
ELMO3		1374995_at	engulfment and cell motility 3, ced-12 homolog (C. elegans) (predicted)	BI291645	3.9	4	219411_at	NM_024712	1.43	10	1434489_at	AI481208	2.041	17
EPB41L3	A	1370503_s_at	erythrocyte protein band 4.1-like 3	BM383451	3.08	4	206710_s_at	NM_012307	2.14	10	1440595_at	BB036542	7.606	17
EPB41L3	A	1368515_at	erythrocyte protein band 4.1-like 3	NM_053927	2.52	4	212681_at	AI770004	3.1	14	1426010_a_at	AF177146	4.595	15
ERO1L	B	1389297_at	ERO1-like (S. cerevisiae)	AI146215	2.61	6	218498_s_at	NM_014584	1.41	12	1449324_at	BM234652	2.092	13
ERO1LB	B		ERO1-like beta (S. cerevisiae)	not on RG230.20 array			220012_at	NM_019891	14	14	1449948_at	NM_026184	18.01	17
ERO1LB	B		ERO1-like beta (S. cerevisiae)	not on RG230.20 array			220012_at	NM_019891	14	14	1434714_at	BB234316	18.2	17
ERO1LB	B		ERO1-like beta (S. cerevisiae)	not on RG230.20 array			220012_at	NM_019891	14	14	1425705_a_at	AF489856	12.61	17
ERP29	C	1387031_at	endoplasmic reticulum protein 29	U36482	2.23	6	201216_at	NM_006817	2.41	14	1453634_a_at	AK013303	6.92	17
FBXL15	C	1374114_at	F-box and leucine-rich repeat protein 15 (predicted)	BI282971	1.86	6	218938_at	NM_024326	1.71	12	1417855_at	NM_133694	5.81	17
FIP1L1	C	1389621_at	FIP1 like 1 (S. cerevisiae) (predicted)	BE113454	1.45	4	221007_s_at	NM_030917	1.43	8	1428280_at	BM199874	3.019	17
FIP1L1	C	1389621_at	FIP1 like 1 (S. cerevisiae) (predicted)	BE113454	1.45	4	221007_s_at	NM_030917	1.43	8	1452720_a_at	BM199874	1.883	15
FKBP2	C	1375669_at	FK506 binding protein 2 (predicted)	BI285619	2.1	6	203391_at	NM_004470	2.17	14	1450694_at	NM_008020	1.951	17
FOSB	C	1375043_at	FBJ murine osteosarcoma viral oncogene homolog	BF415939	14.6	6	202768_at	NM_006732	5.04	10	1422134_at	NM_008036	12.58	17
FOSB	C	1375043_at	FBJ murine osteosarcoma viral oncogene homolog	BF415939	14.6	6	202768_at	NM_006732	5.04	10	1423100_at	AV026617	7.402	17
FOXA2		1368711_at	forkhead box A2	NM_012743	10.6	4	210103_s_at	AB028021	4.43	14	1422833_at	NM_010446	10.66	17
FOXA2		1368711_at	forkhead box A2	NM_012743	10.6	4	40284_at	AB028021	2.88	14	1422833_at	NM_010446	10.66	17
G6PC2	B		glucose-6-phosphatase, catalytic, 2	not on RG230.20 array			221453_at	NM_021176	8.76	14	1423529_at	Z47787	51.7	17
GAD1	A	1368344_at	glutamate decarboxylase 1	NM_017007	44.4	6	205278_at	NM_000817	2.48	8	1416561_at	AF326547	2.981	13
GAD1	A	1370760_a_at	glutamate decarboxylase 1	M38350	7.6	6	205278_at	NM_000817	2.48	8	1416561_at	AF326547	2.981	13
GADD45G		1388792_at	growth arrest and DNA-damage-inducible 45 gamma (predicted)	AI599423	3.34	6	204121_at	NM_006705	1.65	12	1453851_a_at	AK007410	8.5	17
GARNL1	C	1368994_a_at	GTPase activating RANGAP domain-like 1	AF041107	1.43	6	214855_s_at	AL050050	1.27	8	1458117_at	BB468955	2.343	17
GARNL4	B		GTPase activating Rap/RanGAP domain-like 4	not on RG230.20 array			213280_at	AK000478	5.06	14	1434754_at	AU067654	6.699	17
GARS		1388715_at	glycyl-tRNA synthetase (predicted)	AA818089	1.72	6	208693_s_at	D30658	1.58	12	1423784_at	BC021747	1.525	17
GARS	C	1388715_at	glycyl-tRNA synthetase (predicted)	AA818089	1.72	6	208693_s_at	D30658	1.58	12	1423784_at	BC021747	1.525	17
GCGR		1370522_at	glucagon receptor	L04796	3.45	4	210565_at	U03469	1.36	12	1450127_a_at	NM_008101	3.092	15
GCK		1387312_a_at	glucokinase	NM_012565	1.86	4	211167_s_at	M69051	2.05	14	1419146_a_at	BC011139	14.77	17
GCK		1387312_a_at	glucokinase	NM_012565	1.86	4	211167_s_at	M69051	2.05	14	1425303_at	L38990	8.96	17
GFPT1	C	1377314_at	glutamine fructose-6-phosphate transaminase 1	BE104391	1.36	4	202722_s_at	NM_002056	3.65	14	1444329_at	BB533749	2.147	17
GFPT1	C	1373373_at	Glutamine fructose-6-phosphate transaminase 1	BI295011	2.54	6	202721_s_at	BE645771	1.64	12	1444329_at	BB533749	2.147	17
GGA2	C	1372408_at	golgi associated, gamma adaptin ear containing, ARF binding protein 2	BI278620	1.81	6	210658_s_at	BC000284	1.7	10	1428141_at	AK004632	2.12	13
GGA2	C	1372408_at	golgi associated, gamma adaptin ear containing, ARF binding protein 2	BI278620	1.81	6	208913_at	AA868560	1.52	10	1428141_at	AK004632	2.12	13
GGA2	C	1372408_at	golgi associated, gamma adaptin ear containing, ARF binding protein 2	BI278620	1.81	6	208915_s_at	AF190863	1.16	10	1428141_at	AK004632	2.12	13
GGA2	C	1372408_at	golgi associated, gamma adaptin ear containing, ARF binding protein 2	BI278620	1.81	6	214190_x_at	AI799984	1.14	12	1428141_at	AK004632	2.12	13
GLA		1373199_at	Galactosidase, alpha (mapped)	AA943573	1.57	4	214430_at	NM_000169	1.14	10	1449006_at	NM_013463	2.331	17
GLG1	C	1387367_at	golgi apparatus protein 1	NM_017211	2.31	6	214730_s_at	AK025457	1.51	12	1448580_at	BG074127	2.054	17
GLG1	C	1387367_at	golgi apparatus protein 1	NM_017211	2.31	6	212045_at	N32761	1.15	12	1448580_at	BG074127	2.054	17
GLP1R	A	1369699_at	glucagon-like peptide 1 receptor	NM_012728	8.09	6	208391_s_at	NM_002062	1.6	14	1422330_at	NM_021332	7.109	17

GLP1R	A	1369699_at	glucagon-like peptide 1 receptor	NM_012728	8.09	6	208401_s_at	U01157	1.35	12	1422330_at	NM_021332	7.109	17
GLP1R	A	1369699_at	glucagon-like peptide 1 receptor	NM_012728	8.09	6	208390_s_at	U01104	1.16	12	1422330_at	NM_021332	7.109	17
GLP1R	A	1369699_at	glucagon-like peptide 1 receptor	NM_012728	8.09	6	211232_x_at	L23503	1.19	12	1458719_at	BM233846	1.828	17
GLRX2		1374340_at	Glutaredoxin 2 (thioltransferase)	AA848536	1.38	4	219933_at	NM_016066	1.26	8	1422998_a_at	NM_023505	1.754	15
GLRX2		1373675_at	glutaredoxin 2 (thioltransferase) (predicted)	AW525635	1.36	4	219933_at	NM_016066	1.26	8	1422998_a_at	NM_023505	1.754	15
GMPPA		1372859_at	GDP-mannose pyrophosphorylase A	AI598958	1.58	6	218070_s_at	NM_013335	1.34	8	1417578_a_at	NM_133708	1.878	17
GOLGA4	C	1394347_at	golgi autoantigen golgin subtype a4; tGolgin-1	BF568007	2.98	4	201567_s_at	NM_002078	1.89	10	1448803_at	NM_018748	6.103	17
GOLGA4	C	1394347_at	golgi autoantigen golgin subtype a4; tGolgin-1	BF568007	2.98	4	201567_s_at	NM_002078	1.89	10	1460213_at	NM_018748	2.014	15
GOLGA5	C	1376650_at	golgi autoantigen, golgin subfamily a, 5 (predicted)	BF399323	1.66	6	218241_at	NM_005113	1.84	12	1418447_at	NM_013747	1.771	17
GORASP2	C	1370325_at	golgi reassembly stacking protein 2	AF110267	1.49	6	208843_s_at	BC001408	1.57	12	1424710_a_at	BC016455	1.879	17
GORASP2	C	1370325_at	golgi reassembly stacking protein 2	AF110267	1.49	6	207812_s_at	NM_015530	1.36	12	1424710_a_at	BC016455	1.879	17
GORASP2	C	1370325_at	golgi reassembly stacking protein 2	AF110267	1.49	6	208842_s_at	W93787	1.34	14	1424710_a_at	BC016455	1.879	17
GOSR2		1374900_at	golgi SNAP receptor complex member 2	BM388807	1.8	6	213144_at	AI074611	1.19	12	1419371_s_at	NM_019650	1.636	17
GOSR2		1375438_at	golgi SNAP receptor complex member 2	AA964687	1.65	6	213180_s_at	BE895285	1.19	8	1419371_s_at	NM_019650	1.636	17
GPD2		1387670_at	glycerol-3-phosphate dehydrogenase 2	U08027	1.4	2	210007_s_at	U36310	1.75	12	1452741_s_at	BQ175968	4.985	17
GPD2		1369666_at	Glycerol-3-phosphate dehydrogenase 2	U08027	1.34	0	210007_s_at	U36310	1.75	12	1452741_s_at	BQ175968	4.985	17
GSPT1		1371585_at	G1 to S phase transition 1	BE108192	1.41	6	201912_s_at	NM_002094	1.65	10	1455173_at	AW537663	1.648	13
GSPT1		1371585_at	G1 to S phase transition 1	BE108192	1.41	6	215438_x_at	BE906054	1.43	8	1455173_at	AW537663	1.648	13
GSPT1		1371585_at	G1 to S phase transition 1	BE108192	1.41	6	217595_at	AV701723	1.24	10	1455173_at	AW537663	1.648	13
HADH	B	1370237_at	L-3-hydroxyacyl-Coenzyme A dehydrogenase, short chain	AA799574	5.96	6	201036_s_at	NM_005327	4.88	14	1436756_x_at	BB114220	2.089	13
HADH	B	1370237_at	L-3-hydroxyacyl-Coenzyme A dehydrogenase, short chain	AA799574	5.96	6	211569_s_at	AF001903	4.71	14	1455972_x_at	AV018774	2.261	13
HDFFRP3	B	1370458_at	hepatoma-derived growth factor, related protein 3	AF389347	1.41	2	209524_at	AK001280	3.21	8	1435977_at	BM240385	2.504	13
HDFFRP3	B	1374151_at	Hepatoma-derived growth factor, related protein 3	BI289940	1.59	0	209524_at	AK001280	3.21	8	1450924_at	BB291880	1.796	13
HDLBP	C	1398859_at	high density lipoprotein binding protein	U90725	1.52	4	221767_x_at	AA515560	1.47	10	1449615_s_at	C77256	9.396	17
HDLBP	C	1388918_at	High density lipoprotein binding protein	BG378074	1.73	6	200643_at	NM_005336	1.25	8	1415988_at	BG065877	9.246	17
HDLBP	C	1388918_at	High density lipoprotein binding protein	BG378074	1.73	6	200643_at	NM_005336	1.25	8	1419806_at	C77256	7.508	17
HERPUD1	C	1367741_at	homocysteine-inducible, endoplasmic reticulum stress-inducible, ubiqu	NM_053523	4.43	6	217168_s_at	AF217990	1.72	14	1435626_a_at	AI835088	2.46	15
HERPUD1	C	1367741_at	homocysteine-inducible, endoplasmic reticulum stress-inducible, ubiqu	NM_053523	4.43	6	217168_s_at	AF217990	1.72	14	1435626_a_at	AI835088	2.46	15
HIC2		1376400_at	hypermethylated in cancer 2 (predicted)	BG380445	1.08	4	212965_at	BE465318	1.21	12	1454944_at	BE648070	2.167	15
HLXB9			homeo box HB9	not on RG230.20 array			214614_at	AI738662	3.6	14	1460299_at	NM_019944	11.54	17
HMG3	C	1371989_at	high mobility group nucleosomal binding domain 3	BM386357	2.76	4	209377_s_at	AF274949	1.25	12	1434875_a_at	AV018952	3.707	15
HMG3	C	1371989_at	high mobility group nucleosomal binding domain 3	BM386357	2.76	4	209377_s_at	AF274949	1.25	12	1431777_a_at	AK002970	2.612	15
HNRPC		1371505_at	heterogeneous nuclear ribonucleoprotein C (predicted)	BG381750	1.44	4	200751_s_at	BE898861	1.41	10	1418693_at	NM_016884	4.408	17
HNRPC	C	1371505_at	heterogeneous nuclear ribonucleoprotein C (predicted)	BG381750	1.44	4	200014_s_at	NM_004500	1.24	10	1418693_at	NM_016884	4.408	17
HNRPD	C	1372810_at	heterogeneous nuclear ribonucleoprotein D-like (predicted)	BF281256	1.5	4	209067_s_at	D89092	1.4	8	1428224_at	AV101245	1.976	17
HSP90B1	C	1388331_at	heat shock protein 90kDa beta (Grp94), member 1 (tumor rejection anti	BG057543	2.67	6	200598_s_at	AI582238	2.78	14	1438040_a_at	BE995678	3.137	15
HSP90B1	C	1388331_at	heat shock protein 90kDa beta (Grp94), member 1 (tumor rejection anti	BG057543	2.67	6	216449_x_at	AK025862	2.72	14	1438040_a_at	BE995678	3.137	15
HSP90B1	C	1388331_at	heat shock protein 90kDa beta (Grp94), member 1 (tumor rejection anti	BG057543	2.67	6	200599_s_at	NM_003299	1.75	14	1438040_a_at	BE995678	3.137	15
HSPA4L		1371783_at	heat shock 70kDa protein 4-like	AI009936	1.59	6	205543_at	NM_014278	1.73	12	1425993_a_at	D67017	2.093	13
HSPA5 / G	C	1370283_at	heat shock 70kD protein 5	M14050	2.92	6	211936_at	AF216292	1.95	14	1427464_s_at	AJ002387	2.179	11
HSPA5 / G	C	1370283_at	heat shock 70kD protein 5	M14050	2.92	6	211936_at	AF216292	1.95	14	1416064_a_at	NM_022310	1.762	15
HYOU1	A	1371442_at	hypoxia up-regulated 1	BI282904	4.53	6	200825_s_at	NM_006389	1.83	14	1451913_a_at	BC019785	6.251	17
HYOU1	A	1370665_at	hypoxia up-regulated 1	U41853	3.87	6	200825_s_at	NM_006389	1.83	14	1451913_a_at	BC019785	6.251	17
IAPP	B	1387660_at	islet amyloid polypeptide	M25390	106	6	207062_at	NM_000415	59.8	14	1437323_a_at	BB434117	167.2	17
IAPP	B	1387660_at	islet amyloid polypeptide	M25390	106	6	207062_at	NM_000415	59.8	14	1423510_at	BM569571	129.3	17
IAPP	B	1387660_at	islet amyloid polypeptide	M25390	106	6	207062_at	NM_000415	59.8	14	1423509_a_at	BM569571	102.7	17
IAPP	B	1387660_at	islet amyloid polypeptide	M25390	106	6	207062_at	NM_000415	59.8	14	1456004_x_at	BB434117	52.23	17
ICA1		1367787_at	islet cell autoantigen 1	NM_030844	9.43	6	210547_x_at	L21181	1.49	14	1417901_a_at	NM_010492	4.004	17
ICA1		1367787_at	islet cell autoantigen 1	NM_030844	9.43	6	207949_s_at	NM_004968	1.4	12	1417901_a_at	NM_010492	4.004	17
ICA1		1367787_at	islet cell autoantigen 1	NM_030844	9.43	6	214191_at	W67992	1.14	12	1431644_a_at	AK005965	2.774	13

IDS	B	1375843_at	Iduronate 2-sulfatase	BM383777	2.13	2	202439_s_at	NM_000202	5.13	14	1434751_at	BB493523	11.21	17
IDS	B	1375843_at	Iduronate 2-sulfatase	BM383777	2.13	2	212221_x_at	AV703259	4.47	10	1434751_at	BB493523	11.21	17
IDS	B	1375843_at	Iduronate 2-sulfatase	BM383777	2.13	2	202438_x_at	BF346014	4.36	12	1434751_at	BB493523	11.21	17
IDS	B	1375843_at	Iduronate 2-sulfatase	BM383777	2.13	2	212223_at	AI926544	4.31	12	1434751_at	BB493523	11.21	17
IDS	B	1375843_at	Iduronate 2-sulfatase	BM383777	2.13	2	206342_x_at	NM_006123	3.1	14	1434751_at	BB493523	11.21	17
IDS	B	1375843_at	Iduronate 2-sulfatase	BM383777	2.13	2	217432_s_at	AF179281	1.74	14	1434751_at	BB493523	11.21	17
IDS	B	1375843_at	Iduronate 2-sulfatase	BM383777	2.13	2	210666_at	AF050145	1.22	12	1434751_at	BB493523	11.21	17
IDS	B	1375843_at	Iduronate 2-sulfatase	BM383777	2.13	2	211782_at	BC006170	1.63	14	1455481_at	BM241722	2.978	15
IGF1R		1368123_at	insulin-like growth factor 1 receptor	NM_052807	1.97	6	203628_at	H05812	2.13	10	1452108_at	BE980124	6.634	17
IGF1R		1390671_at	Insulin-like growth factor 1 receptor	AI044666	1.35	2	203628_at	H05812	2.13	10	1452108_at	BE980124	6.634	17
IGF2		1367571_a_at	insulin-like growth factor 2	NM_031511	2.28	4	202410_x_at	NM_000612	1.66	10	1448152_at	NM_010514	4.061	15
IGF2		1367571_a_at	insulin-like growth factor 2	NM_031511	2.28	4	210881_s_at	M17863	1.62	10	1424112_at	BG092290	2.121	17
IKBKAP	C	1390504_at	Inhibitor of kappa light polypeptide enhancer in B-cells, kinase comple	AW524470	1.35	6	202491_s_at	NM_003640	1.31	8	1424142_at	AF367244	1.961	17
IL13RA1		1388711_at	interleukin 13 receptor, alpha 1	BF282650	1.82	4	211612_s_at	U62858	2.12	14	1427164_at	BB730912	3.887	17
IL13RA1		1370728_at	interleukin 13 receptor, alpha 1	AY044251	1.62	4	201888_s_at	U81379	2.18	12	1427164_at	BB730912	3.887	17
IL13RA1		1370728_at	interleukin 13 receptor, alpha 1	AY044251	1.62	4	210904_s_at	U81380	1.3	10	1427164_at	BB730912	3.887	17
INS	B	1370077_at	insulin 2	NM_019130	56.7	6	206598_at	NM_000207	105	14	1422446_x_at	NM_008387	46.45	17
INS1 / INS2	B	1387815_at	insulin 1 /// insulin 2	NM_019129	35.2	6	206598_at	NM_000207	105	14	1422447_at	NM_008386	66.14	17
INSM1	B		insulinoma-associated 1	not on RG230.20 array			206502_s_at	NM_002196	25.3	14	1455865_at	BB468410	54.21	17
INSM1	B		insulinoma-associated 1	not on RG230	0	0	206502_s_at	NM_002196	25.3	14	1421399_at	NM_016889	14.43	17
IQGAP1	C	1388764_at	IQ motif containing GTPase activating protein 1 (predicted)	BM387072	1.57	2	200791_s_at	NM_003870	2.26	10	1431395_a_at	BI645591	3.016	15
IQGAP1	C	1388764_at	IQ motif containing GTPase activating protein 1 (predicted)	BM387072	1.57	2	210840_s_at	D29640	2.16	10	1431396_at	BI645591	3.019	15
IRAK1	C	1373379_at	interleukin-1 receptor-associated kinase 1 (predicted)	BF396350	2.01	6	201587_s_at	NM_001569	1.7	10	1460649_at	BE949425	1.83	13
IRF6		1377379_at	interferon regulatory factor 6 (predicted)	BF410603	4.36	6	202597_at	AU144284	1.65	12	1418301_at	NM_016851	1.665	11
IRF6		1386568_at	interferon regulatory factor 6 (predicted)	BF557891	3.51	4	202597_at	AU144284	1.65	12	1418301_at	NM_016851	1.665	11
ITM2C		1371334_at	integral membrane protein 2C (predicted)	BG668228	1.88	-2	221004_s_at	NM_030926	2.13	10	1415961_at	NM_022417	2.191	15
JUN	C	1374404_at	c-Jun / v-jun sarcoma virus 17 oncogene homolog (avian)	BI288619	7.79	4	201465_s_at	BC002646	3.97	12	1448694_at	NM_010591	2.41	15
JUN	C	1389528_s_at	c-Jun / v-jun sarcoma virus 17 oncogene homolog (avian)	BI288619	5.27	6	201466_s_at	NM_002228	3.6	10	1417409_at	NM_010591	3.658	15
JUN	C	1369788_s_at	c-Jun / v-jun sarcoma virus 17 oncogene homolog (avian)	NM_021835	3.1	4	201464_x_at	BG491844	2.87	14	1448694_at	NM_010591	2.41	15
KCNH2		1368343_at	potassium voltage-gated channel, subfamily H (eag-related), member :	NM_053949	1.61	4	205262_at	NM_000238	4.11	14	1449544_a_at	NM_013569	2.737	15
KCNH2		1368343_at	potassium voltage-gated channel, subfamily H (eag-related), member :	NM_053949	1.61	4	210036_s_at	AB044806	2.44	14	1449544_a_at	NM_013569	2.737	15
KIF5B		1372444_at	kinesin family member 5B	AW918352	2.62	6	201992_s_at	NM_004521	1.46	10	1418431_at	BI328541	4.809	17
KIF5B		1372444_at	kinesin family member 5B	AW918352	2.62	6	201991_s_at	BF223224	1.28	8	1418429_at	BI328541	2.029	13
KIF5C	A	1373977_at	Kinesin family member 5C (predicted)	BE108253	5.73	2	203130_s_at	NM_004522	7.19	8	1457255_x_at	BB099427	8.989	17
KIF5C	A	1373977_at	Kinesin family member 5C (predicted)	BE108253	5.73	2	203129_s_at	BF059313	6.16	8	1422945_a_at	AI844677	4.754	17
KIF5C	A	1373977_at	Kinesin family member 5C (predicted)	BE108253	5.73	2	203129_s_at	BF059313	6.16	8	1450804_at	AI844677	1.783	17
LEPR		1369739_at	leptin receptor	D85558	1.41	6	209894_at	U50748	2.22	10	1425873_a_at	U58861	2.541	17
LEPR		1369739_at	leptin receptor	D85558	1.41	6	211354_s_at	U52913	1.42	8	1425873_a_at	U58861	2.541	17
LEPR		1369739_at	leptin receptor	D85558	1.41	6	211356_x_at	U66495	1.33	8	1425873_a_at	U58861	2.541	17
LEPR		1369739_at	leptin receptor	D85558	1.41	6	211355_x_at	U52914	1.31	8	1425873_a_at	U58861	2.541	17
LMAN1	A	1368848_at	lectin, mannose-binding, 1	NM_053886	3.9	6	203293_s_at	NM_005570	1.84	14	1428130_at	BG071597	5.81	17
LMAN1	A	1368848_at	lectin, mannose-binding, 1	NM_053886	3.9	6	203293_s_at	NM_005570	1.84	14	1428129_at	BG071597	4.626	17
LMAN1	A	1368848_at	lectin, mannose-binding, 1	NM_053886	3.9	6	203293_s_at	NM_005570	1.84	14	1452671_s_at	BG071597	3.926	17
LMAN2		1373815_at	lectin, mannose-binding 2 (predicted)	BI304064	1.49	4	200805_at	NM_006816	1.68	10	1423074_at	AK004952	1.527	13
LNPEP		1370743_a_at	leucyl/cystinyl aminopeptidase	AY079189	1.7	6	207904_s_at	NM_005575	1.1	12	1438432_at	BE850004	1.948	17
LONP1		1375802_at	Ion peptidase 1, mitochondrial (Protease, serine, 15)	AI011610	1.69	6	209017_s_at	U02389	1.38	14	1428365_a_at	AK004820	2.054	17
LRBA		1374495_at	LPS-responsive beige-like anchor (predicted)	BM388121	2.57	6	212692_s_at	W60686	1.15	10	1436690_at	BB494139	3.505	17
LRBA		1374495_at	LPS-responsive beige-like anchor (predicted)	BM388121	2.57	6	212692_s_at	W60686	1.15	10	1425717_at	AF188506	2.313	17
LRBA		1374495_at	LPS-responsive beige-like anchor (predicted)	BM388121	2.57	6	214109_at	AI659561	1.18	10	1449099_at	NM_030695	1.941	13
LRP12		1374108_at	low density lipoprotein-related protein 12 (predicted)	BM388942	1.43	4	220254_at	NM_013437	9.99	10	1433864_at	AV254798	2.807	17

LRP12	1374108_at	low density lipoprotein-related protein 12 (predicted)	BM388942	1.43	4	220253_s_at	NM_013437	2.71	12	1433864_at	AV254798	2.807	17
LRP12	1374108_at	low density lipoprotein-related protein 12 (predicted)	BM388942	1.43	4	219631_at	NM_024937	2.19	12	1433864_at	AV254798	2.807	17
LRRC8D	1373431_at	leucine rich repeat containing 8 family, member D (alias: leucine rich re	AI101427	1.77	4	218684_at	NM_018103	1.31	12	1447491_at	BE987722	2.488	17
LRRRC8D	1373431_at	leucine rich repeat containing 8 family, member D (alias: leucine rich re	AI101427	1.77	4	218684_at	NM_018103	1.31	12	1447491_at	BB315861	1.996	15
MAGED1	1367744_at	melanoma antigen, family D, 2	NM_080479	2.83	2	208682_s_at	AF126181	2.23	14	1426306_a_at	AF319976	1.534	13
MAGED1	1367744_at	melanoma antigen, family D, 2	NM_080479	2.83	2	213627_at	AI924630	1.41	10	1426306_a_at	AF319976	1.534	13
MAGED2	1386895_at	melanoma antigen, family D, 1	NM_053409	4.44	6	209014_at	AF217963	1.71	8	1450062_a_at	NM_019791	3.699	17
MAP1B	A 1371003_at	microtubule-associated protein 1b	BG378086	1.51	2	212233_at	AL523076	4.6	8	1450397_at	BB731480	11.28	17
MAP1B	A 1371003_at	microtubule-associated protein 1b	BG378086	1.51	2	212233_at	AL523076	4.6	8	1421851_at	BB731480	6.253	17
MAP1B	A 1371003_at	microtubule-associated protein 1b	BG378086	1.51	2	212233_at	AL523076	4.6	8	1421850_at	BB731480	2.08	13
MAP1B	A 1373363_at	microtubule-associated protein 1b	BI281702	2.39	0	214577_at	BG164365	1.93	8	1444595_at	BF148767	2.656	17
MAP2	1388152_at	microtubule-associated protein 2	BG374290	20.5	2	210015_s_at	U89330	2.84	10	1443239_at	BB386302	2.065	13
MAP2	1368411_a_at	microtubule-associated protein 2	X74211	1.84	4	210015_s_at	U89330	2.84	10	1443239_at	BB386302	2.065	13
MBNL2	A 1375140_at	muscleblind-like 2 (predicted)	AA818999	1.52	2	203640_at	BE328496	1.61	10	1436858_at	BB003847	1.557	7
MED6	C 1372269_at	mediator of RNA polymerase II transcription, subunit 6 homolog (yeast	AI229534	2.36	6	210104_at	AF074723	1.24	12	1458343_x_at	AV070715	1.543	17
MEFAP3	1390063_at	microfibrillar-associated protein 3	BG380473	2.5	6	214588_s_at	R38475	1.22	10	1434562_at	AW536327	6.121	17
MINPP1	1388126_at	multiple inositol polyphosphate histidine phosphatase 1	BG380493	2.17	4	209585_s_at	AF084943	2.83	14	1455787_x_at	AV339366	3.048	17
MLX	1371929_at	MAX-like protein X (transcription factor-like 4) (predicted)	BI278563	1.74	6	217909_s_at	BF056105	1.23	10	1418437_a_at	NM_011550	3.64	17
MYO1D	B 1388917_at	myosin ID	BI279786	3.86	6	212338_at	AA621962	7.66	14	1458550_at	BB200258	2.347	17
MYO1D	B 1369779_at	myosin ID	NM_012983	1.71	6	212338_at	AA621962	7.66	14	1458550_at	BB200258	2.347	17
MYO5A	1368450_at	myosin Va	NM_022178	1.75	4	204527_at	NM_000259	1.31	10	1419754_at	NM_010864	1.5	13
NAP1L1	C 1371872_at	Nucleosome assembly protein 1-like 1	BE108905	1.36	0	204528_s_at	NM_004537	1.87	10	1420477_at	BG064031	1.73	13
NAP1L1	C 1371872_at	Nucleosome assembly protein 1-like 1	BE108905	1.36	0	213864_s_at	AI985751	1.81	10	1420477_at	BG064031	1.73	13
NAP1L1	C 1371872_at	Nucleosome assembly protein 1-like 1	BE108905	1.36	0	212967_x_at	AW148801	1.6	10	1420477_at	BG064031	1.73	13
NAP1L1	C 1371872_at	Nucleosome assembly protein 1-like 1	BE108905	1.36	0	208752_x_at	AI888672	1.55	12	1420477_at	BG064031	1.73	13
NCALD	A 1372953_at	neurocalcin delta	AI101330	3.49	2	211685_s_at	AF251061	1.9	8	1417568_at	BG071381	1.688	11
NCBP2	1383965_at	nuclear cap binding protein subunit 2 (predicted)	BI282103	2.73	2	201517_at	BC001255	1.23	10	1423046_s_at	BE285362	1.859	17
NCBP2	1383965_at	nuclear cap binding protein subunit 2 (predicted)	BI282103	2.73	2	201521_s_at	NM_007362	1.13	10	1423046_s_at	BE285362	1.859	17
NEFM	A 1367845_at	neurofilament 3, medium	NM_017029	3.27	2	205113_at	NM_005382	10.8	10	1457015_at	BB258427	3.819	15
NEK1	1389996_at	NIMA (never in mitosis gene a)-related expressed kinase 1 (predicted)	AI406369	1.48	4	213331_s_at	AV700007	2.02	12	1453612_at	AV254337	5.595	17
NEK1	1389996_at	NIMA (never in mitosis gene a)-related expressed kinase 1 (predicted)	AI406369	1.48	4	213331_s_at	AV700007	2.02	12	1434267_at	BG069735	1.783	13
NEO1	1376786_a_at	Neogenin	BF397709	2.79	6	204321_at	NM_002499	1.59	12	1447693_s_at	BB350308	2.818	17
NEUROD1	A 1394135_at	Neurogenic differentiation 1	AI639109	14.7	6	206282_at	NM_002500	21	12	1426413_at	BM116592	4.207	13
NEUROD1	A 1387288_at	neurogenic differentiation 1	NM_019218	7.09	6	206282_at	NM_002500	21	12	1426412_at	BM116592	3.089	13
NFIC	1371176_at	nuclear factor I/C	AB012233	1.35	4	206929_s_at	NM_005597	1.19	10	1450661_x_at	NM_008688	2.342	17
NKX2-2	B 1385387_at	NK2 transcription factor related, locus 2 (Drosophila) (predicted)	not on RG230A array			206915_at	NM_002509	43.8	14	1421112_at	NM_010919	40.65	17
NKX6-1	B 1368998_at	NK6 transcription factor related, locus 1 (Drosophila)	NM_031737	18.8	6	221366_at	NM_006168	2.51	14	1425828_at	AF357883	79.27	17
NPTX2	B	neuronal pentraxin II	not on RG230.20 array			213479_at	U26662	32.3	14	1420720_at	NM_016789	2.606	13
NPY	1387154_at	neuropeptide Y	NM_012614	2.12	6	206001_at	NM_000905	1.93	10	1419127_at	NM_023456	1.489	9
NUCB2	C 1370000_at	nucleobindin 2	NM_021663	6.77	6	203675_at	NM_005013	6.34	14	1418355_at	NM_016773	6.486	17
NUP98	C 1368747_at	nucleoporin 98	NM_031074	1.43	4	210793_s_at	U41815	1.67	10	1439154_at	BB473634	1.954	17
P2RY1	1370606_at	purinergic receptor P2Y, G-protein coupled 1	U22830	19.9	6	207455_at	NM_002563	1.17	10	1421456_at	NM_008772	7.131	17
PAFAH1B1	A 1369641_at	platelet-activating factor acetylhydrolase, isoform 1b, alpha2 subunit	BM392366	1.58	6	200816_s_at	NM_000430	1.68	8	1422793_at	BB539054	2.035	13
PAFAH1B1	A 1369642_at	platelet-activating factor acetylhydrolase, isoform 1b, alpha2 subunit	BM392366	1.45	0	200816_s_at	NM_000430	1.68	8	1422793_at	BB539054	2.035	13
PAM	B 1370502_at	peptidylglycine alpha-amidating monooxygenase	AW534339	3.73	6	214620_x_at	BF038548	6.34	14	1418908_at	NM_013626	2.265	15
PAM	B 1367687_a_at	peptidylglycine alpha-amidating monooxygenase	AW534339	3.77	6	202336_s_at	NM_000919	5.59	14	1418908_at	NM_013626	2.265	15
PAM	B 1367687_a_at	peptidylglycine alpha-amidating monooxygenase	AW534339	3.77	6	212958_x_at	AI022882	5.36	14	1418908_at	NM_013626	2.265	15
PAPSS2	B 1376047_at	3'-phosphoadenosine 5'-phosphosulfate synthase 2 (predicted)	BI285321	5.84	6	203058_s_at	AW299958	10.3	14	1421987_at	BF786072	7.396	17
PAPSS2	B 1376047_at	3'-phosphoadenosine 5'-phosphosulfate synthase 2 (predicted)	BI285321	5.84	6	203060_s_at	AF074331	8.86	14	1434510_at	BF780807	3.808	15
PAPSS2	B 1376047_at	3'-phosphoadenosine 5'-phosphosulfate synthase 2 (predicted)	BI285321	5.84	6	203059_s_at	NM_004670	4.08	14	1421989_s_at	BF786072	5.349	17

PAX6	B	1369242_at	paired box gene 6	NM_013001	15.4	4	205646_s_at	NM_000280	17.4	14	1452526_a_at	AF443223	9.312	15
PAX6	B	1369242_at	paired box gene 6	NM_013001	15.4	4	205646_s_at	NM_000280	17.4	14	1419271_at	BC011272	6.937	15
PAX6	B	1369242_at	paired box gene 6	NM_013001	15.4	4	205646_s_at	NM_000280	17.4	14	1425960_s_at	AF457142	4.103	17
PCLO	A	1387399_at	piccolo (presynaptic cytomatrix protein)	AF138789	1.43	6	213558_at	AB011131	2.69	8	1452423_at	AW493746	6.142	17
PCLO	A	1387399_at	piccolo (presynaptic cytomatrix protein)	AF138789	1.43	6	210650_s_at	BC001304	1.64	8	1419392_at	NM_011995	4.19	15
PCSK1	B	1368559_at	proprotein convertase subtilisin/kexin type 1	M83745	22.3	6	205825_at	NM_000439	17.3	14	1421396_at	M69196	28.62	17
PCSK1	B	1387247_at	proprotein convertase subtilisin/kexin type 1	M83745	21.6	6	205825_at	NM_000439	17.3	14	1421396_at	M69196	28.62	17
PCSK2	B	1387155_at	proprotein convertase subtilisin/kexin type 2	NM_012746	15.5	4	204869_at	AL031664	7.31	14	1447991_at	AI839700	44.29	17
PCSK2	B	1387155_at	proprotein convertase subtilisin/kexin type 2	NM_012746	15.5	4	204870_s_at	NM_002594	15.4	14	1448312_at	NM_008792	17.11	17
PCSK2	B	1387155_at	proprotein convertase subtilisin/kexin type 2	NM_012746	15.5	4	204869_at	AL031664	7.31	14	1447992_s_at	AI839700	24.52	17
PCSK2	B	1387155_at	proprotein convertase subtilisin/kexin type 2	NM_012746	15.5	4	204869_at	AL031664	7.31	14	1428305_at	AK018159	21.61	17
PCSK2	B	1387155_at	proprotein convertase subtilisin/kexin type 2	NM_012746	15.5	4	204869_at	AL031664	7.31	14	1444147_at	BB357975	5.453	17
PDGFRL		1374616_at	platelet-derived growth factor receptor-like (predicted)	BM384311	1.35	4	205226_at	NM_006207	2.04	8	1428896_at	AK004179	2.982	15
PDIA3 / GR	C	1398788_at	protein disulfide isomerase associated 3 (glucose regulated protein, 58	NM_017319	2.67	6	208612_at	D83485	2.16	14	1445330_at	BG916492	2.195	17
PDIA6	C	1370859_at	protein disulfide isomerase associated 6 (thioredoxin domain containin	BI284965	2.79	6	216640_s_at	AK026926	2.31	14	1423648_at		3.716	17
PDIA6	C	1370859_at	protein disulfide isomerase associated 6 (thioredoxin domain containin	BI284965	2.79	6	207668_x_at	NM_005742	2.31	14	1423648_at		0 3.716	17
PDIA6	C	1370859_at	protein disulfide isomerase associated 6 (thioredoxin domain containin	BI284965	2.79	6	208639_x_at	BC001312	2.26	14	1423648_at		0 3.716	17
PDIA6	C	1370859_at	protein disulfide isomerase associated 6 (thioredoxin domain containin	BI284965	2.79	6	208638_at	BE910010	1.75	14	1423648_at		0 3.716	17
PDX1		1369516_at	pancreatic and duodenal homeobox gene 1	NM_022852	3.42	6	210937_s_at	U35632	1.18	10	1422174_at	AK020261	47.04	17
PDX1		1369516_at	pancreatic and duodenal homeobox gene 1	NM_022852	3.42	6	210937_s_at	U35632	1.18	10	1422173_at	AK020261	12.43	17
PFKFB2	B	1388044_at	6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 2	L27084	4.33	6	209992_at	AB044805	2.26	14	1422092_at	BC018418	9.607	17
PFKFB2	B	1369400_a_at	6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 2	NM_080477	1.73	6	207931_s_at	NM_006212	1.36	12	1422090_a_at	BC018418	5.204	17
PFKFB2	B	1369400_a_at	6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 2	NM_080477	1.73	6	207931_s_at	NM_006212	1.36	12	1422091_at	BC018418	3.504	15
PGCP		1368399_a_at	plasma glutamate carboxypeptidase	NM_031640	1.82	2	208454_s_at	NM_016134	1.5	14	1416441_at	BB468025	3.014	17
PGCP		1368399_a_at	plasma glutamate carboxypeptidase	NM_031640	1.82	2	203501_at	NM_006102	1.3	10	1416441_at	BB468025	3.014	17
PGRMC2	C	1388531_at	progesterone receptor membrane component 2 (predicted)	BF283382	1.67	4	201701_s_at	NM_006320	2.21	12	1448233_at	BE630020	2.349	15
PGRMC2	C	1388531_at	progesterone receptor membrane component 2 (predicted)	BF283382	1.67	4	213227_at	BE879873	1.77	14	1452882_at	BF322962	2.126	17
PICALM		1369452_a_at	phosphatidylinositol binding clathrin assembly protein	NM_053554	1.4	6	215832_x_at	AV722190	1.48	14	1443218_at	BB206107	2.015	17
PITX3		1370222_at	paired-like homeodomain transcription factor 3	AI145840	1.53	4	208277_at	NM_005029	1.31	12	1449917_at	NM_008852	2.733	17
PJA2		1375421_a_at	praja 2, RING-H2 motif containing	AI600019	1.93	2	201133_s_at	AA142966	1.43	10	1424442_a_at	BC017130	3.494	17
PJA2	B	1387903_at	praja 2, RING-H2 motif containing	D32249	1.4	2	201133_s_at	AA142966	1.43	10	1424442_a_at	BC017130	3.494	17
PLCB4	B	1386962_at	phospholipase C, beta 4	NM_024353	1.54	-2	203895_at	AL535113	2.63	12	1421987_at	BF786072	7.396	17
POFUT2	C	1390590_at	protein O-fucosyltransferase 2 (predicted)	AA965207	1.68	6	209578_s_at	BC000626	1.6	12	1435056_x_at	BB699319	2.464	15
POFUT2	C	1388717_at	protein O-fucosyltransferase 2 (predicted)	BM384163	1.7	6	209578_s_at	BC000626	1.6	12	1416573_at	BC018194	1.854	17
PPP1R1A	B	1386968_at	protein phosphatase 1, regulatory (inhibitor) subunit 1A	NM_022676	10	6	205478_at	NM_006741	8.52	12	1422605_at	NM_021391	4.05	13
PPP2R1B		1373959_at	protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), bet	AI411788	1.85	2	202886_s_at	M65254	2.2	14	1428265_at	AK010754	1.682	15
PPP2R1B		1373959_at	protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), bet	AI411788	1.85	2	202884_s_at	NM_002716	1.46	10	1428265_at	AK010754	1.682	15
PRKACB	C	1370746_at	protein kinase, cAMP dependent, catalytic, beta (predicted)	D10770	2.23	2	202741_at	AA130247	2.91	8	1420610_at	AV024339	3.263	15
PRKACB	C	1371835_at	protein kinase, cAMP dependent, catalytic, beta (predicted)	AI411907	1.7	0	202742_s_at	NM_002731	3.24	10	1420611_at	AV024339	2.209	15
PRKAR1A	C	1386905_at	protein kinase, cAMP dependent regulatory, type I, alpha	NM_013181	1.57	4	200604_s_at	M18468	2.42	10	1447635_at	BB410911	3.689	17
PRKAR1A	C	1386905_at	protein kinase, cAMP dependent regulatory, type I, alpha	NM_013181	1.57	4	200605_s_at	NM_002734	1.78	10	1425550_a_at	BC005697	1.875	15
PRNP	A	1370156_at	prion protein	BI278802	6.31	6	215707_s_at	AV725328	1.21	8	1416130_at	BE630020	2.294	13
PRPS1	C	1368165_at	phosphoribosyl pyrophosphate synthetase 1	M29392	1.75	6	208447_s_at	NM_002764	1.28	10	1416052_at	AK011304	2.404	17
PRPS1	C	1387085_at	phosphoribosyl pyrophosphate synthetase 1	M29392	1.82	6	209440_at	BC001605	1.85	10	1448192_s_at	AK011304	1.606	15
PTK2	A	1394338_x_at	PTK2 protein tyrosine kinase 2	BF290004	1.35	6	208820_at	AL037339	1.48	10	1430827_x_at	AV301702	2.484	17
PTK2	A	1394338_x_at	PTK2 protein tyrosine kinase 2	BF290004	1.35	6	207821_s_at	NM_005607	1.21	10	1430827_a_at	AV301702	2.484	17
PTPRF		1368035_a_at	protein tyrosine phosphatase, receptor type, F	X83505	3.23	2	200635_s_at	AU145351	1.36	8	1420842_at	BF235516	2.247	15
PTPRF		1368035_a_at	protein tyrosine phosphatase, receptor type, F	X83505	3.23	2	200635_s_at	AU145351	1.36	8	1420841_at	BF235516	1.817	13
PTPRF		1368036_at	protein tyrosine phosphatase, receptor type, F	M60103	1.79	4	200637_s_at	AI762627	1.68	12	1420843_at	BF235516	2.462	15
PTPRN		1370362_at	protein tyrosine phosphatase, receptor type, N	D45414	4.6	6	204945_at	NM_002846	8.66	14	1416588_at	NM_008985	5.366	15

PTPRN2	A	1370182_at	protein tyrosine phosphatase, receptor type, N polypeptide 2	AI574884	12.5	6	203029_s_at	NM_002847	7	14	1425724_at	U57345	1.428	17
PTPRN2	A	1370182_at	protein tyrosine phosphatase, receptor type, N polypeptide 2	AI574884	12.5	6	203030_s_at	AF007555	5.32	14	1425724_at	U57345	1.428	17
PTPRN2	A	1370182_at	protein tyrosine phosphatase, receptor type, N polypeptide 2	AI574884	12.5	6	211534_x_at	U65065	1.44	12	1425724_at	U57345	1.428	17
RAB14	C	1367884_at	RAB14, member RAS oncogene family	NM_053589	1.36	0	211503_s_at	AF112206	1.26	10	1419245_at	AV339290	2.42	17
RAB1A	C	1398841_at	RAB1, member RAS oncogene family	AI411122	1.62	6	207791_s_at	NM_004161	1.95	12	1416082_at	AW108405	1.627	15
RAB1A	C	1373022_at	RAB1, member RAS oncogene family	AI231860	1.94	4	208724_s_at	BC000905	1.37	14	1416082_at	AW108405	1.627	15
RAB2A	B	1370087_at	RAB2, member RAS oncogene family	NM_031718	1.42	6	208731_at	AU158062	1.82	10	1418623_at	NM_021518	1.91	15
RAB2A	B	1370087_at	RAB2, member RAS oncogene family	NM_031718	1.42	6	208732_at	AI743756	1.34	8	1418623_at	NM_021518	1.91	15
RAB2A	B	1370087_at	RAB2, member RAS oncogene family	NM_031718	1.42	6	208733_at	AW301641	1.31	10	1418623_at	NM_021518	1.91	15
RAB2A	B	1370087_at	RAB2, member RAS oncogene family	NM_031718	1.42	6	208734_x_at	M28213	1.62	14	1419946_s_at	C80220	1.566	15
RAB3B		1370061_at	RAB3B, member RAS oncogene family	NM_031091	1.94	4	205924_at	BC005035	4.64	14	1422583_at	NM_023537	2.52	13
RAB3B		1370061_at	RAB3B, member RAS oncogene family	NM_031091	1.94	4	205925_s_at	NM_002867	2.63	14	1422583_at	NM_023537	2.52	13
RANBP2		1375371_at	similar to Ran-binding protein 2 (predicted)	BI296662	1.39	4	201713_s_at	D42063	2.44	14	1440104_at	BB208180	5.203	17
RANBP2		1375371_at	similar to Ran-binding protein 2 (predicted)	BI296662	1.39	4	201711_x_at	AI681120	1.62	8	1440104_at	BB208180	5.203	17
RANBP2		1375371_at	similar to Ran-binding protein 2 (predicted)	BI296662	1.39	4	201712_s_at	NM_006267	1.36	14	1440104_at	BB208180	5.203	17
RBBP7	C	1398768_at	retinoblastoma binding protein 7	NM_031816	1.45	4	201092_at	NM_002893	1.44	10	1439131_at	BB022577	2.616	17
RBM39	C	1398993_at	RNA-binding region (RNP1, RRM) containing 2 (predicted)	AI045458	1.63	2	208720_s_at	AI890947	1.24	10	1438398_at	BB203348	3.594	17
RBM39	C	1399101_at	RNA-binding region (RNP1, RRM) containing 2 (predicted)	AA849715	1.51	2	208720_s_at	AI890947	1.24	10	1456386_at	BB203348	2.437	15
RBM39	C	1381967_at	RNA-binding region (RNP1, RRM) containing 2 (predicted)	BE114972	1.48	2	208720_s_at	AI890947	1.24	10	1438420_at	AV019076	1.886	13
RBM39	C	1379737_a_at	RNA-binding region (RNP1, RRM) containing 2 (predicted)	BG372903	1.44	2	208720_s_at	AI890947	1.24	10	1438397_a_at	BB203348	1.785	15
RGS4	A	1368505_at	regulator of G-protein signaling 4	U27767	5.5	6	204338_s_at	NM_005613	7.48	12	1416287_at	NM_009062	8.283	15
RGS4	A	1368505_at	regulator of G-protein signaling 4	U27767	5.5	6	204339_s_at	BC000737	5.6	12	1416286_at	NM_009062	9.081	15
RGS4	A	1368506_at	regulator of G-protein signaling 4	U27767	5.22	6	204337_at	AL514445	9.12	12	1448285_at	NM_009062	3.413	15
ROBO1	A	1383767_at	Roundabout homolog 1 (Drosophila)	AW524430	1.4	6	213194_at	BF059159	2.32	12	1427231_at	BG065230	2.391	11
RPH3AL	B	1387176_at	rabphilin 3A-like (without C2 domains)	NM_133591	5.17	4	221614_s_at	BC005153	1.69	14	1431695_at	AK018335	2.094	13
RRAGA	C	1367736_at	Ras-related GTP binding A	NM_053973	1.71	6	201628_s_at	NM_006570	1.33	8	1428905_at	AI118026	1.591	15
RRAGD	B	1373427_at	Ras-related GTP binding D (predicted)	BI288816	1.41	2	221523_s_at	AL138717	5.61	12	1431164_at	AK017818	3.236	17
RRAGD	B	1373427_at	Ras-related GTP binding D (predicted)	BI288816	1.41	2	221524_s_at	AF272036	2.98	12	1431164_at	AK017818	3.236	17
SC4MOL		1368275_at	sterol-C4-methyl oxidase-like	NM_080886	2.24	4	209146_at	AV704962	2.23	8	1459627_at	BG070057	3.978	17
SCFD1	C	1368158_at	sec1 family domain containing 1	NM_019364	2.59	6	215548_s_at	AB020724	1.99	14	1428335_a_at	AK014070	1.852	15
SCG2	B	1368044_at	secretogranin 2	NM_022669	67.3	4	204035_at	NM_003469	29	14	1450708_at	NM_009129	22.92	17
SCG3	B	1368656_at	secretogranin III	NM_053856	31.6	4	219196_at	NM_013243	53.4	14	1448628_at	NM_009130	14.54	17
SCG5	B	1367992_at	secretory granule neuroendocrine protein 1	NM_013175	19.4	4	203889_at	NM_003020	24.8	14	1423150_at	AK019337	6.412	15
SCN3A	A	1369691_at	sodium channel, voltage-gated, type III, alpha polypeptide /// hypotheti	NM_013119	2.31	4	210432_s_at	AF225986	2.48	8	1439204_at	BB096886	2.978	13
SEC22B		1389138_at	SEC22 vesicle trafficking protein-like 1 (S. cerevisiae)	AA945574	2.51	6	209207_s_at	BC001364	1.83	14	1449063_at	BC009024	2.326	17
SEC22B		1389138_at	SEC22 vesicle trafficking protein-like 1 (S. cerevisiae)	AA945574	2.51	6	209206_at	AV701283	1.56	12	1449063_at	BC009024	2.326	17
SEC24D	C	1373610_at	SEC24 related gene family, member D (S. cerevisiae) (predicted)	AI232694	6.26	6	202375_at	NM_014822	4.46	14	1426972_at	AK009425	4.549	17
SEC61B	C	1382615_at	Sec61 alpha 1 subunit (S. cerevisiae)	BI284366	2.67	6	217716_s_at	NM_013336	1.53	12	1416190_a_at	BC003707	7.606	15
SEC61B	C	1382615_at	Sec61 alpha 1 subunit (S. cerevisiae)	BI284366	2.67	6	217716_s_at	NM_013336	1.53	12	1416191_at	BC003707	5.271	15
SEC61B	C	1371368_at	Sec61 alpha 1 subunit (S. cerevisiae)	BI279499	3.81	6	217716_s_at	NM_013336	1.53	12	1434986_a_at	AI327023	1.852	15
SEC61B	C	1382615_at	Sec61 alpha 1 subunit (S. cerevisiae)	BI284366	2.67	6	217716_s_at	NM_013336	1.53	12	1416189_a_at	BC003707	2.089	15
SEC61B	C	1388519_at	Sec61 beta subunit (predicted)	BI281906	2.66	6	203133_at	NM_006808	1.54	14	1417083_at	NM_024171	1.563	15
SEC63	C	1390115_at	SEC63-like (S. cerevisiae) (predicted)	BM384667	1.86	6	201916_s_at	NM_007214	1.73	12	1424926_at	C76103	4.32	17
SEC63	C	1390115_at	SEC63-like (S. cerevisiae) (predicted)	BM384667	1.86	6	201914_s_at	AK001465	1.59	12	1424925_at	C76103	2.988	17
SEC63	C	1390115_at	SEC63-like (S. cerevisiae) (predicted)	BM384667	1.86	6	201914_s_at	AK001465	1.59	12	1424924_at	C76103	1.778	13
SEPT2	C	1388389_at	septin 2	BF282620	1.54	4	200015_s_at	NM_004404	1.32	10	1423473_at	AV304911	1.705	17
SEPT2	C	1368984_at	septin 2	NM_057148	1.92	6	200778_s_at	AI191427	1.61	14	1423473_at	AV304911	1.705	17
SEZ6L2		1374825_at	seizure related 6 homolog (mouse)-like 2 (predicted)	BG380338	2.24	6	218720_x_at	NM_012410	1.24	8	1434641_x_at	AW121511	6.872	15
SEZ6L2		1374825_at	seizure related 6 homolog (mouse)-like 2 (predicted)	BG380338	2.24	6	218720_x_at	NM_012410	1.24	8	1423983_at	BC011475	5.546	15
SFRS10	C	1370189_at	splicing factor, arginine/serine-rich 10 (transformer 2 homolog, Drosop	BI290979	1.66	4	200892_s_at	BC000451	1.21	8	1446496_at	BB219075	1.782	17

SFRS10	C	1370188_at	splicing factor, arginine/serine-rich 10 (transformer 2 homolog, Drosop	BI290979	1.79	4	200892_s_at	BC000451	1.21	8	1441466_at	BG066615	1.508	15
SLC18A1	A	1387999_at	solute carrier family 18 (vesicular monoamine), member 1	M97380	1.4	6	207074_s_at	NM_003053	1.17	10	1458777_at	BG068849	7.459	17
SLC2A2		1387228_at	solute carrier family 2 (facilitated glucose transporter), member 2	NM_012879	28.6	6	206535_at	NM_000340	5.47	8	1449067_at	NM_031197	4.311	13
SLC35B1	B	1377143_at	solute carrier family 35, member B1	BM388792	2.42	6	202433_at	NM_005827	1.45	14	1448769_at	NM_016972	2.512	17
SLC7A2		1369460_at	solute carrier family 7 (cationic amino acid transporter, y+ system), me	NM_022619	2.35	4	207626_s_at	NM_003046	1.62	12	1426008_a_at	M62838	3.764	15
SLC7A2		1369460_at	solute carrier family 7 (cationic amino acid transporter, y+ system), me	NM_022619	2.35	4	207626_s_at	NM_003046	1.62	12	1422648_at	BF533509	3.524	15
SLC7A2		1369460_at	solute carrier family 7 (cationic amino acid transporter, y+ system), me	NM_022619	2.35	4	207626_s_at	NM_003046	1.62	12	1440506_at	BB433596	1.831	15
SLC7A8	B	1387057_at	solute carrier family 7 (cationic amino acid transporter, y+ system), me	NM_053442	4.59	4	217248_s_at	AL365343	2.56	12	1417929_at	NM_016972	2.66	17
SLC7A8	B	1387057_at	solute carrier family 7 (cationic amino acid transporter, y+ system), me	NM_053442	4.59	4	202752_x_at	NM_012244	1.79	12	1417929_at	NM_016972	2.66	17
SLC7A8	B	1387057_at	solute carrier family 7 (cationic amino acid transporter, y+ system), me	NM_053442	4.59	4	216092_s_at	AL365347	1.65	12	1417929_at	NM_016972	2.66	17
SMARCA2	C	1376337_at	SWI/SNF related, matrix associated, actin dependent regulator of chro	AI170385	1.59	4	206542_s_at	AV725365	2.29	10	1452333_at	BM230202	1.781	17
SMARCA2	C	1376337_at	SWI/SNF related, matrix associated, actin dependent regulator of chro	AI170385	1.59	4	206544_x_at	NM_003070	2.27	12	1452333_at	BM230202	1.781	17
SMARCA2	C	1376337_at	SWI/SNF related, matrix associated, actin dependent regulator of chro	AI170385	1.59	4	217707_x_at	AI535683	1.58	10	1452333_at	BM230202	1.781	17
SND1	C	1367790_at	staphylococcal nuclease domain containing 1	NM_022694	2.05	6	201622_at	NM_014390	1.72	14	1437036_at	BB240087	5.873	17
SND1	C	1367790_at	staphylococcal nuclease domain containing 1	NM_022694	2.05	6	201622_at	NM_014390	1.72	14	1437037_x_at	BB240087	3.261	17
SND1	C	1367790_at	staphylococcal nuclease domain containing 1	NM_022694	2.05	6	201622_at	NM_014390	1.72	14	1416038_at	NM_019776	1.729	13
SNRPN III	A	1368098_a_at	small nuclear ribonucleoprotein N	M29294	1.56	2	206042_x_at	NM_022804	3.04	10	1415896_x_at		3.635	17
SNRPN III	A	1368098_a_at	small nuclear ribonucleoprotein N	M29294	1.56	2	221974_at	AW770748	2.61	10	1435716_x_at		1.064	9
SOCS2		1369577_at	suppressor of cytokine signaling 2	NM_058208	2.42	6	203372_s_at	AB004903	2.13	12	1449109_at	NM_007706	1.914	13
SOCS2		1369577_at	suppressor of cytokine signaling 2	NM_058208	2.42	6	203373_x_at	NM_003877	1.47	12	1449109_at	NM_007706	1.914	13
SORL1	C	1393933_at	sortilin-related receptor, L(DLR class) A repeats-containing (predicted)	AW144823	4.02	4	212560_at	AV728268	3.99	10	1445889_at	BE980129	1.997	17
SORL1	C	1377457_a_at	sortilin-related receptor, L(DLR class) A repeats-containing (predicted)	AA850618	3.38	4	212560_at	AV728268	3.99	10	1460390_at	BI648081	1.512	15
SORL1	C	1386269_x_at	sortilin-related receptor, L(DLR class) A repeats-containing (predicted)	AI177589	3.36	4	212560_at	AV728268	3.99	10	1460390_at	BI648081	1.512	15
SORL1	C	1390710_x_at	sortilin-related receptor, L(DLR class) A repeats-containing (predicted)	AA850618	2.02	4	212560_at	AV728268	3.99	10	1460390_at	BI648081	1.512	15
SORL1	C	1377458_at	sortilin-related receptor, L(DLR class) A repeats-containing (predicted)	AA850618	1.94	4	212560_at	AV728268	3.99	10	1460390_at	BI648081	1.512	15
SQLE		1387017_at	squalene epoxidase	NM_017136	1.81	-2	209218_at	AF098865	1.84	10	1415993_at	NM_009270	2.336	15
SQLE		1387017_at	squalene epoxidase	NM_017136	1.81	-2	213562_s_at	BF979497	1.49	8	1415993_at	NM_009270	2.336	15
SQSTM1	C	1375374_at	Sequestosome 1	BF400606	3	4	201471_s_at	NM_003900	2.23	14	1444021_at	AW741388	3.21	17
SQSTM1	C	1389381_at	sequestosome 1	AI411586	2.56	4	213112_s_at	N30649	1.33	14	1444021_at	AW741388	3.21	17
SQSTM1	C	1371353_at	sequestosome 1	BG663093	1.56	4	201471_s_at	NM_003900	2.23	14	1444021_at	AW741388	3.21	17
SRP72	C	1375552_at	signal recognition particle,72 kDa subunit	BG372976	3.25	6	208095_s_at	NM_001222	2.01	14	1428876_at	BM942773	2.247	17
SRP72	C	1375552_at	signal recognition particle,72 kDa subunit	BG372976	3.25	6	208800_at	AV702627	1.54	12	1428876_at	BM942773	2.247	17
SRP72	C	1375552_at	signal recognition particle,72 kDa subunit	BG372976	3.25	6	208801_at	BE856385	1.52	12	1428876_at	BM942773	2.247	17
SRP72	C	1375552_at	signal recognition particle,72 kDa subunit	BG372976	3.25	6	208802_at	AI493872	1.19	10	1428877_at	BM942773	1.931	17
SRP9	C	1372445_at	Signal recognition particle 9 kDa protein (SRP9)	AI171783	2.25	6	201273_s_at	NM_003133	1.3	10	1448753_at	BI661964	1.878	15
SRPR	C	1371343_at	signal recognition particle receptor ('docking protein') (predicted)	BM391232	2.71	6	200918_s_at	NM_003139	2.02	14	1423670_a_at	BC021839	3.12	17
SRPR	C	1371343_at	signal recognition particle receptor ('docking protein') (predicted)	BM391232	2.71	6	200917_s_at	BG474541	1.28	10	1423670_a_at	BC021839	3.12	17
SSR1	C	1373090_at	signal sequence receptor, alpha (predicted)	AI230697	2.5	6	200889_s_at	AI016620	2.67	14	1441327_a_at	AV325051	3.343	17
SSR1	C	1373090_at	signal sequence receptor, alpha (predicted)	AI230697	2.5	6	200890_s_at	AW006345	2.09	14	1441327_a_at	AV325051	3.343	17
SSR1	C	1373090_at	signal sequence receptor, alpha (predicted)	AI230697	2.5	6	200891_s_at	NM_003144	2.12	14	1448843_at	BG077348	2.228	13
SSR2	C	1373185_at	signal sequence receptor, beta (predicted)	BE329347	1.94	4	200652_at	NM_003145	1.49	8	1449930_a_at	NM_025448	1.555	13
SSR4	C	1367690_at	signal sequence receptor 4	NM_017199	2.49	6	201004_at	NM_006280	2.84	14	1427096_s_at	AV051818	2.458	17
SSTR2	A	1371174_s_at	somatostatin receptor 2	X98234	1.4	4	217455_s_at	AF184174	1.97	12	1422256_at	NM_009217	1.947	15
ST18	B	1370658_a_at	suppression of tumorigenicity 18	AF031942	4.87	6	206135_at	NM_014682	11.7	12	1455123_at	BB178719	22.07	17
ST18	B	1370658_a_at	suppression of tumorigenicity 18	AF031942	4.87	6	206135_at	NM_014682	11.7	12	1436793_at	AV347235	2.785	17
STAU2	A	1390558_at	staufen, RNA binding protein, homolog 2 (Drosophila)	AI502229	2.63	4	204226_at	NM_014393	1.27	8	1425533_a_at	AJ244015	2.576	17
STAU2	A	1390558_at	staufen, RNA binding protein, homolog 2 (Drosophila)	AI502229	2.63	4	204226_at	NM_014393	1.27	8	1450916_at	AI843206	2.557	15
STCH	C	1372368_at	Stress 70 protein chaperone, microsomal-associated, 60kD human hor	AI230431	3.57	6	202558_s_at	NM_006948	2.72	14	1453172_at	BE533039	17.34	17
STCH	C	1368844_at	stress 70 protein chaperone, microsomal-associated, 60kD human hor	NM_019271	2.42	6	202557_at	AI718418	2.62	12	1430026_at	AK021006	8.187	17
STCH	C	1368844_at	stress 70 protein chaperone, microsomal-associated, 60kD human hor	NM_019271	2.42	6	202557_at	AI718418	2.62	12	1433772_at	BE650268	2.426	15

STCH	C	1368844_at	stress 70 protein chaperone, microsome-associated, 60kD human hom	NM_019271	2.42	6	202557_at	AI718418	2.62	12	1429502_at	BE533039	2.156	17
STXBP1	A	1370518_a_at	syntaxin binding protein 1	U06069	4.21	4	202260_s_at	NM_003165	5.62	8	1420506_a_at	AF326545	9.613	17
STXBP1	A	1398330_at	syntaxin binding protein 1	BM388715	2.12	4	202260_s_at	NM_003165	5.62	8	1420506_a_at	AF326545	9.613	17
STXBP1	A	1370519_at	syntaxin binding protein 1	U06069	2.74	4	202260_s_at	NM_003165	5.62	8	1420505_a_at	AF326545	2.406	13
SYP	A	1368276_at	synaptophysin	NM_012664	1.75	2	213200_at	U93305	2.59	8	1448280_at	NM_009305	1.691	13
TAF9B	B	1368429_at	TAF9-like RNA polymerase II, TATA box binding protein (TBP)-associa	NM_133615	1.49	2	221618_s_at	AF220509	1.44	12	1434960_at	AW555571	2.848	17
TAF9B	B	1368429_at	TAF9-like RNA polymerase II, TATA box binding protein (TBP)-associa	NM_133615	1.49	2	221618_s_at	AF220509	1.44	12	1447829_x_at	BB431174	2.221	17
TARS	C	1372145_at	threonyl-tRNA synthetase	BI289683	1.59	2	201263_at	NM_003191	2.24	12	1460323_at	AK010256	1.944	17
TCTA		1372507_at	T-cell leukemia translocation altered gene	AA901093	1.57	6	203054_s_at	NM_022171	1.33	14	1420123_at	C85065	2.286	17
TERF2IP		1388780_at	telomeric repeat binding factor 2, interacting protein (predicted)	BF395792	2.17	6	201174_s_at	NM_018975	2.12	8	1419381_at	NM_020584	1.675	13
TGOLN2		1371028_at	trans-golgi network protein 2	AW917192	2.32	6	212040_at	BG249599	1.94	14	1423307_s_at	AI314055	1.617	15
TGOLN2		1389965_at	trans-golgi network protein 2	AA799818	1.63	4	203833_s_at	BF061845	1.65	14	1423307_s_at	AI314055	1.617	15
TGOLN2		1387755_s_at	trans-golgi network protein 2	NM_138840	1.63	6	212043_at	W72053	1.57	12	1423307_s_at	AI314055	1.617	15
TGOLN2		1389965_at	trans-golgi network protein 2	AA799818	1.63	4	203834_s_at	NM_006464	1.23	10	1423307_s_at	AI314055	1.617	15
TLE1		1374425_at	transducin-like enhancer of split 1, homolog of Drosophila E(spl) (predi	BG380534	2.09	4	203220_s_at	AI951720	1.22	10	1422751_at	NM_011599	1.616	11
TLOC1	C	1374188_at	translocation protein 1 (predicted)	AI103194	1.47	6	208942_s_at	BE866511	1.79	12	1455698_at	AK018223	1.495	7
TLOC1	C	1374188_at	translocation protein 1 (predicted)	AI103194	1.47	6	208943_s_at	U93239	1.59	14	1455698_at	BQ175470	1.495	7
TM4SF4	B	1368706_at	transmembrane 4 superfamily member 4	NM_053785	11.6	6	209937_at	BC001386	15.8	14	1424962_at	BC010814	9.872	17
TM9SF4		1371647_at	transmembrane protein TM9SF3 (predicted)	AA963367	1.58	6	212194_s_at	AI418892	1.53	14	1423204_at	BB367487	7.14	17
TMOD2		1369541_at	tropomodulin 2	NM_031613	1.66	4	219701_at	NM_014548	1.85	10	1430153_at	AK018223	11.38	17
TMOD2		1390482_at	Tropomodulin 2	BF285511	2.06	6	219701_at	NM_014548	1.85	10	1431326_a_at	AK017072	5.517	15
TMOD2		1393418_at	tropomodulin 2	BF567833	1.93	4	219701_at	NM_014548	1.85	10	1435060_at	BB667124	3.117	15
TPD52	B	1391880_at	Tumor protein D52 (predicted)	AA893670	1.65	2	201688_s_at	BG389015	5.99	14	1419493_a_at	BC002036	1.967	15
TPD52	B	1391880_at	Tumor protein D52 (predicted)	AA893670	1.65	2	201689_s_at	BE974098	3.95	14	1419493_a_at	BC002036	1.967	15
TPD52	B	1391880_at	Tumor protein D52 (predicted)	AA893670	1.65	2	201690_s_at	AA524023	3.92	14	1419493_a_at	BC002036	1.967	15
TPD52	B	1391880_at	Tumor protein D52 (predicted)	AA893670	1.65	2	201691_s_at	NM_005079	2.04	14	1419493_a_at	BC002036	1.967	15
TRIB1	C	1389987_at	tribbles homolog 1 (Drosophila)	AI511059	4.96	6	202241_at	NM_025195	3.48	10	1424881_at	AV237242	3.652	15
TRIB1	C	1371019_at	tribbles homolog 1 (Drosophila)	BM387324	2.08	4	202241_at	NM_025195	3.48	10	1424880_at	AV237242	4.079	15
TSN		1367728_at	translin	NM_021762	1.88	6	201513_at	AI659180	1.62	12	1448516_at	BB818311	2.941	17
TSN		1367728_at	translin	NM_021762	1.88	6	201504_s_at	AI435302	1.44	10	1448516_at	BB818311	2.941	17
TSN		1367728_at	translin	NM_021762	1.88	6	201515_s_at	NM_004622	1.35	12	1448516_at	BB818311	2.941	17
TSPYL4	A	1372381_at	TSPY-like 4	BM383457	1.84	0	212928_at	AL050331	1.69	8	1424029_at	BC017540	1.556	13
TSPYL4	A	1390097_at	TSPY-like 4 (predicted)	BI281738	1.92	0	212928_at	AL050331	1.69	8	1424029_at	BC017540	1.556	13
TXNDC1	C	1372067_at	thioredoxin domain containing 1 (predicted)	AI179567	1.91	2	208097_s_at	NM_030755	2.45	14	1437143_a_at	AV310544	2.14	15
TXNDC1	C	1372067_at	thioredoxin domain containing 1 (predicted)	AI179567	1.91	2	209476_at	AL080080	1.68	14	1437143_a_at	AV310544	2.14	15
TXNDC1	C	1372298_at	thioredoxin domain containing 4 (endoplasmic reticulum) (predicted)	BE098766	2.43	6	208959_s_at	BC005374	1.96	14	1423246_at	BI100077	3.851	17
TXNL1	C	1388862_at	Thioredoxin-like (32kD)	AI412143	1.47	4	201588_at	NM_004786	1.3	10	1435235_at	BI662855	2.469	17
TXNL1	C	1368936_at	thioredoxin-like (32kD)	NM_080887	1.44	6	201588_at	NM_004786	1.3	10	1435235_at	BI662855	2.469	17
TXNL1	C	1380191_s_at	thioredoxin-like (32kD)	BF555110	1.37	6	201588_at	NM_004786	1.3	10	1435235_at	BI662855	2.469	17
TXNL1	C	1391789_at	thioredoxin-like (32kD)	BF555110	1.89	6	201588_at	NM_004786	1.3	10	1437905_at	AV106191	1.883	13
UBL3	B	1374413_at	ubiquitin-like 3 (predicted)	BM386522	1.8	4	201534_s_at	AF044221	2.11	14	1423461_a_at	AV328436	1.907	15
UBL3	B	1374413_at	ubiquitin-like 3 (predicted)	BM386522	1.8	4	201535_at	NM_007106	1.48	8	1423461_a_at	AV328436	1.907	15
UBQLN2	C	1372131_at	ubiquilin 2 (predicted)	BF398140	1.57	2	215884_s_at	AK001029	1.86	10	1420844_at	AV171029	2.878	17
UBXD2		1388444_at	UBX domain containing 2 (predicted)	BI291633	1.81	6	212006_at	AU149908	2.07	14	1426485_at	AI788596	7.221	17
UBXD2		1388444_at	UBX domain containing 2 (predicted)	BI291633	1.81	6	212008_at	N29889	2.29	14	1444057_at	BB253461	1.784	13
UFC1	C	1371758_at	Ufm1-conjugating enzyme 1	BM986259	1.69	6	217797_at	NM_016406	1.73	14	1416327_at	NM_025388	2.16	17
USP47	C	1399036_at	ubiquitin specific protease 47 (predicted)	AA964374	1.53	4	221518_s_at	BE966019	1.21	8	1426976_at	BC069271	1.501	11
VAT1		1374905_at	Vesicle amine transport protein 1 homolog (T californica)	AA925303	1.57	6	208626_s_at	BC001913	1.82	14	1438165_x_at	BB559097	2.079	13
VLDLR		1369098_at	very low density lipoprotein receptor	NM_013155	2.13	4	209822_s_at	L22431	4.46	14	1438258_at	BE647363	1.907	11
VLDLR		1387455_a_at	very low density lipoprotein receptor	NM_013155	2.02	4	209822_s_at	L22431	4.46	14	1438258_at	BE647363	1.907	11

WDR6	C	1373399_at	WD repeat domain 6 (predicted)	BI291997	2.82	4	217734_s_at	NM_018031	1.62	10	1455940_x_at	BB453609	2.105	17
WFS1		1387356_at	Wolfram syndrome 1	NM_031823	3.45	6	202908_at	NM_006005	2.27	14	1448411_at	NM_011716	4.078	17
WFS1		1368839_at	Wolfram syndrome 1	NM_031823	3	6	202908_at	NM_006005	2.27	14	1448411_at	NM_011716	4.078	17
WNT4	B	1368641_at	wingless-related MMTV integration site 4	NM_053402	7.47	4	208606_s_at	NM_030761	15.1	14	1450782_at	NM_009523	3.741	15
YWHAQ	C	1370168_at	tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation pr	BF281342	1.72	-2	200693_at	NM_006826	1.75	10	1437608_x_at	BB414446	3.509	17
YWHAQ	C	1387862_at	tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation pr	BF281342	1.58	-2	212426_s_at	BF033313	1.95	10	1445341_at	BG074147	1.971	15
ZNF451	C	1389047_at	Zinc finger protein 451	AI009167	2.17	4	212557_at	AB011148	1.42	10	1456415_at	BI083675	2.433	17
ZNF451	C	1389047_at	Zinc finger protein 451	AI009167	2.17	4	212557_at	AB011148	1.42	10	1456350_at	BI174076	1.683	15

Islet-abundant and , in rat tissue data set, islet endocrine non-beta (alpha cell) -enriched

symbol	probe set	gene (Rat)	Accession	av FC	FCeta sco	probe set	Accession	av FC	FCeta sco	probe set	Accession	av FC	FCeta sco
ANXA4	1368908_at	annexin A4	NM_024155	1.45	4	201301_s_at	BC000182	3.74	10	1421223_a_at	NM_013471	1.689	9
ANXA4	1389305_at	annexin A4	BM385237	1.34	2	201302_at	NM_001153	3.69	8	1421223_a_at	NM_013471	1.689	9
ALCAM	1370043_at	activated leukocyte cell adhesion molecule	NM_031753	1.55	-2	201951_at	BF242905	9.6	14	1426301_at	U95030	2.038	15
ALCAM	1375263_at	Activated leukocyte cell adhesion molecule	BI296498	1.41	4	201952_at	AA156721	5.39	14	1426301_at	U95030	2.038	15
ABCC8	1369632_a_at	ATP-binding cassette, sub-family C (CFTR/MRP), member 8	AB052294	2.69	4	210246_s_at	AF087138	22.3	14	1457066_at	BF466569	9.954	17
ABCC8	1369632_a_at	ATP-binding cassette, sub-family C (CFTR/MRP), member 8	AB052294	2.69	4	210246_s_at	AF087138	22.3	14	1455765_a_at	BB515948	3.333	17
BTG3	1368072_at	B-cell translocation gene 3	NM_019290	5.1	4	213134_x_at	AI765445	6.48	14	1449007_at	NM_009770	2.914	17
BTG3	1368072_at	B-cell translocation gene 3	NM_019290	5.1	4	205548_s_at	NM_006806	5.44	14	1449007_at	NM_009770	2.914	17
BCLAF1	1373868_at	BCL2-associated transcription factor 1 (predicted)	AI717547	1.8	4	214499_s_at	AF249273	2.05	8	1438089_a_at	BE853331	3.675	17
BCLAF1	1373868_at	BCL2-associated transcription factor 1 (predicted)	AI717547	1.8	4	201083_s_at	AA740754	1.79	12	1438089_a_at	BE853331	3.675	17
BCLAF1	1373868_at	BCL2-associated transcription factor 1 (predicted)	AI717547	1.8	4	201084_s_at	NM_014739	1.43	12	1438089_a_at	BE853331	3.675	17
MTHFD2	1372808_at	Bifunctional methylenetetrahydrofolate dehydrogenase/cyclohydrolase	AW251324	2.75	4	201761_at	NM_006636	2.44	10	1419254_at	BG076333	2.997	17
MTHFD2	1372808_at	Bifunctional methylenetetrahydrofolate dehydrogenase/cyclohydrolase	AW251324	2.75	4	201761_at	NM_006636	2.44	10	1419253_at	BG076333	2.726	17
MTHFD2	1372808_at	Bifunctional methylenetetrahydrofolate dehydrogenase/cyclohydrolase	AW251324	2.75	4	201761_at	NM_006636	2.44	10	1436704_x_at	AV215673	1.748	13
CDH1	1386947_at	cadherin 1	NM_031334	8.43	4	201131_s_at	NM_004360	9.41	8	1448261_at	NM_009864	1.876	13
CDH1	1386947_at	cadherin 1	NM_031334	8.43	4	201130_s_at	L08599	5	12	1448261_at	NM_009864	1.876	13
CPD	1369215_a_at	carboxypeptidase D	NM_012836	1.4	4	201942_s_at	D85390	4.1	12	1418018_at	NM_007754	3.633	17
CPD	1369215_a_at	carboxypeptidase D	NM_012836	1.4	4	201940_at	AA897514	4.05	14	1418018_at	NM_007754	3.633	17
CPD	1369215_a_at	carboxypeptidase D	NM_012836	1.4	4	201941_at	BE349147	3.88	14	1418018_at	NM_007754	3.633	17
CPD	1369215_a_at	carboxypeptidase D	NM_012836	1.4	4	201943_s_at	NM_001304	4.61	14	1447392_s_at	BB710395	2.409	15
CBX6	1374703_at	Chromobox homolog 6	BI304209	1.48	0	202047_s_at	AI458128	2.33	10	1424407_s_at	BC019942	1.524	7
CBX6	1375307_at	chromobox homolog 6 (predicted)	BI275772	1.62	0	202048_s_at	NM_014292	1.74	10	1424407_s_at	BC019942	1.524	7
CHGB	1368034_at	chromogranin B	NM_012526	13.9	2	204260_at	NM_001819	31.8	14	1415885_at	NM_007694	10.9	17
DAP	1369941_at	death-associated protein	NM_022526	2.15	2	201095_at	NM_004394	2.85	14	1451112_s_at	BC024876	1.964	15
DPP6	1376183_at	Dipeptidylpeptidase 6	AI101484	3.21	0	207789_s_at	NM_001936	2.94	10	1439748_at	AV328546	5.194	17
DPP6	1369390_a_at	dipeptidylpeptidase 6	NM_022850	2.57	2	207789_s_at	NM_001936	2.94	10	1439748_at	AV328546	5.194	17
DUSP8	1372385_at	dual specificity phosphatase 8 (predicted)	AA957292	1.26	4	206374_at	NM_004420	1.42	8	1418714_at	NM_008748	2.079	13
DYRK3	1383649_a_at	Dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 3	BI291080	1.92	2	210151_s_at	AF186773	1.42	12	1424229_at	BC006704	2.015	15
EGR1	1368321_at	early growth response 1	NM_012551	4.86	4	201694_s_at	NM_001964	4.72	14	1417065_at	NM_007913	4.923	17
EGR1	1368321_at	early growth response 1	NM_012551	4.86	4	201693_s_at	AV733950	2.23	14	1417065_at	NM_007913	4.923	17
EPM2AIP1	1372568_at	EPM2A (laforin) interacting protein 1 (predicted)	AI411270	1.73	4	202909_at	NM_014805	1.58	10	1434106_at	AV340515	4.85	17
EPM2AIP1	1372568_at	EPM2A (laforin) interacting protein 1 (predicted)	AI411270	1.73	4	202909_at	NM_014805	1.58	10	1425624_at	BC018474	4.455	15
EPM2AIP1	1389100_at	EPM2A (laforin) interacting protein 1 (predicted)	BF409831	2.76	4	202909_at	NM_014805	1.58	10	1454835_at	AV340515	1.908	15
EPM2AIP1	1372568_at	EPM2A (laforin) interacting protein 1 (predicted)	AI411270	1.73	4	202909_at	NM_014805	1.58	10	1434105_at	AV340515	1.766	13
GCG	1369888_at	glucagon	NM_012707	39.6	4	206422_at	NM_002054	60	14	1425952_a_at	AF276754	47.58	17
GNAS	1387906_a_at	GNAS complex locus	AF107845	8.34	2	217057_s_at	AF107846	26.1	14	1443007_at	AW545941	1.858	13
GNAS	1387906_a_at	GNAS complex locus	AF107845	8.34	2	214157_at	AA401492	14.9	14	1427789_s_at	AI561892	4.001	17

GNAS	1387906_a_at	GNAS complex locus	AF107845	8.34	2	211858_x_at	AF088184	2.99	14	1427789_s_at	AI561892	4.001	17
GNAS	1387906_a_at	GNAS complex locus	AF107845	8.34	2	200981_x_at	NM_016592	2.9	14	1427789_s_at	AI561892	4.001	17
GNAS	1387906_a_at	GNAS complex locus	AF107845	8.34	2	212273_x_at	AI591100	2.84	14	1427789_s_at	AI561892	4.001	17
GNAS	1387906_a_at	GNAS complex locus	AF107845	8.34	2	200780_x_at	NM_000516	2.83	14	1427789_s_at	AI561892	4.001	17
GNAS	1387906_a_at	GNAS complex locus	AF107845	8.34	2	214548_x_at	AF064092	2.75	14	1443375_at	BM239335	1.976	17
GNAS	1369897_s_at	GNAS complex locus /// XLas protein	BI277035	1.4	2	217673_x_at	AA650558	2.29	14	1444767_at	AV253089	2.531	17
GCH1	1387221_at	GTP cyclohydrolase 1	NM_024356	5.85	4	204224_s_at	NM_000161	18.8	14	1429692_s_at	BB698398	9.306	17
GCH1	1373513_at	GTP cyclohydrolase 1	AI012419	6.23	4	204224_s_at	NM_000161	18.8	14	1420499_at	NM_008102	3.857	17
GCH1	1380230_at	GTP cyclohydrolase 1	AI639457	4.62	4	204224_s_at	NM_000161	18.8	14	1420499_at	NM_008102	3.857	17
GCH1	1368503_at	GTP cyclohydrolase 1	NM_024356	1.89	4	204224_s_at	NM_000161	18.8	14	1420499_at	NM_008102	3.857	17
GNAI1	1387505_at	guanine nucleotide binding protein, alpha inhibiting 1	NM_013145	1.41	0	209576_at	AL049933	3.43	12	1427510_at	U38501	2.084	17
GNAI1	1387505_at	guanine nucleotide binding protein, alpha inhibiting 1	NM_013145	1.41	0	209576_at	AL049933	3.43	12	1427510_at	U38501	2.084	17
IER2	1372389_at	immediate early response 2	BF420059	2.06	4	202081_at	NM_004907	3.84	12	1416442_at	NM_010499	3.838	15
IGSF1	1379252_at	Immunoglobulin superfamily, member 4A	AW522833	1.42	-2	209030_s_at	NM_014333	4.28	14	1440412_at	BB312687	1.857	15
IGSF1	1379252_at	Immunoglobulin superfamily, member 4A	AW522833	1.42	-2	209031_at	AL519710	3.59	12	1440412_at	BB312687	1.857	15
IGSF1	1379252_at	Immunoglobulin superfamily, member 4A	AW522833	1.42	-2	209032_s_at	AF132811	2.4	14	1440412_at	BB312687	1.857	15
IRS2	1371091_at	insulin receptor substrate 2	not on RG230A array			209184_s_at	BF700086	2.44	14	1443969_at	BE199054	10.08	17
IRS2	1397999_at	Insulin receptor substrate 2	not on RG230A array			209185_s_at	AF073310	1.51	12	1443969_at	BE199054	10.08	17
ISL1	1369681_at	ISL1 transcription factor, LIM/homeodomain 1	NM_017339	1.5	2	206104_at	NM_002202	26.9	14	1450723_at	BQ176915	20.03	17
ISL1	1369681_at	ISL1 transcription factor, LIM/homeodomain 1	NM_017339	1.5	2	206104_at	NM_002202	26.9	14	1422720_at	BQ176915	8.868	17
JUNB	1387788_at	Jun-B oncogene	NM_021836	2	4	201473_at	NM_002229	3.05	12	1415899_at	NM_008416	2.173	15
NDN	1371947_at	necdin (predicted)	BG671865	1.7	0	209550_at	U35139	1.82	12	1437853_x_at	BB074430	1.908	13
NEO1	1390468_at	Neogenin	BF397709	1.42	2	204321_at	NM_002499	1.59	12	1447693_s_at	BB350308	2.818	17
NFIC	1371176_at	nuclear factor I/C	AB012233	1.35	4	213298_at	X12492	1.05	10	1450661_x_at	NM_008688	2.342	17
NR4A2	1387410_at	nuclear receptor subfamily 4, group A, member 2	U72345	1.5	4	216248_s_at	S77154	3.27	10	1455034_at	BB703394	2.905	15
NR4A2	1387410_at	nuclear receptor subfamily 4, group A, member 2	U72345	1.5	4	204622_x_at	NM_006186	2.93	10	1450749_a_at	NM_013613	2.527	15
NR4A2	1387410_at	nuclear receptor subfamily 4, group A, member 2	U72345	1.5	4	204621_s_at	AI935096	2.17	12	1447863_s_at	BB322941	1.991	17
NOL5A	1368173_at	nucleolar protein 5	NM_021754	1.78	2	200874_s_at	BE796327	1.57	14	1450986_at	BB729616	2.348	15
PPY	1369196_at	pancreatic polypeptide	NM_012626	4.39	4	210670_at	M15788	42.4	14	1420440_at	NM_008918	78.13	17
PGBD5	1389597_at	piggyBac transposable element derived 5 (predicted)	BI293445	1.35	0	219225_at	NM_024554	1.96	8	1460570_at	AV151597	1.918	15
PCSK1N	1367835_at	proprotein convertase subtilisin/kexin type 1 inhibitor	NM_019279	9.6	2	218952_at	NM_013271	10.8	14	1416965_at	AF181560	5.882	15
RYBP	1372674_at	RING1 and YY1 binding protein (predicted)	BE107155	1.42	2	201845_s_at	AB029551	2.02	8	1428219_at	AK010548	2.064	15
RYBP	1372674_at	RING1 and YY1 binding protein (predicted)	BE107155	1.42	2	201846_s_at	NM_012234	1.46	8	1428219_at	AK010548	2.064	15
RYBP	1372674_at	RING1 and YY1 binding protein (predicted)	BE107155	1.42	2	201844_s_at	W84482	1.29	10	1428219_at	AK010548	2.064	15
SCN3B	1370850_at	sodium channel, voltage-gated, type III, beta	AA685184	1.75	0	204722_at	AW007335	1.28	10	1426328_a_at	AY049036	1.972	15
SLC7A1	1368391_at	solute carrier family 7 (cationic amino acid transporter, y+ system), me	AB066224	2.29	4	206566_at	NM_003045	1.69	14	1454992_at	BB264620	1.915	15
SLC7A1	1368392_at	solute carrier family 7 (cationic amino acid transporter, y+ system), me	AB066224	1.42	4	212295_s_at	AA148507	1.78	12	1454992_at	BB264620	1.915	15
TIPARP	1374446_at	TCDD-inducible poly(ADP-ribose) polymerase (predicted)	AI179464	1.49	4	212665_at	AL556438	2.52	12	1452160_at	BB707122	8.334	17
TIPARP	1374446_at	TCDD-inducible poly(ADP-ribose) polymerase (predicted)	AI179464	1.49	4	212665_at	AL556438	2.52	12	1452161_at	BB707122	2.728	13
TIPARP	1374446_at	TCDD-inducible poly(ADP-ribose) polymerase (predicted)	AI179464	1.49	4	212665_at	AL556438	2.52	12	1426721_s_at	BB707122	2.648	17
TRIB3	1386321_s_at	tribbles homolog 3 (Drosophila)	H31287	3.09	4	218145_at	NM_021158	1.62	12	1456225_x_at	BB508622	2.256	15
TRIB3	1370695_s_at	tribbles homolog 3 (Drosophila)	AB020967	2.41	4	218145_at	NM_021158	1.62	12	1456225_x_at	BB508622	2.256	15
TRIB3	1370694_at	tribbles homolog 3 (Drosophila)	AB020967	1.57	4	218145_at	NM_021158	1.62	12	1456225_x_at	BB508622	2.256	15
TNFRSF12A	1371785_at	tumor necrosis factor receptor superfamily, member 12a	BI303379	1.91	2	218368_s_at	NM_016639	1.51	12	1418572_x_at	NM_013749	4.485	17
TACSTD1	1388199_at	tumor-associated calcium signal transducer 1	BG376410	10.2	2	201839_s_at	NM_002354	10.7	12	1447899_x_at	AV099587	3.826	17
WBP11	1372575_at	WW domain binding protein 11 (predicted)	AA892364	1.67	4	217822_at	NM_016312	1.22	10	1447401_at	AW909375	1.908	15
ZBTB10	1376125_at	Zinc finger and BTB domain containing 10	BI289112	1.8	4	219312_s_at	NM_023929	2.16	10	1436067_at	BG068032	2.246	17
ZBTB10	1387453_at	zinc finger and BTB domain containing 10	NM_024489	1.88	4	219312_s_at	NM_023929	2.16	10	1436068_at	BG068032	1.62	17
ZKSCAN1	1373106_at	Zinc finger protein 36, C3H type-like 2	AI178069	1.47	2	214900_at	AC004522	1.18	12	1452519_a_at	X14678	1.701	15
ZKSCAN1	1389851_at	zinc finger protein 36, C3H type-like 2 (predicted)	AA943730	1.37	6	214670_at	AA653300	1.81	12	1452519_a_at	X14678	1.701	15

ZDHHC14	1372989_at	Zinc finger, DHHC domain containing 14	BI296586	1.39	2	219247_s_at	NM_024630	1.24	12	1438151_x_at	BB544336	11.38	17
ZDHHC14	1372989_at	Zinc finger, DHHC domain containing 14	BI296586	1.39	2	219247_s_at	NM_024630	1.24	12	1437614_x_at	AV223474	1.705	15
SST	1367762_at	somatostatin	NM_012659	17.1	4	213921_at	NM_001048	57.4	14	1417954_at	NM_009215	22.47	17

Islet-abundant and , in human tissue data set, pancreatic duct cell-enriched as compared to beta cell fraction

symbol	probe set	gene (Rat)	Accession	av FCeta sco	probe set	Accession	av FCeta sco	probe set	Accession	av FCeta sco			
BACE2	1377390_at	beta-site APP-cleaving enzyme 2	BG381587	1.74	4	217867_x_at	NM_012105	1.74	8	1435581_at	BE947462	7.756	17
BACE2	1377390_at	beta-site APP-cleaving enzyme 2	BG381587	1.74	4	217867_x_at	NM_012105	1.74	8	1437846_x_at	BB348062	7.746	17
BACE2	1377390_at	beta-site APP-cleaving enzyme 2	BG381587	1.74	4	217867_x_at	NM_012105	1.74	8	1416673_at	NM_019517	5.473	17
BACE2	1377390_at	beta-site APP-cleaving enzyme 2	BG381587	1.74	4	217867_x_at	NM_012105	1.74	8	1438645_x_at	BB558905	3.612	15
CXCL1	1387316_at	chemokine (C-X-C motif) ligand 1	NM_030845	6.38	6	204470_at	NM_001511	2.7	8	1457644_s_at	BB554288	14.79	17
CXCL1	1387316_at	chemokine (C-X-C motif) ligand 1	NM_030845	6.38	6	204470_at	NM_001511	2.7	8	1441855_x_at	BB554288	11.86	17
CXCL1	1387316_at	chemokine (C-X-C motif) ligand 1	NM_030845	6.38	6	204470_at	NM_001511	2.7	8	1419209_at	NM_008176	5.93	17
DUSP1	1368147_at	dual specificity phosphatase 1	BE110108	1.84	4	201044_x_at	AA530892	3.09	10	1448830_at	NM_013642	3.849	17
DUSP1	1368146_at	dual specificity phosphatase 1	BE110108	1.77	4	201041_s_at	NM_004417	2.16	10	1448830_at	NM_013642	3.849	17
KLF6	1387060_at	Kruppel-like factor 6 (core promoter element binding protein)	NM_031642	5.62	6	208961_s_at	AB017493	3.47	10	1418280_at	NM_011803	1.794	15
KLF6	1388986_at	Kruppel-like factor 6 (Core promoter element binding protein)	AI598339	3.76	4	208961_s_at	AB017493	3.47	10	1433508_at	AV025472	1.903	13
RIPK4	1389648_at	Receptor-interacting serine-threonine kinase 4 (predicted)	AI170382	3.17	4	221215_s_at	NM_020639	1.8	10	1418488_s_at	AF302127	3.269	17
RIPK4	1389648_at	Receptor-interacting serine-threonine kinase 4 (predicted)	AI170382	3.17	4	221215_s_at	NM_020639	1.8	10	1418487_at	AF302127	2.111	11