

RefSeq ID	ProbeSets	Gene Symbol	Gene Description	Entrez Gene ID	Sequence annotation	Seed matches location(s)	Ago-2 binding specific enrichment (replicate 1)	Ago-2 binding specific enrichment (replicate 2)	OE lysate log2 fold change (replicate 1)	OE lysate log2 fold change (replicate 2)	Probability	Pulled down in Karginov?
NM_005646	202813_at	TARBP1	Homo sapiens TAR (HIV-1) RNA binding protein 1 (TARBP1), mRNA.	6894	TR(1..5130)CDS(1..4866)	4868..4874,5006..5013	3.73	2.53	-1.54	-0.44	1 Yes	
NM_001665	203175_at	RHOG	Homo sapiens ras homolog gene family, member G (rho G) (RHOG), mRNA.	391	TR(1..1332)CDS(159..734)	810..817,782..788,790..796,873..879	3.56	2.78	-1.62	-1	1 Yes	
NM_002742	205880_at	PRKD1	Homo sapiens protein kinase D1 (PRKD1), mRNA.	5587	TR(1..3679)CDS(182..2920)	3538..3544,3202..3208	4.15	1.83	-2.55	-0.42	1 Yes	
NM_003068	213139_at	SNAI2	Homo sapiens snail homolog 2 (Drosophila) (SNAI2), mRNA.	6591	TR(1..2101)CDS(165..971)	1410..1417,1814..1820,1610..1616	3.5	2.79	-1.38	-0.31	1 Yes	
NM_006270	212647_at	RRAS	Homo sapiens related RAS viral (r-ras) oncogene homolog (RRAS), mRNA.	6237	TR(1..1013)CDS(46..702)	871..877	3.82	2.29	-1.54	-0.55	1 Yes	
NM_025188	219923_at,242056_at	TRIM45	Homo sapiens tripartite motif-containing 45 (TRIM45), mRNA.	80263	TR(1..3584)CDS(589..2331)	3408..3414,2437..2444,3425..3431,2781..2787	3.87	1.89	-0.62	-0.09	1 Yes	
NM_024684	221600_s_at,221599_at	C11orf67	Homo sapiens chromosome 11 open reading frame 67 (C11orf67), mRNA.	28971	TR(1..557)CDS(99..467)	327..333	4.35	3.54	-1.06	-0.56	1 Yes	
NM_031458	223220_s_at	PARP9	Homo sapiens poly (ADP-ribose) polymerase family, member 9 (PARP9), mRNA.	83666	TR(1..3243)CDS(229..2793)	2951..2957	4.16	2.03	-1.8	-0.36	1 No	
NM_001914	215726_s_at	CYB5A	Homo sapiens cytochrome b5 type A (microsomal) (CYB5A), transcript variant 2, mRNA.	1528	TR(1..844)CDS(112..408)	730..736	3.44	2.25	-0.99	-0.46	1 No	
NM_057179	229404_at	TWIST2	Homo sapiens twist homolog 2 (Drosophila) (TWIST2), mRNA.	117581	TR(1..1381)CDS(132..614)	1002..1008	3.4	2.51	-1.49	-0.84	1 Yes	
NM_014397	223158_s_at,223159_s_at	NEK6	Homo sapiens NIMA (never in mitosis gene a)-related kinase 6 (NEK6), mRNA.	10783	TR(1..2596)CDS(146..1087)	1127..1133,1101..1108	3.28	1.93	-0.7	-0.23	1 Yes	
NM_018441	221142_s_at	PECR	Homo sapiens peroxisomal trans-2-enoyl-CoA reductase (PECR), mRNA.	55825	TR(1..1863)CDS(69..980)	1012..1018,1772..1778	3.24	2.79	-0.54	-0.51	1 No	
NM_016361	218795_at	ACP6	Homo sapiens acid phosphatase 6, lysophosphatidic (ACP6), mRNA.	51205	TR(1..1832)CDS(465..1751)	1791..1797	3.22	2.35	-0.71	-0.44	1 No	
NM_178865	224762_at	SERINC2	Homo sapiens serine incorporator 2 (SERINC2), mRNA.	347735	TR(1..2011)CDS(151..1518)	1677..1683,1943..1950,587..593	3.38	1.6	-0.88	-0.01	1 Yes	
NM_016006	218739_at	ABHD5	Homo sapiens abhydrolase domain containing 5 (ABHD5), mRNA.	51099	TR(1..1536)CDS(111..1160)	1328..1334,246..252	3.12	1.8	-1.87	-0.56	1 Yes	
NM_001078650	218531_at	TMEM134	Homo sapiens transmembrane protein 134 (TMEM134), transcript variant 2, mRNA.	80194	TR(1..885)CDS(69..611)	791..798	3.06	2.29	-0.82	-0.54	1 Yes	
NM_004886	215148_s_at	APBA3	Homo sapiens amyloid beta (A4) precursor protein-binding, family A, member 3 (X11-like 2) (APBA3), mRNA.	9546	TR(1..2158)CDS(178..1905)	253..259,1930..1936	3.05	2.3	-0.5	-0.13	1 No	
XM_001128702	238567_at	SGPP2	PREDICTED: Homo sapiens sphingosine-1-phosphate phosphatase 2 (SGPP2), mRNA.	130367	TR(1..1190)CDS(1..1017)	928..935	3.05	1.81	-0.45	-0.28	1 No	
NM_022044	218681_s_at	SDF2L1	Homo sapiens stromal cell-derived factor 2-like 1 (SDF2L1), mRNA.	23753	TR(1..876)CDS(85..750)	476..482,842..848	3.04	2.06	-1.05	-0.37	1 Yes	
NM_002306	208949_s_at	LGALS3	Homo sapiens lectin, galactoside-binding, soluble, 3 (LGALS3), transcript variant 1, mRNA.	3958	TR(1..1080)CDS(152..904)	497..504	2.97	1.99	-0.42	-0.31	1 Yes	
XM_377476	1568987_at	MGC57346	PREDICTED: Homo sapiens hypothetical LOC401884 (MGC57346), mRNA.	401884	TR(1..1879)CDS(1..351)	450..457,371..377	3.04	1.49	-0.54	-0.2	1 No	
NM_014481	204408_at	APEX2	Homo sapiens APEX nuclease (apurinic/apyrimidinic endonuclease) 2 (APEX2), nuclear gene encoding mitochondrial protein, mRNA.	27301	TR(1..1991)CDS(77..1633)	1811..1818	2.93	2.2	-0.72	-0.47	1 Yes	
NM_197939	223592_s_at,223591_at	RNF135	Homo sapiens ring finger protein 135 (RNF135), transcript variant 2, mRNA.	84282	TR(1..2032)CDS(137..769)	1923..1929	2.79	2.05	-1.13	-0.5	1 Yes	
NM_001077426	219711_at	ZNF586	Homo sapiens zinc finger protein 586 (ZNF586), transcript variant 2, mRNA.	54807	TR(1..2088)CDS(187..819)	1422..1428	2.75	1.89	-0.63	-0.02	1 No	
NM_022821	57163_at,218028_at	ELOVL1	Homo sapiens elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like 1 (ELOVL1), mRNA.	64834	TR(1..1501)CDS(129..968)	1244..1250	2.74	1.9	-1.31	-0.78	1 Yes	
NM_021198	217844_at	CTDSP1	Homo sapiens CTD (carboxy-terminal domain, RNA polymerase II, polypeptide A) small phosphatase 1 (CTDSP1), transcript variant 1, mRNA.	58190	TR(1..2655)CDS(337..1122)	1518..1524,2211..2217,1793..1799,2577..2583,1329..1335	2.74	1.72	-1.9	-1.13	1 Yes	
NM_015690	231806_s_at	STK36	Homo sapiens serine/threonine kinase 36, fused homolog (Drosophila) (STK36), mRNA.	27148	TR(1..4887)CDS(220..4167)	4539..4545,4436..4442	2.65	1.61	-1.11	-0.35	1 Yes	
NM_015044	208915_s_at,208913_at,208914_at,210658_s_at	GGA2	Homo sapiens golgi associated, gamma adaptin ear containing, ARF binding protein 2 (GGA2), mRNA.	23062	TR(1..5203)CDS(77..1918)	319..325,2313..2319,2582..2589	2.64	1.64	-1.19	-0.46	1 Yes	
NM_005697	218143_s_at	SCAMP2	Homo sapiens secretory carrier membrane protein 2 (SCAMP2), mRNA.	10066	TR(1..1313)CDS(75..1064)	664..670,1129..1136	2.62	1.87	-1.27	-0.58	1 Yes	
NM_022337	219412_at	RAB38	Homo sapiens RAB38, member RAS oncogene family (RAB38), mRNA.	23682	TR(1..1412)CDS(48..683)	1296..1302	2.92	1.33	-1.82	-0.7	1 No	
NM_001079519	227029_at	C14orf24	Homo sapiens chromosome 14 open reading frame 24 (C14orf24), transcript variant 2, mRNA.	283635	TR(1..3276)CDS(460..1101)	3198..3205	2.6	1.38	-1.66	-0.83	1 No	
NM_022365	222621_at,218409_s_at,222620_s_at	DNAJC1	Homo sapiens DnaJ (Hsp40) homolog, subfamily C, member 1 (DNAJC1), mRNA.	64215	TR(1..2213)CDS(288..1952)	2052..2059,981..987	2.58	1.88	-1.15	-0.64	1 Yes	
NM_001081003	224387_at	COMMD5	Homo sapiens COMM domain containing 5 (COMMD5), transcript variant 2, mRNA.	28991	TR(1..1422)CDS(234..908)	1305..1311	2.56	1.64	-0.16	0.04	1 No	
NM_006663	218849_s_at	PPP1R13L	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 13 like (PPP1R13L), mRNA.	10848	TR(1..3117)CDS(67..2553)	3024..3030,2742..2748	2.86	1.29	-0.69	-0.26	1 Yes	
NM_130468	226314_at	CHST14	Homo sapiens carbohydrate (N-acetyl)galactosamine 4-O) sulfotransferase 14 (CHST14), mRNA.	113189	TR(1..2156)CDS(201..1331)	1803..1809	2.5	1.54	-0.75	-0.32	1 No	
NM_004937	36566_at,204925_at	CTNS	Homo sapiens cystinosis, nephropathic (CTNS), transcript variant 2, mRNA.	1497	TR(1..4431)CDS(594..1697)	2514..2521	2.48	1.73	-0.77	-0.48	1 Yes	
NM_024310	219566_at	PLEKHF1	Homo sapiens pleckstrin homology domain containing, family F (with FYVE domain) member 1 (PLEKHF1), mRNA.	79156	TR(1..1774)CDS(103..942)	1088..1094	2.48	1.44	0.09	-0.04	1 Yes	
NM_018090	220731_s_at	NECAP2	Homo sapiens NECAP endocytosis associated 2 (NECAP2), mRNA.	55707	TR(1..2037)CDS(25..816)	1743..1750	2.48	1.3	-1.54	-0.84	1 No	
NM_007162	221866_at,50221_at	TFEB	Homo sapiens transcription factor EB (TFEB), mRNA.	7942	TR(1..2446)CDS(303..1733)	2327..2333	2.44	1.65	-0.34	-0.17	1 Yes	
NM_015917	217751_at	GSTK1	Homo sapiens glutathione S-transferase kappa 1 (GSTK1), mRNA.	373156	TR(1..1017)CDS(44..724)	938..944	2.43	1.55	-0.68	-0.22	1 Yes	
NM_006016	208405_s_at,208654_s_at	CD164	Homo sapiens CD164 molecule, sialomucin (CD164), mRNA.	8763	TR(1..3026)CDS(82..675)	2327..2334,1326..1333	2.55	1.25	-2.28	-1.06	1 No	
NM_001010924	212771_at	C10orf38	Homo sapiens chromosome 10 open reading frame 38 (C10orf38), mRNA.	221061	TR(1..3964)CDS(8..2680)	3422..3428	2.38	1.69	-0.77	-0.36	1 Yes	
NM_006858	203679_at	TMED1	Homo sapiens transmembrane emp24 protein transport domain containing 1 (TMED1), mRNA.	11018	TR(1..1374)CDS(117..800)	1088..1095	2.36	1.5	-1.03	-0.53	1 No	
NM_001845	211980_at,211981_at	COL4A1	Homo sapiens collagen, type IV, alpha 1 (COL4A1), mRNA.	1282	TR(1..6549)CDS(123..5132)	5457..5463,5250..5257	2.32	1.44	-1.45	-0.68	1 No	
NM_004781	201337_s_at,201336_at,211749_s_at	VAMP3	Homo sapiens vesicle-associated membrane protein 3 (cellulobrevin) (VAMP3), mRNA.	9341	TR(1..2225)CDS(116..418)	558..564,855..862,620..626	2.31	1.34	-2.07	-0.94	1 Yes	
NM_032890	228184_at	DISP1	Homo sapiens dispatched homolog 1 (Drosophila) (DISP1), mRNA.	84976	TR(1..4743)CDS(148..4722)	3543..3549,755..761,3602..3608,2325..2331	2.29	1.7	-0.78	-0.05	1 No	
NM_005702	212087_s_at	ERAL1	Homo sapiens Era G-protein-like 1 (E. coli) (ERAL1), mRNA.	26284	TR(1..1856)CDS(11..1324)	1515..1521,1100..1107	2.26	1.43	-0.01	-0.1	1 Yes	
NM_004390	202295_s_at	CTSH	Homo sapiens cathepsin H (CTSH), transcript variant 1, mRNA.	1512	TR(1..1494)CDS(98..1105)	1318..1325	2.19	1.59	-0.76	-0.13	1 Yes	
NM_001012334	209035_at	MDK	Homo sapiens midkine (neurite growth-promoting factor 2) (MDK), transcript variant 1, mRNA.	4192	TR(1..997)CDS(223..654)	744..750	2.24	1.18	-0.24	-0.05	1 Yes	
NM_030935	208104_s_at	TSC22D4	Homo sapiens TSC22 domain family, member 4 (TSC22D4), mRNA.	81628	TR(1..2399)CDS(756..1943)	1963..1969	2.17	1.62	-0.11	-0.01	1 Yes	
NM_001084	202185_at	PLOD3	Homo sapiens procollagen-lysine, 2-oxoglutarate 5-dioxygenase 3 (PLOD3), mRNA.	8985	TR(1..2995)CDS(457..2673)	2708..2714	2.15	1.41	-1.25	-0.73	1 Yes	
NM_001085354	219038_at	MORC4	Homo sapiens MORC family CW-type zinc finger 4 (MORC4), transcript variant 2, mRNA.	79710	TR(1..3777)CDS(244..2946)	3140..3146,2434..2441	2.47	1.12	-0.94	-0.12	1 No	

NM_006117	218025_s_at	PECI	Homo sapiens peroxisomal D3,D2-enoyl-CoA isomerase (PECI), transcript variant 1, mRNA.	10455 TR(1..1410)CDS(118..1212)	1299..1305	2.26	1.13	-1.04	-0.41	1 No
NM_017814	43977_at	TMEM161A	Homo sapiens transmembrane protein 161A (TMEM161A), mRNA.	54929 TR(1..1795)CDS(31..1470)	746..752	2.08	1.2	-0.24	-0.14	1 No
NM_005009	212739_s_at	NME4	Homo sapiens non-metastatic cells 4, protein expressed in (NME4), mRNA.	4833 TR(1..1059)CDS(32..595)	755..761	2.06	1.15	-1.12	-0.48	1 Yes
NM_014319	218604_at	LEMD3	Homo sapiens LEM domain containing 3 (LEMD3), mRNA.	23592 TR(1..4785)CDS(7..2742)	4228..4235	2.03	1.35	-0.47	-0.4	1 No
NM_018845	219125_s_at	RAG1AP1	Homo sapiens recombination activating gene 1 activating protein 1 (RAG1AP1), mRNA.	55974 TR(1..1315)CDS(60..725)	1243..1249,977..983,1266..1272	2.11	1.1	-0.33	-0.15	1 Yes
NM_015158	213005_s_at	ANKRD15	Homo sapiens ankyrin repeat domain 15 (ANKRD15), transcript variant 1, mRNA.	23189 TR(1..5083)CDS(136..4194)	5014..5021	2.02	1.4	-1.23	-0.51	1 Yes
NM_001078175	201802_at,201801_s_at	SLC29A1	Homo sapiens solute carrier family 29 (nucleoside transporters), member 1 (SLC29A1), nuclear gene encoding mitochondrial protein, transcript variant 2, mRNA.	2030 TR(1..2253)CDS(242..1612)	2179..2185,852..858	2.02	1.19	-0.11	0.08	1 No
NM_006410	207180_s_at,210253_at,209448_at	HTATIP2	Homo sapiens HIV-1 Tat interactive protein 2, 30kDa (HTATIP2), transcript variant 2, mRNA.	10553 TR(1..1404)CDS(67..795)	1273..1279	2.01	1.35	-0.52	-0.26	1 Yes
NM_139029	204306_s_at	CD151	Homo sapiens CD151 molecule (Raph blood group) (CD151), transcript variant 4, mRNA.	977 TR(1..1570)CDS(141..902)	1042..1048	2.05	1.11	-0.66	-0.25	1 Yes
NM_012434	223441_at,221041_s_at	SLC17A5	Homo sapiens solute carrier family 17 (anion/sugar transporter), member 5 (SLC17A5), mRNA.	26503 TR(1..3292)CDS(88..1575)	2176..2182,1636..1642,1085..1091	1.99	1.27	-1.35	-0.89	1 Yes
NM_012110	219492_at	CHIC2	Homo sapiens cysteine-rich hydrophobic domain 2 (CHIC2), mRNA.	26511 TR(1..1151)CDS(322..819)	868..874	1.99	1.23	-0.68	-0.25	1 Yes
NM_014587	226913_s_at	SOX8	Homo sapiens SRY (sex determining region Y)-box 8 (SOX8), mRNA.	30812 TR(1..3069)CDS(116..1456)	2758..2764	1.99	1.16	-0.08	-0.53	1 No
NM_178507	225510_at	OAF	Homo sapiens OAF homolog (Drosophila) (OAF), mRNA.	220323 TR(1..1957)CDS(242..1063)	1673..1679,1273..1279	1.98	1.21	-0.99	-0.46	1 Yes
NM_148178	225403_at	C9orf23	Homo sapiens chromosome 9 open reading frame 23 (C9orf23), transcript variant 1, mRNA.	138716 TR(1..896)CDS(78..569)	619..625	1.96	1.24	-0.17	-0.19	1 Yes
NM_138401	227864_s_at	FAM125A	Homo sapiens family with sequence similarity 125, member A (FAM125A), mRNA.	93343 TR(1..1216)CDS(34..855)	1106..1112,195..202	1.95	1.19	-0.16	-0.18	1 No
NM_001035	214044_at,207557_s_at	RYR2	Homo sapiens ryanodine receptor 2 (cardiac) (RYR2), mRNA.	6262 TR(1..16365)CDS(121..15024)	15486..15493,3518..3524	1.95	1.29	-0.42	-0.18	1 No
NM_138410	226017_at	CMTM7	Homo sapiens CKLF-like MARVEL transmembrane domain containing 7 (CMTM7), transcript variant 1, mRNA.	112616 TR(1..1369)CDS(237..764)	1141..1148	2.04	1.06	-0.13	0	1 No
NM_004862	200704_at,200706_s_at	LITAF	Homo sapiens lipopolysaccharide-induced TNF factor (LITAF), mRNA.	9516 TR(1..2368)CDS(234..719)	1209..1215,1107..1113	2.09	1.05	-1.46	-0.66	1 Yes
NM_006478	208258_s_at,209729_at	GAS2L1	Homo sapiens growth arrest-specific 2 like 1 (GAS2L1), transcript variant 1, mRNA.	10634 TR(1..2557)CDS(195..2240)	2446..2452	1.93	1.2	-0.11	-0.15	1 Yes
NM_002668	201136_at	PLP2	Homo sapiens proteolipid protein 2 (colonic epithelium-enriched) (PLP2), mRNA.	5355 TR(1..945)CDS(76..534)	654..660	1.92	1.26	-0.95	-0.41	1 Yes
NM_014415	204847_at	ZBTB11	Homo sapiens zinc finger and BTB domain containing 11 (ZBTB11), mRNA.	27107 TR(1..5049)CDS(159..3320)	2979..2985,4669..4675	2.54	1.03	-1.1	-0.32	1 No
NM_018214	218816_at	LRRC1	Homo sapiens leucine rich repeat containing 1 (LRRC1), mRNA.	55227 TR(1..3180)CDS(278..1852)	2003..2010,2446..2453	2.44	1.03	-2.05	-0.63	1 Yes
NM_006714	213624_at	SMPDL3A	Homo sapiens sphingomyelin phosphodiesterase, acid-like 3A (SMPDL3A), mRNA.	10924 TR(1..1759)CDS(69..1430)	1569..1575	2.54	1.03	-1.28	-0.19	1 No
NM_006764	209100_at	IFRD2	Homo sapiens interferon-related developmental regulator 2 (IFRD2), mRNA.	7866 TR(1..2045)CDS(215..1543)	1777..1783	1.91	1.1	-0.96	-0.5	1 Yes
NM_022075	222212_s_at	LASS2	Homo sapiens LAG1 homolog, ceramide synthase 2 (LASS2), transcript variant 2, mRNA.	29956 TR(1..2465)CDS(348..1490)	1870..1877	1.88	1.25	-1.08	-0.46	1 No
NM_024092	201361_at	TMEM109	Homo sapiens transmembrane protein 109 (TMEM109), mRNA.	79073 TR(1..2097)CDS(73..804)	1431..1438	1.87	1.41	-0.94	-0.68	1 No
NM_006014	219061_s_at	LAGE3	Homo sapiens L antigen family, member 3 (LAGE3), mRNA.	8270 TR(1..1816)CDS(343..774)	541..547	1.87	1.14	0.19	0.04	1 No
NM_174934	236359_at	SCN4B	Homo sapiens sodium channel, voltage-gated, type IV, beta (SCN4B), mRNA.	6330 TR(1..4485)CDS(150..836)	2073..2080	1.86	1.13	-0.82	-0.72	1 No
NM_023943	223524_s_at	TMEM108	Homo sapiens transmembrane protein 108 (TMEM108), mRNA.	66000 TR(1..2162)CDS(193..1920)	1135..1141	2.55	1.01	-0.36	-0.05	1 No
NM_006631	214686_at	ZNF266	Homo sapiens zinc finger protein 266 (ZNF266), mRNA.	10781 TR(1..3618)CDS(1272..2921)	1545..1551	2.46	1	-0.75	0.02	1 No
NM_000021	203460_s_at,207782_s_at	PSEN1	Homo sapiens presenilin 1 (Alzheimer disease 3) (PSEN1), mRNA.	5663 TR(1..2763)CDS(249..1652)	1502..1509	2.08	0.99	-0.52	-0.11	1 No
NM_024792	218898_at	FAM57A	Homo sapiens family with sequence similarity 57, member A (FAM57A), mRNA.	79850 TR(1..2096)CDS(42..815)	2035..2041	1.81	1.21	-1.25	-0.6	1 Yes
NM_005192	209714_s_at,1555758_a_at	CDKN3	Homo sapiens cyclin-dependent kinase inhibitor 3 (CDK2-associated dual specificity phosphatase) (CDKN3), mRNA.	1033 TR(1..836)CDS(62..700)	446..452	2.58	0.99	-0.39	0.03	1 No
NM_004568	211474_s_at	SERPINB6	Homo sapiens serpin peptidase inhibitor, clade B (ovalbumin), member 6 (SERPINB6), mRNA.	5269 TR(1..1665)CDS(377..1507)	409..415	2.43	0.99	-1.08	-0.34	1 Yes
NM_024033	220949_s_at	C7orf49	Homo sapiens chromosome 7 open reading frame 49 (C7orf49), mRNA.	78996 TR(1..1498)CDS(267..656)	772..778	1.9	1.01	-0.22	-0.1	1 No
NM_001154	200782_at	ANXA5	Homo sapiens annexin A5 (ANXA5), mRNA.	308 TR(1..1630)CDS(193..1155)	1253..1259	1.8	1.29	-0.47	-0.22	1 No
NM_199344	214838_at	SFT2D2	Homo sapiens SFT2 domain containing 2 (SFT2D2), mRNA.	375035 TR(1..1155)CDS(73..555)	651..657	2.13	0.98	-1.02	-0.29	1 No
NM_004447	202609_at	EPS8	Homo sapiens epidermal growth factor receptor pathway substrate 8 (EPS8), mRNA.	2059 TR(1..3912)CDS(262..2730)	3826..3832	1.81	1.04	-0.92	-0.22	1 No
NM_017514	203623_at,1553139_s_at	PLXNA3	Homo sapiens plexin A3 (PLXNA3), mRNA.	55558 TR(1..6762)CDS(173..5788)	243..249,6598..6605,5954..5960	1.79	1.05	-0.06	-0.01	1 No
NM_021158	1555788_a_at,218145_at	TRIB3	Homo sapiens tribbles homolog 3 (Drosophila) (TRIB3), mRNA.	57761 TR(1..2469)CDS(507..1583)	1677..1683	1.76	1.34	-0.4	-0.39	1 Yes
NM_152792	235514_at	SASP	Homo sapiens skin aspartic protease (SASP), mRNA.	151516 TR(1..2168)CDS(574..1605)	214..2121	1.74	1.26	-1.07	-0.23	1 No
NM_006902	205991_s_at,226695_at	PRRX1	Homo sapiens paired related homeobox 1 (PRRX1), transcript variant pmx-1a, mRNA.	5396 TR(1..4071)CDS(48..701)	1289..1296,1770..1777	1.74	1.28	-0.95	-0.97	1 No
NM_001033859	200710_at	ACADVL	Homo sapiens acyl-Coenzyme A dehydrogenase, very long chain (ACADVL), nuclear gene encoding mitochondrial protein, transcript variant 2, mRNA.	37 TR(1..2254)CDS(152..2053)	2084..2090	1.79	1	-0.51	-0.14	1 Yes
XM_001131046	209191_at	TUBB6	PREDICTED: Homo sapiens tubulin, beta 6 (TUBB6), mRNA.	84617 TR(1..1698)CDS(1..1212)	1580..1586	1.83	0.98	-0.73	-0.27	1 Yes
NM_020159	223197_s_at	SMARCAD1	Homo sapiens SWI/SNF-related, matrix-associated actin-dependent regulator of chromatin, subfamily a, containing DEAD/H box 1 (SMARCAD1), mRNA.	56916 TR(1..4934)CDS(78..3158)	3549..3556	2.55	0.95	-0.73	-0.31	1 No
NM_003870	200791_s_at,210840_s_at,213446_s_at	IQGAP1	Homo sapiens IQ motif containing GTPase activating protein 1 (IQGAP1), mRNA.	8826 TR(1..7219)CDS(102..5075)	1970..1976,5950..5957,6247..6253	1.71	1.14	-1.18	-0.51	1 No
NM_022060	218581_at	ABHD4	Homo sapiens abhydrolase domain containing 4 (ABHD4), mRNA.	63874 TR(1..2543)CDS(70..1098)	2003..2009	1.72	1.04	-0.11	-0.29	1 Yes
NM_004364	204039_at	CEBPA	Homo sapiens CCAAT/enhancer binding protein (C/EBP), alpha (CEBPA), mRNA.	1050 TR(1..2385)CDS(1..1077)	2059..2065,1418..1424,1361..1367	1.73	1.02	-0.78	-0.6	1 Yes
NM_001654	201895_at	ARAF	Homo sapiens v-raf murine sarcoma 3611 viral oncogene homolog (ARAF), mRNA.	369 TR(1..2442)CDS(134..1954)	1946..1953,1653..1659,476..482,2280..2286	1.7	1.07	-0.99	-0.47	1 Yes
NM_005327	201036_s_at,201035_s_at,211569_s_at	HADH	Homo sapiens hydroxyacyl-Coenzyme A dehydrogenase (HADH), nuclear gene encoding mitochondrial protein, mRNA.	3033 TR(1..1929)CDS(150..1094)	1345..1351	1.86	0.95	-0.85	-0.4	1 No
NM_003098	203516_at	SNTA1	Homo sapiens syntrophin, alpha 1 (dystrophin-associated protein A1, 59kDa, acidic component) (SNTA1), mRNA.	6640 TR(1..2345)CDS(273..1790)	2024..2030	1.69	1.63	-0.1	-0.43	1 No
NM_139012	202530_at	MAPK14	Homo sapiens mitogen-activated protein kinase 14 (MAPK14), transcript variant 2, mRNA.	1432 TR(1..3757)CDS(363..1445)	2049..2056	2.08	0.93	-1.13	-0.55	1 No
NM_003945	200096_s_at	ATP6V0E1	Homo sapiens ATPase, H+ transporting, lysosomal 9kDa, V0 subunit e1 (ATP6V0E1), mRNA.	8992 TR(1..894)CDS(102..347)	496..502,468..474,171..177	1.66	1.05	-1.37	-0.68	1 No

NM_138355	228730_s_at	SCRN2	Homo sapiens secernin 2 (SCRN2), mRNA.	90507 TR(1..1488)CDS(40..1317)	704..710,1444..1450	1.66	1.02	-0.02	0.04	1 No
NM_182760	226850_at	SUMF1	Homo sapiens sulfatase modifying factor 1 (SUMF1), mRNA.	285362 TR(1..2148)CDS(26..1150)	1894..1900	2.27	0.92	-1.9	-0.78	1 Yes
NM_018290	223738_s_at,225366_at,225367_at	PGM2	Homo sapiens phosphoglucomutase 2 (PGM2), mRNA.	55276 TR(1..3192)CDS(14..1852)	2144..2151,1174..1180	2.08	0.92	-1.42	-0.5	1 No
NM_020376	39854_r_at	PNPLA2	Homo sapiens patatin-like phospholipase domain containing 2 (PNPLA2), mRNA.	57104 TR(1..2113)CDS(203..1717)	1906..1913	2.19	0.9	-0.12	-0.01	1 No
NM_017556	1554795_a_at	FBLIM1	Homo sapiens filamin binding LIM protein 1 (FBLIM1), transcript variant 1, mRNA.	54751 TR(1..3363)CDS(300..1421)	1800..1806,847..853,2103..2109	2.03	0.9	-0.14	-0.01	1 Yes
NM_024779	218942_at	PIP4K2C	Homo sapiens phosphatidylinositol-5-phosphate 4-kinase, type II, gamma (PIP4K2C), mRNA.	79837 TR(1..2937)CDS(17..1282)	1558..1564	1.59	1.11	-0.65	-0.51	1 No
NM_181726	227337_at	ANKRD37	Homo sapiens ankyrin repeat domain 37 (ANKRD37), mRNA.	353322 TR(1..934)CDS(239..715)	224..230	1.58	1.2	0.05	-0.13	1 No
NM_012249	212122_at	RHOQ	Homo sapiens ras homolog gene family, member Q (RHOQ), mRNA.	23433 TR(1..4542)CDS(320..937)	2322..2328,3352..3359,3521..3527	2.23	0.89	-1.76	-0.67	1 No
NM_031307	221277_s_at	PUS3	Homo sapiens pseudouridylyl synthase 3 (PUS3), mRNA.	83480 TR(1..1851)CDS(88..1533)	1792..1798,1286..1292,452..458	2.02	0.88	-0.81	-0.17	1 No
NM_002196	206502_s_at	INSM1	Homo sapiens insulinoma-associated 1 (INSM1), mRNA.	3642 TR(1..2838)CDS(148..1680)	1710..1716	1.55	1.05	-0.25	-0.09	1 No
NM_152707	214140_at,209910_at	SLC25A16	Homo sapiens solute carrier family 25 (mitochondrial carrier; Graves disease autoantigen), member 16 (SLC25A16), nuclear gene encoding mitochondrial protein, mRNA.	8034 TR(1..2560)CDS(453..1451)	1289..1296	1.54	1.06	-0.56	-0.27	1 No
NM_004926	211965_at	ZFP36L1	Homo sapiens zinc finger protein 36, C3H type-like 1 (ZFP36L1), mRNA.	677 TR(1..3022)CDS(131..1147)	1516..1522	1.54	1.35	-0.03	-0.7	1 Yes
NM_032256	223594_at	TMEM117	Homo sapiens transmembrane protein 117 (TMEM117), mRNA.	84216 TR(1..2706)CDS(67..1611)	1917..1923	1.63	0.92	-1.18	-0.29	1 No
NM_198080	225782_at,1554126_at,1554127_s_at,238583_at	MSRB3	Homo sapiens methionine sulfoxide reductase B3 (MSRB3), transcript variant 1, mRNA.	253827 TR(1..4242)CDS(62..640)	4089..4095,267..273	1.75	0.88	-0.18	-0.06	1 Yes
NM_006320	201701_s_at	PGRMC2	Homo sapiens progesterone receptor membrane component 2 (PGRMC2), mRNA.	10424 TR(1..1873)CDS(6..677)	1394..1401	1.61	0.91	-0.82	-0.43	1 Yes
NM_014445	200969_at,200970_s_at	SERP1	Homo sapiens stress-associated endoplasmic reticulum protein 1 (SERP1), mRNA.	27230 TR(1..3181)CDS(507..707)	2352..2359,895..901	2.18	0.86	-1.65	-0.56	1 No
NM_018226	218301_at	RNPEPL1	Homo sapiens arginyl aminopeptidase (aminopeptidase B)-like 1 (RNPEPL1), mRNA.	57140 TR(1..2948)CDS(594..2078)	2436..2443,2122..2128	1.8	0.87	-0.24	-0.38	1 Yes
NM_030805	221274_s_at	LMAN2L	Homo sapiens lectin, mannose-binding 2-like (LMAN2L), mRNA.	81562 TR(1..2412)CDS(25..1071)	2242..2248,1301..1307,311..317,1575..1581	1.75	0.87	-0.45	-0.15	1 No
NM_005827	202433_at	SLC35B1	Homo sapiens solute carrier family 35, member B1 (SLC35B1), mRNA.	10237 TR(1..1186)CDS(88..1056)	240..246	1.82	0.86	-0.03	-0.04	1 No
NM_014305	208249_s_at	TGDS	Homo sapiens TDP-glucose 4,6-dehydratase (TGDS), mRNA.	23483 TR(1..1952)CDS(122..1174)	450..456,1073..1079,1621..1627	2.44	0.85	-1.04	-0.16	1 No
NM_001098832	225319_s_at	FAM104A	Homo sapiens family with sequence similarity 104, member A (FAM104A), transcript variant 1, mRNA.	84923 TR(1..2905)CDS(89..712)	1950..1957	1.62	0.88	-0.57	-0.1	1 Yes
NM_031279	221008_s_at	AGXT2L1	Homo sapiens alanine-glyoxylate aminotransferase 2-like 1 (AGXT2L1), mRNA.	64850 TR(1..2107)CDS(156..1655)	2009..2016,1987..1993	1.46	1.01	-0.97	-0.9	1 No
NM_000075	202246_s_at	CDK4	Homo sapiens cyclin-dependent kinase 4 (CDK4), mRNA.	1019 TR(1..1474)CDS(228..1139)	1270..1276	1.46	0.96	-0.49	-0.23	1 No
NM_020195	213398_s_at	C14orf124	Homo sapiens chromosome 14 open reading frame 124 (C14orf124), mRNA.	56948 TR(1..1247)CDS(33..914)	890..896	1.68	0.85	0.09	0.09	1 Yes
NM_015666	226606_s_at,226598_s_at	GTPBP5	Homo sapiens GTP binding protein 5 (putative) (GTPBP5), mRNA.	26164 TR(1..2949)CDS(39..1259)	1338..1344	1.59	0.87	-0.1	-0.11	1 No
NM_004789	211219_s_at,206140_at	LHX2	Homo sapiens LIM homeobox 2 (LHX2), mRNA.	9355 TR(1..2416)CDS(740..1960)	2265..2272	1.43	1	-0.22	-0.38	1 No
NM_019895	1554149_at,208925_at	CLDND1	Homo sapiens claudin domain containing 1 (CLDND1), transcript variant 2, mRNA.	56650 TR(1..2244)CDS(288..1049)	2001..2007	1.73	0.84	-1.49	-0.69	1 Yes
NM_001008393	238015_at	LOC201725	Homo sapiens hypothetical protein LOC201725 (LOC201725), mRNA.	201725 TR(1..3503)CDS(226..567)	1429..1436	1.64	0.85	-0.9	-0.64	1 No
NM_017785	221685_s_at	CCDC99	Homo sapiens coiled-coil domain containing 99 (CCDC99), mRNA.	54908 TR(1..2683)CDS(280..2097)	2227..2233	2.25	0.82	-1.49	-0.36	1 No
NM_145810	224428_s_at	CDCA7	Homo sapiens cell division cycle associated 7 (CDCA7), transcript variant 2, mRNA.	83879 TR(1..2602)CDS(132..1247)	2192..2198	1.54	0.88	-0.94	-0.36	1 Yes
NM_012280	213937_s_at,205324_s_at	FTSJ1	Homo sapiens FtsJ homolog 1 (E. coli) (FTSJ1), transcript variant 1, mRNA.	24140 TR(1..1951)CDS(324..1313)	1020..1026	1.76	0.83	-0.19	-0.01	1 Yes
NM_020360	56197_at,218828_at	PLSCR3	Homo sapiens phospholipid scramblase 3 (PLSCR3), mRNA.	57048 TR(1..1746)CDS(210..1097)	1656..1663,423..429	2.15	0.82	-0.5	-0.01	1 Yes
NM_152695	228968_at,241066_at	ZNF449	Homo sapiens zinc finger protein 449 (ZNF449), mRNA.	203523 TR(1..4034)CDS(141..1697)	2694..2700,3694..3701	1.52	0.88	-0.76	-0.25	1 No
NM_006289	203254_s_at	TLN1	Homo sapiens talin 1 (TLN1), mRNA.	7094 TR(1..8187)CDS(91..7716)	7897..7903,4808..4814	2.16	0.81	-0.98	-0.45	1 Yes
NM_173560	1552673_at	RFXDC1	Homo sapiens regulatory factor X domain containing 1 (RFXDC1), mRNA.	222546 TR(1..3538)CDS(64..2850)	3172..3178	1.38	1.04	-1.1	-0.65	1 No
NM_015221	212838_at	DNMBP	Homo sapiens dynamin binding protein (DNMBP), mRNA.	23268 TR(1..6419)CDS(93..4826)	2934..2940	2.02	0.8	-0.62	-0.13	1 Yes
NM_003029	201469_s_at	SHC1	Homo sapiens SHC (Src homology 2 domain containing) transforming protein 1 (SHC1), transcript variant 2, mRNA.	6464 TR(1..3076)CDS(121..1545)	1975..1981	1.85	0.8	-0.21	-0.16	1 No
NM_001080522	234936_s_at,1559409_a_at	KIAA1345	Homo sapiens KIAA1345 protein (KIAA1345), mRNA.	57545 TR(1..4890)CDS(73..4758)	4592..4598,3725..3732,3750..3757	1.66	0.8	-0.03	-0.01	1 Yes
NM_003627	204394_at	SLC43A1	Homo sapiens solute carrier family 43, member 1 (SLC43A1), mRNA.	8501 TR(1..2525)CDS(257..1936)	1518..1524,1351..1357,875..881	1.36	0.92	0.28	-0.07	1 Yes
NM_138485	223486_at,221046_s_at	GTPBP8	Homo sapiens GTP-binding protein 8 (putative) (GTPBP8), transcript variant 2, mRNA.	29083 TR(1..1265)CDS(48..803)	945..951	2.2	0.78	-1.16	-0.36	1 No
NM_022152	217730_at	TMBIM1	Homo sapiens transmembrane BAX inhibitor motif containing 1 (TMBIM1), mRNA.	64114 TR(1..2375)CDS(133..1068)	2173..2179	1.81	0.78	-0.78	-0.45	1 No
NM_032166	1552937_s_at	ATRIP	Homo sapiens ATR interacting protein (ATRIP), transcript variant 2, mRNA.	84126 TR(1..2428)CDS(33..2327)	1721..1728	1.85	0.78	-0.3	0.15	1 No
NM_194281	229582_at	C18orf37	Homo sapiens chromosome 18 open reading frame 37 (C18orf37), transcript variant 2, mRNA.	125476 TR(1..987)CDS(118..696)	809..815	1.56	0.81	-0.75	-0.38	1 No
NM_014168	221570_s_at	METTL5	Homo sapiens methyltransferase like 5 (METTL5), mRNA.	29081 TR(1..956)CDS(247..876)	209..215	1.96	0.74	-0.48	-0.06	1 Yes
NM_003331	205546_s_at	TYK2	Homo sapiens tyrosine kinase 2 (TYK2), mRNA.	7297 TR(1..4224)CDS(342..3905)	4022..4029	1.41	0.83	-0.1	-0.06	1 Yes
NM_001013649	238768_at	LOC388969	Homo sapiens hypothetical LOC388969 (LOC388969), mRNA.	388969 TR(1..2849)CDS(46..546)	2123..2129,2039..2046,1730..1736	1.85	0.73	-1.06	-0.53	1 No
NM_012401	208890_s_at	PLXNB2	Homo sapiens plexin B2 (PLXNB2), mRNA.	23654 TR(1..6369)CDS(107..5623)	5905..5912,1768..1774	1.53	0.78	-0.21	-0.13	1 No
NM_023927	218706_s_at	GRAMD3	Homo sapiens GRAM domain containing 3 (GRAMD3), mRNA.	65983 TR(1..2912)CDS(199..1497)	1826..1832	1.63	0.75	-1.19	0.04	1 Yes
NM_013375	218405_at	ABT1	Homo sapiens activator of basal transcription 1 (ABT1), mRNA.	29777 TR(1..2255)CDS(32..850)	1344..1350	1.21	0.91	-0.05	-0.14	1 No
NM_003645	205768_s_at,205769_at	SLC27A2	Homo sapiens solute carrier family 27 (fatty acid transporter), member 2 (SLC27A2), mRNA.	11001 TR(1..2343)CDS(204..2066)	860..866	2.01	0.72	-0.41	-0.03	1 No
NM_013232	203415_at	PDCC6	Homo sapiens programmed cell death 6 (PDCC6), mRNA.	10016 TR(1..1151)CDS(101..676)	1056..1063,89..95	1.7	0.73	-0.65	-0.16	1 Yes
NM_030818	208094_s_at	CCDC130	Homo sapiens coiled-coil domain containing 130 (CCDC130), mRNA.	81576 TR(1..1679)CDS(218..1408)	1590..1596	1.13	0.97	0.1	-0.12	1 No
NM_018706	209916_at,227094_at	DHTKD1	Homo sapiens dehydrogenase E1 and transketolase domain containing 1 (DHTKD1), mRNA.	55526 TR(1..5202)CDS(100..2859)	4974..4981	1.46	0.78	-0.62	-0.33	1 No

NM_031934	1555630_a_at,224710_at	RAB34	Homo sapiens RAB34, member RAS oncogene family (RAB34), mRNA.	83871 TR(1..1740)CDS(603..1382)	1707..1714	1.22	0.89	-0.72	-0.48	1 Yes
NM_015345	212793_at	DAAM2	Homo sapiens dishevelled associated activator of morphogenesis 2 (DAAM2), mRNA.	23500 TR(1..6215)CDS(184..3387)	5698..5704	1.17	0.91	-0.61	-0.36	1 Yes
NM_016946	221664_s_at,223000_s_at,224097_s_at	F11R	Homo sapiens F11 receptor (F11R), transcript variant 1, mRNA.	50848 TR(1..3660)CDS(71..970)	612..618,1058..1064,1003..1009	1.86	0.71	-0.95	-0.63	1 No
NM_138821	202336_s_at	PAM	Homo sapiens peptidylglycine alpha-amidating monooxygenase (PAM), transcript variant 3, mRNA.	5066 TR(1..3636)CDS(374..2974)	3356..3362	1.69	0.72	-1.57	-0.37	1 No
XM_001128922	229238_at	LOC400566	PREDICTED: Homo sapiens hypothetical gene supported by AK128660 (LOC400566), mRNA.	400566 TR(1..1814)CDS(1..1263)	1440..1446	1.24	0.86	-0.15	0.34	1 No
NM_144635	221904_at,232806_s_at	FAM131A	Homo sapiens family with sequence similarity 131, member A (FAM131A), mRNA.	131408 TR(1..3680)CDS(1365..2372)	2143..2149	1.69	0.71	0.19	-0.02	1 No
NM_198504	1558322_a_at	PAQR9	Homo sapiens progesterin and adipoQ receptor family member IX (PAQR9), mRNA.	344838 TR(1..2119)CDS(1..1134)	2083..2089	1.05	1.59	-0.61	-0.87	1 No
XM_001127234	214440_at	NAT1	PREDICTED: Homo sapiens N-acetyltransferase 1 (arylamine N-acetyltransferase) (NAT1), mRNA.	9 TR(1..1835)CDS(381..1253)	816..822	1.75	0.7	-0.52	-0.19	1 No
NM_014447	214483_s_at,218230_at	ARFIP1	Homo sapiens ADP-ribosylation factor interacting protein 1 (arfaptin 1) (ARFIP1), transcript variant 3, mRNA.	27236 TR(1..2860)CDS(142..1167)	2076..2082	2.22	0.68	-1.35	-0.48	1 No
NM_005149	206838_at	TBX19	Homo sapiens T-box 19 (TBX19), mRNA.	9095 TR(1..2882)CDS(52..1398)	2494..2501,2777..2783	1.11	0.9	-0.06	-0.01	1 No
NM_017953	218932_at	C1orf181	Homo sapiens chromosome 1 open reading frame 181 (C1orf181), mRNA.	54680 TR(1..2821)CDS(135..1547)	2489..2495	1.65	0.7	-1.12	-0.36	1 No
NM_006809	201870_at	TOMM34	Homo sapiens translocase of outer mitochondrial membrane 34 (TOMM34), nuclear gene encoding mitochondrial protein, mRNA.	10953 TR(1..2066)CDS(141..1070)	690..696,1629..1635	1.48	0.74	-0.95	-0.48	1 No
NM_006633	203474_at	IQGAP2	Homo sapiens IQ motif containing GTPase activating protein 2 (IQGAP2), mRNA.	10788 TR(1..5769)CDS(223..4950)	3296..3302,5101..5107,3428..3434	2.42	0.67	-1.43	-0.22	1 No
NM_080430	226051_at	SELM	Homo sapiens selenoprotein M (SELM), mRNA.	140606 TR(1..739)CDS(61..498)	684..690	1.89	0.67	0.39	0.14	1 No
NM_152285	226405_s_at	ARRDC1	Homo sapiens arrestin domain containing 1 (ARRDC1), mRNA.	92714 TR(1..1621)CDS(75..1376)	996..1002,1440..1446	1.39	0.77	-0.04	-0.14	1 Yes
NM_002632	209652_s_at	PGF	Homo sapiens placental growth factor, vascular endothelial growth factor-related protein (PGF), mRNA.	5228 TR(1..1758)CDS(347..859)	1034..1040	1.64	0.7	0.25	0	1 Yes
NM_003387	202664_at,202665_s_at,202663_at	WIPF1	Homo sapiens WAS/WASL interacting protein family, member 1 (WIPF1), transcript variant 1, mRNA.	7456 TR(1..4600)CDS(101..1612)	980..986,4224..4231,554..560	1.46	0.74	-0.75	-0.5	1 No
NM_024607	1552670_a_at,222662_at	PPP1R3B	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 3B (PPP1R3B), mRNA.	79660 TR(1..5564)CDS(166..1023)	3261..3267,3680..3686,3074..3080,5395..5401	1.47	0.74	-0.91	-0.33	1 No
NM_004004	223278_at	GJB2	Homo sapiens gap junction protein, beta 2, 26kDa (GJB2), mRNA.	2706 TR(1..2331)CDS(216..896)	1390..1397	1.25	0.82	-0.79	-0.65	1 No
NM_181425	205565_s_at	FXN	Homo sapiens frataxin (FXN), nuclear gene encoding mitochondrial protein, transcript variant 2, mRNA.	2395 TR(1..2270)CDS(356..946)	1789..1795	1.54	0.71	0.06	0	1 No
NM_015392	218086_at	NPDC1	Homo sapiens neural proliferation, differentiation and control, 1 (NPDC1), mRNA.	56654 TR(1..1554)CDS(215..1192)	59..65	1.96	0.66	0.18	0.04	1 No
NM_017839	239598_s_at	AYTL1	Homo sapiens acyltransferase like 1 (AYTL1), mRNA.	54947 TR(1..1937)CDS(141..1775)	1607..1613	1.99	0.66	-0.08	0.13	1 No
NM_182540	227485_at	DDX26B	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 26B (DDX26B), mRNA.	203522 TR(1..3793)CDS(335..2920)	3091..3097,3383..3389,1491..1497	1.9	0.65	-0.17	-0.01	1 No
NM_003821	209544_at,209545_s_at	RIPK2	Homo sapiens receptor-interacting serine-threonine kinase 2 (RIPK2), mRNA.	8767 TR(1..2588)CDS(315..1937)	1677..1683,2202..2209	1.53	0.7	-0.91	-0.18	1 No
NM_016227	203429_s_at	C1orf9	Homo sapiens chromosome 1 open reading frame 9 (C1orf9), transcript variant 2, mRNA.	51430 TR(1..5919)CDS(125..4342)	4707..4713	2.05	0.64	-1.74	-0.56	1 No
NM_021951	220493_at	DMRT1	Homo sapiens doublesex and mab-3 related transcription factor 1 (DMRT1), mRNA.	1761 TR(1..2229)CDS(150..1271)	1636..1643	2.32	0.64	-1.2	-0.36	1 No
NM_012458	218188_s_at	TIMM13	Homo sapiens translocase of inner mitochondrial membrane 13 homolog (yeast) (TIMM13), nuclear gene encoding mitochondrial protein, mRNA.	26517 TR(1..1977)CDS(345..632)	829..835	1.55	0.68	-0.06	-0.03	1 Yes
NM_019605	229674_at	SERTAD4	Homo sapiens SERTA domain containing 4 (SERTAD4), mRNA.	56256 TR(1..2038)CDS(180..1250)	1925..1932	1.39	0.73	-0.35	-0.15	1 No
NM_017448	207022_s_at	LDHC	Homo sapiens lactate dehydrogenase C (LDHC), transcript variant 2, mRNA.	3948 TR(1..1171)CDS(59..1057)	34..40	2.23	0.63	-0.26	0.04	1 No
NM_001081491	208922_s_at	NXF1	Homo sapiens nuclear RNA export factor 1 (NXF1), transcript variant 2, mRNA.	10482 TR(1..4147)CDS(137..1207)	3698..3704	1.48	0.7	0.11	0.07	1 No
NM_138471	227292_at	LOC144097	Homo sapiens hypothetical protein BC007540 (LOC144097), mRNA.	144097 TR(1..2004)CDS(237..1382)	1719..1726	0.85	1.2	-0.04	-0.56	1 No
NM_182574	1557091_at	FLJ36070	Homo sapiens likely ortholog of MEF2-activating SAP transcriptional regulator (FLJ36070), mRNA.	284358 TR(1..1825)CDS(616..1554)	1714..1720	0.84	1.2	0.27	-0.19	1 No
NM_001001664	225658_at,225659_at	LOC339745	Homo sapiens hypothetical protein LOC339745 (LOC339745), mRNA.	339745 TR(1..5721)CDS(380..1558)	4723..4729,4321..4327,1821..1827	1.92	0.62	-1.81	-0.57	1 No
NM_007271	1553117_a_at,202951_at	STK38	Homo sapiens serine/threonine kinase 38 (STK38), mRNA.	11329 TR(1..3593)CDS(287..1684)	2946..2952	1.85	0.62	-1.61	-0.36	1 No
NM_032497	224518_s_at	ZNF559	Homo sapiens zinc finger protein 559 (ZNF559), mRNA.	84527 TR(1..3059)CDS(648..2264)	2697..2703,2244..2250,960..966,425..431	1.99	0.62	-0.21	0.02	1 No
NM_015627	57082_at,221790_s_at	LDLRAP1	Homo sapiens low density lipoprotein receptor adaptor protein 1 (LDLRAP1), mRNA.	26119 TR(1..2935)CDS(115..1041)	2210..2216	1.13	0.82	-0.58	-0.33	1 No
NM_005110	205100_at	GFPT2	Homo sapiens glutamine-fructose-6-phosphate transaminase 2 (GFPT2), mRNA.	9945 TR(1..3011)CDS(99..2147)	2283..2289	1.14	0.8	-0.53	-0.01	1 No
NM_000696	201612_at	ALDH9A1	Homo sapiens aldehyde dehydrogenase 9 family, member A1 (ALDH9A1), mRNA.	223 TR(1..2500)CDS(106..1662)	1843..1849	1.4	0.7	-0.55	-0.24	1 No
NM_138452	213279_at	DHRS1	Homo sapiens dehydrogenase/reductase (SDR family) member 1 (DHRS1), mRNA.	115817 TR(1..1470)CDS(270..1211)	1382..1388,1417..1423	1.35	0.72	-0.69	-0.39	1 Yes
NM_033389	1555423_at,1554114_s_at	SSH2	Homo sapiens slingshot homolog 2 (Drosophila) (SSH2), mRNA.	85464 TR(1..9166)CDS(1..4272)	8801..8807	0.77	0.97	-0.28	-0.23	1 No
NM_016647	218500_at	C8orf55	Homo sapiens chromosome 8 open reading frame 55 (C8orf55), mRNA.	51337 TR(1..2264)CDS(145..771)	2179..2185	1.38	0.7	-0.23	-0.07	1 Yes
NM_005841	212558_at	SPRY1	Homo sapiens sprouty homolog 1, antagonist of FGF signaling (Drosophila) (SPRY1), transcript variant 1, mRNA.	10252 TR(1..2612)CDS(323..1282)	2210..2216,943..949	1.25	0.75	0.01	-0.01	1 No
NM_024815	219665_at	NUDT18	Homo sapiens nudix (nucleoside diphosphate linked moiety X)-type motif 18 (NUDT18), mRNA.	79873 TR(1..1519)CDS(120..1091)	1273..1279	0.95	0.86	-0.03	-0.23	1 No
NM_002826	201482_at	QSOX1	Homo sapiens quiescin Q6 sulfhydryl oxidase 1 (QSOX1), transcript variant 1, mRNA.	5768 TR(1..3316)CDS(76..2319)	3239..3246,3018..3024	1.99	0.58	-0.86	-0.32	1 No
NM_020648	219201_s_at,225406_at	TWSG1	Homo sapiens twisted gastrulation homolog 1 (Drosophila) (TWSG1), mRNA.	57045 TR(1..3693)CDS(106..777)	2482..2488,1216..1222	1.6	0.62	-1.51	-0.63	1 No
NM_001083589	221586_s_at	E2F5	Homo sapiens E2F transcription factor 5, p130-binding (E2F5), transcript variant 3, mRNA.	1875 TR(1..1594)CDS(372..929)	1422..1428	1.41	0.68	-1.53	-0.85	1 Yes
NM_016016	223649_s_at	SLC25A39	Homo sapiens solute carrier family 25, member 39 (SLC25A39), mRNA.	51629 TR(1..1556)CDS(118..1173)	1242..1248	1.29	0.73	-0.25	-0.11	1 No
NM_005943	211673_s_at	MOCS1	Homo sapiens molybdenum cofactor synthesis 1 (MOCS1), transcript variant 1, mRNA.	4337 TR(1..3037)CDS(139..1296)	1533..1540	0.68	1.27	0.02	-0.62	1 Yes
NM_001718	206176_at	BMP6	Homo sapiens bone morphogenetic protein 6 (BMP6), mRNA.	654 TR(1..3105)CDS(179..1720)	1842..1849,1775..1781	1.99	0.57	-0.8	-0.17	1 No
NM_153367	212423_at,212419_at	C10orf56	Homo sapiens chromosome 10 open reading frame 56 (C10orf56), mRNA.	219654 TR(1..4945)CDS(188..913)	4846..4853,1636..1642	1.37	0.69	-0.74	-0.52	1 Yes
NM_001076786	226265_at,219705_at,229982_at,244563_at	QSER1	Homo sapiens glutamine and serine rich 1 (QSER1), mRNA.	79832 TR(1..9271)CDS(268..5475)	6752..6758,6862..6869,6617..6623	1.54	0.63	-0.85	-0.68	1 No
NM_031439	228698_at,224013_s_at	SOX7	Homo sapiens SRY (sex determining region Y)-box 7 (SOX7), mRNA.	83595 TR(1..3219)CDS(80..1246)	1293..1299	1.35	0.69	-0.16	-0.08	1 No
NM_152293	226240_at	MGC21874	Homo sapiens transcriptional adaptor 2 (ADA2 homolog, yeast)-beta (MGC21874), mRNA.	93624 TR(1..4312)CDS(152..1414)	2918..2924	1.08	0.79	-0.85	-0.25	1 No

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NM_024539	219263_at	RNF128	Homo sapiens ring finger protein 128 (RNF128), transcript variant 2, mRNA.	79589 TR(1..2688)CDS(166..1374)	2384..2391	1.46	0.64	-1.13	-0.48	1 No
NM_152261	224759_s_at	C12orf23	Homo sapiens chromosome 12 open reading frame 23 (C12orf23), mRNA.	90488 TR(1..3167)CDS(214..564)	1159..1165,436..442	1.77	0.56	-1.25	-0.5	1 No
NM_001020818	225673_at	MYADM	Homo sapiens myeloid-associated differentiation marker (MYADM), transcript variant 1, mRNA.	91663 TR(1..3243)CDS(315..1283)	1965..1972	1.83	0.55	-0.79	-0.19	1 Yes
NM_172037	226021_at,1552378_s_at	RDH10	Homo sapiens retinol dehydrogenase 10 (all-trans) (RDH10), mRNA.	157506 TR(1..2751)CDS(189..1214)	2509..2515	1.2	0.73	-0.74	-0.47	1 No
NM_018593	222939_s_at	SLC16A10	Homo sapiens solute carrier family 16, member 10 (aromatic amino acid transporter) (SLC16A10), mRNA.	117247 TR(1..2488)CDS(148..1695)	1072..1079	1.59	0.58	-0.2	-0.11	1 No
NM_005857	202939_at	ZMPSTE24	Homo sapiens zinc metalloproteinase (STE24 homolog, S. cerevisiae) (ZMPSTE24), mRNA.	10269 TR(1..3154)CDS(212..1639)	3086..3092	1.58	0.58	-1.09	-0.19	1 No
NM_024544	218246_at	C1orf166	Homo sapiens chromosome 1 open reading frame 166 (C1orf166), mRNA.	79594 TR(1..2442)CDS(144..1202)	330..337	1.52	0.6	0.08	0.08	1 No
NM_079834	213244_at,235073_at	SCAMP4	Homo sapiens secretory carrier membrane protein 4 (SCAMP4), mRNA.	113178 TR(1..2548)CDS(108..797)	1948..1955	1.46	0.62	-0.35	-0.02	1 No
NM_013256	219495_s_at	ZNF180	Homo sapiens zinc finger protein 180 (ZNF180), mRNA.	7733 TR(1..3100)CDS(266..2344)	2120..2126	1.34	0.66	-0.61	-0.14	1 No
NM_001002257	226996_at	LYCAT	Homo sapiens lysocardiolipin acyltransferase (LYCAT), transcript variant 2, mRNA.	253558 TR(1..4872)CDS(119..1249)	4567..4573,4388..4395	1.54	0.58	-1.43	-0.51	1 No
NM_013322	218404_at	SNX10	Homo sapiens sorting nexin 10 (SNX10), mRNA.	29887 TR(1..2591)CDS(216..821)	662..668	1.65	0.54	-0.9	-0.39	0.99 No
NM_006700	243196_s_at,202837_at,35254_at	TRAFD1	Homo sapiens TRAF-type zinc finger domain containing 1 (TRAFD1), mRNA.	10906 TR(1..2618)CDS(55..1803)	1001..1008	1.38	0.63	0.12	-0.03	0.99 No
NM_001010984	223511_at	C1orf124	Homo sapiens chromosome 1 open reading frame 124 (C1orf124), transcript variant 2, mRNA.	83932 TR(1..3854)CDS(449..1201)	2908..2914,3158..3164	1.22	0.7	-0.26	-0.12	0.99 No
NM_004529	204917_s_at,204918_s_at	MLLT3	Homo sapiens myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 3 (MLLT3), mRNA.	4300 TR(1..3439)CDS(260..1966)	2349..2355,906..912	1.86	0.51	-0.65	-0.18	0.99 No
NM_002508	202007_at,202008_s_at	NID1	Homo sapiens nidogen 1 (NID1), mRNA.	4811 TR(1..5903)CDS(103..3846)	4642..4649,3855..3862	1.14	0.72	-1.22	-0.51	0.99 Yes
NM_080652	235037_at	TMEM41A	Homo sapiens transmembrane protein 41A (TMEM41A), mRNA.	90407 TR(1..1395)CDS(54..848)	340..346	1.17	0.71	-0.8	-0.42	0.99 No
NM_000249	202520_s_at	MLH1	Homo sapiens mutL homolog 1, colon cancer, nonpolyposis type 2 (E. coli) (MLH1), mRNA.	4292 TR(1..2524)CDS(61..2331)	1824..1830	1.73	0.51	-0.42	-0.12	0.99 No
NM_001560	201887_at,210904_s_at,211612_s_at,201888_s_at	IL13RA1	Homo sapiens interleukin 13 receptor, alpha 1 (IL13RA1), mRNA.	3597 TR(1..4006)CDS(44..1327)	701..707	1.25	0.66	-0.21	-0.03	0.99 No
NM_007218	209510_at	RNF139	Homo sapiens ring finger protein 139 (RNF139), mRNA.	11236 TR(1..3312)CDS(344..2338)	1411..1417	1.42	0.59	-0.44	-0.13	0.99 Yes
NM_001986	211603_s_at,1554576_a_at	ETV4	Homo sapiens ets variant gene 4 (E1A enhancer binding protein, E1AF) (ETV4), transcript variant 1, mRNA.	2118 TR(1..2298)CDS(129..1583)	497..503	1.75	0.49	-0.23	-0.11	0.99 No
NM_024683	219146_at	C17orf42	Homo sapiens chromosome 17 open reading frame 42 (C17orf42), mRNA.	79736 TR(1..1357)CDS(71..1153)	844..850	1.94	0.47	-0.98	-0.02	0.99 No
NM_145018	228281_at	C11orf82	Homo sapiens chromosome 11 open reading frame 82 (C11orf82), mRNA.	220042 TR(1..3545)CDS(213..3209)	3346..3352,165..171	1.88	0.47	-1.89	-0.44	0.99 No
NM_002524	202647_s_at	NRAS	Homo sapiens neuroblastoma RAS viral (v-ras) oncogene homolog (NRAS), mRNA.	4893 TR(1..1963)CDS(254..823)	1474..1480	1.03	0.74	-0.71	-0.4	0.99 Yes
NM_022725	222713_s_at,218689_at	FANCF	Homo sapiens Fanconi anemia, complementation group F (FANCF), mRNA.	2188 TR(1..3309)CDS(32..1156)	2217..2223	1.04	0.73	-0.06	-0.03	0.99 No
NM_152305	218587_s_at,222681_at	KTELC1	Homo sapiens KTEL (Lys-Tyr-Glu-Leu) containing 1 (KTELC1), mRNA.	56983 TR(1..3538)CDS(71..1249)	1919..1925	1.66	0.5	-1.41	-0.29	0.99 No
NM_032478	225103_at	MRPL38	Homo sapiens mitochondrial ribosomal protein L38 (MRPL38), nuclear gene encoding mitochondrial protein, mRNA.	64978 TR(1..1907)CDS(621..1661)	1755..1761	1.08	0.71	0.15	0.01	0.99 No
NM_021994	1555193_a_at,218645_at	ZNF277P	Homo sapiens zinc finger protein 277 pseudogene (ZNF277P), mRNA.	11179 TR(1..2687)CDS(130..1482)	301..307	1.79	0.47	-0.86	-0.16	0.99 No
NM_001015880	203060_s_at,203058_s_at,203059_s_at	PAPSS2	Homo sapiens 3'-phosphoadenosine 5'-phosphosulfate synthase 2 (PAPSS2), transcript variant 2, mRNA.	9060 TR(1..3874)CDS(264..2123)	3720..3727	1.37	0.6	-1.15	-0.52	0.99 Yes
NM_006373	208626_s_at	VAT1	Homo sapiens vesicle amine transport protein 1 homolog (T. californica) (VAT1), mRNA.	10493 TR(1..2758)CDS(121..1302)	1994..2001,2088..2094	1.3	0.62	0.25	0.08	0.99 No
NM_152743	225437_s_at	C7orf27	Homo sapiens chromosome 7 open reading frame 27 (C7orf27), mRNA.	221927 TR(1..2713)CDS(43..2508)	2159..2165	1.7	0.48	-0.08	0.12	0.99 No
NM_017423	218313_s_at,222587_s_at	GALNT7	Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 7 (GalNAc-T7) (GALNT7), mRNA.	51809 TR(1..4339)CDS(84..2057)	2322..2328	1.67	0.48	-1.73	-0.5	0.99 No
NM_013241	218530_at	FHOD1	Homo sapiens formin homology 2 domain containing 1 (FHOD1), mRNA.	29109 TR(1..3865)CDS(113..3607)	3474..3481	0.79	0.79	-0.07	-0.02	0.99 No
NM_003141	204804_at	TRIM21	Homo sapiens tripartite motif-containing 21 (TRIM21), mRNA.	6737 TR(1..1946)CDS(114..1541)	1420..1426	1.84	0.45	-0.36	-0.01	0.99 No
NM_005484	214086_s_at	PARP2	Homo sapiens poly (ADP-ribose) polymerase family, member 2 (PARP2), transcript variant 1, mRNA.	10038 TR(1..1904)CDS(29..1780)	1128..1134	1.53	0.51	-0.33	-0.15	0.99 No
NM_004184	200629_at	WARS	Homo sapiens tryptophanyl-tRNA synthetase (WARS), transcript variant 1, mRNA.	7453 TR(1..2884)CDS(382..1797)	1293..1300	1.4	0.57	-0.13	0.03	0.99 No
NM_175902	64438_at,219254_at,53071_s_at	FLJ22222	Homo sapiens hypothetical protein FLJ22222 (FLJ22222), transcript variant 2, mRNA.	79701 TR(1..1474)CDS(92..1087)	1363..1370	1.62	0.48	-0.16	-0.04	0.99 No
NM_181846	228769_at	ZSCAN22	Homo sapiens zinc finger and SCAN domain containing 22 (ZSCAN22), mRNA.	342945 TR(1..4648)CDS(148..1623)	2721..2727,2078..2084	3.24	0.43	-1.42	0	0.99 No
NM_022091	212815_at,231269_at	ASCC3	Homo sapiens activating signal cointegrator 1 complex subunit 3 (ASCC3), transcript variant 2, mRNA.	10973 TR(1..3470)CDS(330..665)	2959..2965	1.72	0.45	-1.51	-0.31	0.99 No
NM_181050	212849_at	AXIN1	Homo sapiens axin 1 (AXIN1), transcript variant 2, mRNA.	8312 TR(1..3369)CDS(178..2658)	3091..3097	1.11	0.68	-0.08	0.19	0.99 No
NM_207361	230964_at	FREM2	Homo sapiens FRAS1 related extracellular matrix protein 2 (FREM2), mRNA.	341640 TR(1..16179)CDS(310..9819)	12974..12980,14795..14801,550..556,4385..4391	1.43	0.55	-1.46	-0.56	0.99 No
NM_203374	228364_at	ZNF784	Homo sapiens zinc finger protein 784 (ZNF784), mRNA.	147808 TR(1..2006)CDS(15..986)	214..220	1.37	0.45	0.45	0.13	0.99 No
NM_004538	204749_at	NAP1L3	Homo sapiens nucleosome assembly protein 1-like 3 (NAP1L3), mRNA.	4675 TR(1..2680)CDS(306..1826)	2286..2292	1.59	0.48	-1.42	0.04	0.99 No
NM_145648	225043_at,225057_at	SLC15A4	Homo sapiens solute carrier family 15, member 4 (SLC15A4), mRNA.	121260 TR(1..2802)CDS(54..1787)	1955..1962,817..823	1.29	0.61	-0.82	-0.43	0.99 No
NM_017851	219034_at	PARP16	Homo sapiens poly (ADP-ribose) polymerase family, member 16 (PARP16), mRNA.	54956 TR(1..2551)CDS(255..1226)	1263..1270	1.31	0.6	-0.75	-0.09	0.99 No
NM_199072	217599_s_at,211675_s_at	MDFIC	Homo sapiens MyoD family inhibitor domain containing (MDFIC), mRNA.	29969 TR(1..4607)CDS(264..1331)	2515..2521,2675..2681,2540..2546	0.66	0.81	-0.72	-0.67	0.99 No
NM_018062	218397_at	FANCL	Homo sapiens Fanconi anemia, complementation group L (FANCL), mRNA.	55120 TR(1..1750)CDS(67..1194)	345..351	1.87	0.43	-0.7	-0.09	0.99 No
NM_001839	201445_at	CNN3	Homo sapiens calponin 3, acidic (CNN3), mRNA.	1266 TR(1..2045)CDS(252..1241)	1368..1374	1.46	0.52	-0.51	-0.05	0.99 No
NM_024613	222699_s_at,218640_s_at	PLEKHF2	Homo sapiens pleckstrin homology domain containing, family F (with FYVE domain) member 2 (PLEKHF2), mRNA.	79666 TR(1..2905)CDS(248..997)	1483..1490	1.05	0.69	-0.7	-0.19	0.99 No
XM_001126650	206240_s_at	ZNF136	PREDICTED: Homo sapiens zinc finger protein 136 (ZNF136), mRNA.	7695 TR(1..3223)CDS(141..1343)	1434..1440	1.51	0.5	-0.4	-0.27	0.98 No
NM_012331	219281_at	MSRA	Homo sapiens methionine sulfoxide reductase A (MSRA), mRNA.	4482 TR(1..1502)CDS(213..920)	1285..1292	1.68	0.44	-0.04	0.05	0.98 No
NM_007054	228680_at,213623_at	KIF3A	Homo sapiens kinesin family member 3A (KIF3A), mRNA.	11127 TR(1..6257)CDS(155..2254)	1037..1044,6029..6035	1.84	0.42	-1.35	-0.17	0.98 No
NM_144578	212643_at,212644_s_at,225643_at	C14orf32	Homo sapiens chromosome 14 open reading frame 32 (C14orf32), mRNA.	93487 TR(1..6469)CDS(165..902)	843..849,348..354,6334..6340	1.46	0.52	-0.7	-0.2	0.98 Yes
NM_001211	203755_at	BUB1B	Homo sapiens BUB1 budding uninhibited by benzimidazoles 1 homolog beta (yeast) (BUB1B), mRNA.	701 TR(1..3721)CDS(185..3337)	72..78	1.63	0.45	-0.65	0	0.98 No

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NM_174898	236706_at	LYG1	Homo sapiens lysozyme G-like 1 (LYG1), mRNA.	129530 TR(1..1015)CDS(251..835)	163..169	0.85	0.75	0.02	-0.09	0.98 No
NM_003094	231112_at	SNRPE	Homo sapiens small nuclear ribonucleoprotein polypeptide E (SNRPE), mRNA.	6635 TR(1..1593)CDS(49..327)	898..904,1282..1288	2.79	0.4	-0.8	0.01	0.98 No
NM_020428	224609_at,225175_s_at	SLC44A2	Homo sapiens solute carrier family 44, member 2 (SLC44A2), mRNA.	57153 TR(1..3389)CDS(82..2202)	2770..2776	1.28	0.59	-0.13	0.11	0.98 No
NM_007057	204026_s_at	ZWINT	Homo sapiens ZW10 interactor (ZWINT), transcript variant 1, mRNA.	11130 TR(1..1687)CDS(38..871)	1395..1402	1.17	0.63	-0.53	-0.16	0.98 Yes
NM_022769	218648_at	CRTC3	Homo sapiens CREB regulated transcription coactivator 3 (CRTC3), transcript variant 1, mRNA.	64784 TR(1..5171)CDS(107..1968)	4229..4236,2948..2955	1.18	0.63	-0.72	-0.4	0.98 Yes
NM_133367	226423_at,227626_at	PAQR8	Homo sapiens progesterin and adipoQ receptor family member VIII (PAQR8), mRNA.	85315 TR(1..4641)CDS(156..1220)	2724..2731	1.81	0.4	-0.12	-0.01	0.98 No
NM_015412	225279_s_at,225281_at	C3orf17	Homo sapiens chromosome 3 open reading frame 17 (C3orf17), transcript variant 1, mRNA.	25871 TR(1..4857)CDS(62..1765)	702..708,3179..3186	1.79	0.4	-0.27	-0.01	0.98 No
NM_005665	208297_s_at,209717_at	EVI5	Homo sapiens ecotropic viral integration site 5 (EVI5), mRNA.	7813 TR(1..7403)CDS(11..2443)	6818..6824,7361..7367	1.16	0.64	-0.78	-0.35	0.98 No
NM_020639	221215_s_at,234730_s_at	RIPK4	Homo sapiens receptor-interacting serine-threonine kinase 4 (RIPK4), mRNA.	54101 TR(1..3890)CDS(49..2403)	3783..3789,3584..3590	0.86	0.73	-0.07	-0.11	0.98 Yes
NM_002498	211089_s_at,213116_at	NEK3	Homo sapiens NIMA (never in mitosis gene a)-related kinase 3 (NEK3), transcript variant 1, mRNA.	4752 TR(1..2396)CDS(377..1897)	709..715,906..912	1.5	0.48	-0.27	0.03	0.98 No
NM_173515	242394_at,227481_at	CNKSR3	Homo sapiens CNKSR family member 3 (CNKSR3), mRNA.	154043 TR(1..3277)CDS(506..2173)	2215..2221	1.27	0.58	-0.55	-0.23	0.98 Yes
NM_006754	201260_s_at,201259_s_at	SYPL1	Homo sapiens synaptophysin-like 1 (SYPL1), transcript variant 1, mRNA.	6856 TR(1..2183)CDS(83..862)	2068..2074,1013..1019	1.23	0.6	-0.88	-0.45	0.98 Yes
NM_153442	244617_at	GPR26	Homo sapiens G protein-coupled receptor 26 (GPR26), mRNA.	2849 TR(1..7512)CDS(53..1066)	4324..4330	0.01	0.87	0.14	-0.33	0.98 No
NM_207352	226745_at,228391_at	CYP4V2	Homo sapiens cytochrome P450, family 4, subfamily V, polypeptide 2 (CYP4V2), mRNA.	285440 TR(1..4718)CDS(305..1882)	1898..1904,888..895	1.62	0.42	-0.75	-0.17	0.98 No
NM_147156	212989_at	SGMS1	Homo sapiens sphingomyelin synthase 1 (SGMS1), mRNA.	259230 TR(1..3767)CDS(955..2196)	2940..2946	1.5	0.47	-1.17	-0.39	0.98 No
NM_001139	207381_at	ALOX12B	Homo sapiens arachidonate 12-lipoxygenase, 12R type (ALOX12B), mRNA.	242 TR(1..2502)CDS(262..2367)	2304..2310	1.46	0.49	-0.06	0.01	0.98 No
NM_033035	235737_at	TSLP	Homo sapiens thymic stromal lymphopietin (TSLP), transcript variant 1, mRNA.	85480 TR(1..2652)CDS(200..679)	511..517,211..217,422..428	1.87	0.37	-0.36	-0.22	0.98 No
NM_003270	209109_s_at,209108_at	TSPAN6	Homo sapiens tetraspanin 6 (TSPAN6), mRNA.	7105 TR(1..2069)CDS(104..841)	1786..1792	1.27	0.57	-0.36	-0.14	0.98 Yes
NM_001018160	202268_s_at	APPBP1	Homo sapiens amyloid beta precursor protein binding protein 1 (APPBP1), transcript variant 3, mRNA.	8883 TR(1..1716)CDS(241..1578)	899..906	1.57	0.43	-0.86	0.06	0.98 No
NM_022460	219020_at	HS1BP3	Homo sapiens HCLS1 binding protein 3 (HS1BP3), mRNA.	64342 TR(1..2403)CDS(42..1220)	2277..2284	1.46	0.48	-0.23	0.08	0.98 Yes
NM_178868	235099_at	CMTM8	Homo sapiens CKLF-like MARVEL transmembrane domain containing 8 (CMTM8), mRNA.	152189 TR(1..1185)CDS(295..816)	635..641	1.18	0.61	0.18	-0.07	0.97 No
NM_032139	221522_at	ANKRD27	Homo sapiens ankyrin repeat domain 27 (VPS9 domain) (ANKRD27), mRNA.	84079 TR(1..4461)CDS(157..3309)	4392..4398	1.43	0.49	-0.89	-0.28	0.97 Yes
NM_030784	220993_s_at	GPR63	Homo sapiens G protein-coupled receptor 63 (GPR63), mRNA.	81491 TR(1..1892)CDS(208..1467)	1139..1145	0.94	0.7	-0.06	0.01	0.97 No
NM_003051	209900_s_at,202236_s_at,202234_s_at,202235_at,1557918_s_at	SLC16A1	Homo sapiens solute carrier family 16, member 1 (monocarboxylic acid transporter 1) (SLC16A1), mRNA.	6566 TR(1..3927)CDS(364..1866)	2144..2151	1.37	0.52	-1.03	-0.4	0.97 No
NM_052940	215084_s_at	LRRC42	Homo sapiens leucine rich repeat containing 42 (LRRC42), mRNA.	115353 TR(1..1714)CDS(201..1487)	1071..1077	1.25	0.58	-0.27	-0.01	0.97 No
NM_016422	226106_at	RNF141	Homo sapiens ring finger protein 141 (RNF141), mRNA.	50862 TR(1..4084)CDS(154..846)	3611..3617	1.16	0.61	-1.1	-0.34	0.97 No
NM_016565	220647_s_at	CHCHD8	Homo sapiens coiled-coil-helix-coiled-coil-helix domain containing 8 (CHCHD8), mRNA.	51287 TR(1..833)CDS(106..369)	734..740	0.83	0.72	-0.04	-0.06	0.97 Yes
NM_178422	242123_at	PAQR7	Homo sapiens progesterin and adipoQ receptor family member VII (PAQR7), mRNA.	164091 TR(1..3058)CDS(668..1708)	1125..1131,1029..1035	0.91	0.7	-0.06	-0.37	0.97 No
NM_024528	223439_at	NKAP	Homo sapiens NFkB activating protein (NKAP), mRNA.	79576 TR(1..1608)CDS(168..1415)	1526..1532	1.73	0.36	-1.47	-0.29	0.97 No
NM_024496	223474_at	C14orf4	Homo sapiens chromosome 14 open reading frame 4 (C14orf4), mRNA.	64207 TR(1..4166)CDS(900..3290)	4105..4111	0.88	0.7	0.11	-0.11	0.97 No
NM_058172	1555536_at,225524_at	ANTXR2	Homo sapiens anthrax toxin receptor 2 (ANTXR2), mRNA.	118429 TR(1..3881)CDS(540..2006)	2555..2562,2503..2509	1.74	0.36	-1.1	-0.15	0.97 No
NM_001966	205222_at	EHHADH	Homo sapiens enoyl-Coenzyme A, hydratase/3-hydroxyacyl Coenzyme A dehydrogenase (EHHADH), mRNA.	1962 TR(1..3821)CDS(28..2199)	3686..3692	1.04	0.65	-0.6	-0.43	0.97 No
NM_001042482	221218_s_at,223686_at	TPK1	Homo sapiens thiamin pyrophosphokinase 1 (TPK1), transcript variant 2, mRNA.	27010 TR(1..2302)CDS(104..688)	412..419,719..725	1.51	0.43	-0.19	0.01	0.97 No
NM_032435	228565_at	KIAA1804	Homo sapiens mixed lineage kinase 4 (KIAA1804), mRNA.	84451 TR(1..5809)CDS(262..3372)	3378..3385,1434..1440,3247..3253	1.91	0.33	-1.57	-0.4	0.97 No
NM_018719	225081_s_at	CDCA7L	Homo sapiens cell division cycle associated 7-like (CDCA7L), mRNA.	55536 TR(1..2874)CDS(85..1449)	1832..1839	1.43	0.46	-0.79	-0.28	0.96 Yes
NM_033285	244721_at,225912_at	TP53NP1	Homo sapiens tumor protein p53 inducible nuclear protein 1 (TP53NP1), mRNA.	94241 TR(1..5587)CDS(357..1079)	347..353,5243..5249,3342..3348	1.09	0.62	-1.09	-0.42	0.96 No
NM_001083962	203753_at,212387_at,212382_at,222146_s_at,228837_at,212386_at,212385_at,213891_s_at	TCF4	Homo sapiens transcription factor 4 (TCF4), transcript variant 1, mRNA.	6925 TR(1..8332)CDS(613..2628)	3717..3723,636..642	1.33	0.51	-1.08	-0.49	0.96 No
NM_014278	205543_at	HSPA4L	Homo sapiens heat shock 70kDa protein 4-like (HSPA4L), mRNA.	22824 TR(1..3284)CDS(254..2773)	1607..1614	1.78	0.32	-1.1	-0.09	0.96 No
NM_000159	208369_s_at,203500_at	GCDH	Homo sapiens glutaryl-Coenzyme A dehydrogenase (GCDH), nuclear gene encoding mitochondrial protein, transcript variant 1, mRNA.	2639 TR(1..1839)CDS(78..1394)	1761..1768	0.67	0.74	-0.01	-0.08	0.96 No
NM_001271	1554015_a_at,1554014_at,225077_at	CHD2	Homo sapiens chromodomain helicase DNA binding protein 2 (CHD2), transcript variant 1, mRNA.	1106 TR(1..9374)CDS(576..6062)	209..215,35..42,5120..5127	1.96	0.3	-0.65	0.01	0.96 No
NM_001098634	229439_s_at,222496_s_at,229440_at,218035_s_at	FLJ20273	Homo sapiens RNA-binding protein (FLJ20273), transcript variant 1, mRNA.	54502 TR(1..5160)CDS(711..2492)	3658..3664,2721..2727,3221..3228	1.14	0.59	-0.78	-0.43	0.96 Yes
NM_015251	201855_s_at	ASCIZ	Homo sapiens ATM/ATR-Substrate Chk2-Interacting Zn2+-finger protein (ASCIZ), mRNA.	23300 TR(1..4866)CDS(19..2490)	4807..4813,4500..4506,3909..3915	1.53	0.4	-1.21	-0.25	0.96 No
NM_001040615	231976_at,1554456_a_at,228348_at	LINS1	Homo sapiens lines homolog 1 (Drosophila) (LINS1), transcript variant 5, mRNA.	55180 TR(1..2468)CDS(187..2460)	1101..1107,1149..1155,636..642	1.52	0.4	-0.49	-0.19	0.96 No
NM_019857	219080_s_at	CTPS2	Homo sapiens CTP synthase II (CTPS2), transcript variant 1, mRNA.	56474 TR(1..3844)CDS(255..2015)	203..209,2358..2364,1491..1497	0.53	0.77	-0.48	-0.2	0.96 No
NM_183238	227822_at	ZNF605	Homo sapiens zinc finger protein 605 (ZNF605), mRNA.	90462 TR(1..6238)CDS(326..2251)	4976..4982,2860..2866	2.21	0.29	-1.64	-0.01	0.96 No
NM_022371	233851_s_at	TOR3A	Homo sapiens tor sin family 3, member A (TOR3A), mRNA.	64222 TR(1..2138)CDS(153..1346)	1614..1621	0.7	0.73	0.26	0.12	0.95 No
NM_001100877	226846_at	PHYHD1	Homo sapiens phytanoyl-CoA dioxygenase domain containing 1 (PHYHD1), transcript variant 3, mRNA.	254295 TR(1..1200)CDS(42..854)	916..922	0.87	0.68	-0.18	-0.01	0.95 No
NM_212554	226631_at	LOC399818	Homo sapiens similar to CG9643-PA (LOC399818), mRNA.	399818 TR(1..2638)CDS(38..913)	1337..1343	1.17	0.57	-0.75	-0.18	0.95 No
NM_148173	207621_s_at	PEMT	Homo sapiens phosphatidylethanolamine N-methyltransferase (PEMT), nuclear gene encoding mitochondrial protein, transcript variant 3, mRNA.	10400 TR(1..905)CDS(68..667)	791..797	1.24	0.53	0.01	-0.01	0.95 No
NM_017762	225810_at	MTMR10	Homo sapiens myotubularin related protein 10 (MTMR10), mRNA.	54893 TR(1..4969)CDS(99..2432)	1865..1871,3078..3084	1.85	0.28	-0.73	-0.14	0.95 No
NM_014021	203016_s_at,203015_s_at,203018_s_at,203017_s_at	SSX2IP	Homo sapiens synovial sarcoma, X breakpoint 2 interacting protein (SSX2IP), mRNA.	117178 TR(1..5658)CDS(267..2111)	1756..1762,427..433	1.48	0.4	-0.85	-0.18	0.95 No
NM_198321	207357_s_at	GALNT10	Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetyl(galactosaminyl)transferase 10 (GalNAc-T10) (GALNT10), transcript variant 1, mRNA.	55568 TR(1..5222)CDS(133..1944)	4016..4022,3955..3961	1.13	0.58	-0.25	-0.06	0.95 No
NM_001753	203065_s_at,212097_at	CAV1	Homo sapiens caveolin 1, caveolae protein, 22kDa (CAV1), mRNA.	857 TR(1..2704)CDS(279..815)	1470..1476,1393..1399	1.37	0.46	-1.08	-0.44	0.95 Yes

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NM_005389	210156_s_at,205202_at,208857_s_at	PCMT1	Homo sapiens protein-L-isoaspartate (D-aspartate) O-methyltransferase (PCMT1), mRNA.	5110 TR(1..1599)CDS(74..757)	713..719	1.28	0.5	-0.38	-0.01	0.95 No
NM_022070	65493_at,218991_at	HEATR6	Homo sapiens HEAT repeat containing 6 (HEATR6), mRNA.	63897 TR(1..3935)CDS(18..3563)	1640..1646	1.31	0.48	-0.31	0.1	0.94 No
NM_018179	218987_at	ATF7IP	Homo sapiens activating transcription factor 7 interacting protein (ATF7IP), mRNA.	55729 TR(1..4661)CDS(159..3971)	4599..4606	1.27	0.5	-0.59	-0.15	0.94 No
NM_019000	218532_s_at	FAM134B	Homo sapiens family with sequence similarity 134, member B (FAM134B), transcript variant 2, mRNA.	54463 TR(1..3108)CDS(311..1381)	2921..2927	1.58	0.33	-1.46	-0.41	0.94 No
NM_017546	224695_at	C2orf29	Homo sapiens chromosome 2 open reading frame 29 (C2orf29), mRNA.	55571 TR(1..2590)CDS(50..1582)	2452..2458	1.13	0.57	-0.77	-0.28	0.94 No
NM_144689	238937_at	ZNF420	Homo sapiens zinc finger protein 420 (ZNF420), mRNA.	147923 TR(1..2990)CDS(216..2282)	2705..2711	1.23	0.52	-1.23	-0.32	0.94 No
NM_017672	223324_s_at	TRPM7	Homo sapiens transient receptor potential cation channel, subfamily M, member 7 (TRPM7), mRNA.	54822 TR(1..7255)CDS(265..5862)	2425..2431,2152..2158,4322..4329	1.26	0.5	-1.4	-0.48	0.94 No
NM_005903	225219_at,225223_at,205187_at,235451_at,205188_s_at	SMAD5	Homo sapiens SMAD family member 5 (SMAD5), transcript variant 1, mRNA.	4090 TR(1..7012)CDS(361..1758)	3653..3659,6433..6439,2822..2828	1.26	0.5	-1.2	-0.77	0.94 No
NM_001100590	232366_at,212441_at	KIAA0232	Homo sapiens KIAA0232 (KIAA0232), transcript variant 2, mRNA.	9778 TR(1..7789)CDS(380..4567)	2463..2469	1.25	0.5	-0.95	-0.17	0.94 No
NM_000214	209098_s_at,209097_s_at,216268_s_at	JAG1	Homo sapiens jagged 1 (Alagille syndrome) (JAG1), mRNA.	182 TR(1..5896)CDS(414..4070)	2899..2905,5378..5384,4496..4502	1.3	0.47	-0.13	-0.12	0.93 No
NM_016651	219179_at	DACT1	Homo sapiens dapper, antagonist of beta-catenin, homolog 1 (Xenopus laevis) (DACT1), transcript variant 1, mRNA.	51339 TR(1..3877)CDS(165..2675)	2841..2847	1.3	0.47	-0.84	-0.21	0.93 No
NM_144599	1552696_at,225752_at	NIPA1	Homo sapiens non imprinted in Prader-Willi/Angelman syndrome 1 (NIPA1), mRNA.	123606 TR(1..6565)CDS(26..1015)	1673..1679,632..638	1.89	0.22	-1.08	-0.07	0.93 No
NM_133452	223425_at	RAVER1	Homo sapiens ribonucleoprotein, PTB-binding 1 (RAVER1), mRNA.	125950 TR(1..3606)CDS(81..2351)	2694..2700,2999..3005	1.23	0.51	-0.49	-0.17	0.93 No
NM_032449	242487_at,225323_at	CC2D1B	Homo sapiens coiled-coil and C2 domain containing 1B (CC2D1B), mRNA.	200014 TR(1..5639)CDS(139..2715)	3371..3377,3515..3521,1909..1915	0.75	0.69	-0.53	-0.49	0.93 No
NM_000182	208630_at	HADHA	Homo sapiens hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), alpha subunit (HADHA), nuclear gene encoding mitochondrial protein, mRNA.	3030 TR(1..3048)CDS(131..2422)	2558..2564	1.1	0.56	-0.7	-0.12	0.93 Yes
NM_001414	201632_at	EIF2B1	Homo sapiens eukaryotic translation initiation factor 2B, subunit 1 alpha, 26kDa (EIF2B1), mRNA.	1967 TR(1..1794)CDS(144..1061)	1487..1493	1.19	0.52	-0.63	-0.15	0.93 No
NM_018323	222631_at	PI4K2B	Homo sapiens phosphatidylinositol 4-kinase type 2 beta (PI4K2B), mRNA.	55300 TR(1..3458)CDS(112..1557)	2287..2293,3090..3096,2017..2023	2.11	0.2	-1.97	-0.19	0.93 No
NM_003060	205074_at	SLC22A5	Homo sapiens solute carrier family 22 (organic cation transporter), member 5 (SLC22A5), mRNA.	6584 TR(1..3252)CDS(222..1895)	1143..1149,3185..3191	1.54	0.31	-1.3	-0.13	0.92 No
NM_007215	205811_at	POLG2	Homo sapiens polymerase (DNA directed), gamma 2, accessory subunit (POLG2), mRNA.	11232 TR(1..1595)CDS(84..1541)	646..652,511..517	1.48	0.35	-0.42	-0.14	0.92 No
NM_004957	202945_at	FPGS	Homo sapiens folypolyglutamate synthase (FPGS), nuclear gene encoding mitochondrial protein, transcript variant 1, mRNA.	2356 TR(1..2487)CDS(51..1814)	2182..2188,382..388,1984..1990,1885..1891	1.19	0.51	-0.18	-0.13	0.92 No
NM_016536	206648_at	ZNF571	Homo sapiens zinc finger protein 571 (ZNF571), mRNA.	51276 TR(1..2291)CDS(102..1931)	558..564	1.83	0.2	-1.62	0.02	0.92 No
NM_178544	240429_at	ZNF546	Homo sapiens zinc finger protein 546 (ZNF546), mRNA.	339327 TR(1..4566)CDS(171..2681)	3777..3783	1.37	0.39	-0.25	-0.03	0.9 No
NM_006361	230105_at,209844_at	HOXB13	Homo sapiens homeobox B13 (HOXB13), mRNA.	10481 TR(1..3047)CDS(157..1011)	388..395	1.45	0.34	0.03	0.07	0.9 Yes
NM_002372	205105_at	MAN2A1	Homo sapiens mannosidase, alpha, class 2A, member 1 (MAN2A1), mRNA.	4124 TR(1..5128)CDS(964..4398)	1746..1752	1.53	0.29	-1.51	-0.33	0.9 No
NM_031990	212016_s_at	PTBP1	Homo sapiens polypyrimidine tract binding protein 1 (PTBP1), transcript variant 2, mRNA.	5725 TR(1..3300)CDS(88..1740)	2691..2697,2069..2076,3078..3084	0.99	0.58	-0.72	-0.39	0.9 No
NM_032825	1561687_a_at,1557260_a_at	ZNF382	Homo sapiens zinc finger protein 382 (ZNF382), mRNA.	84911 TR(1..2830)CDS(114..1766)	1273..1279	1.33	0.41	-0.75	-0.03	0.9 No
NM_002206	216331_at,209663_s_at	ITGA7	Homo sapiens integrin, alpha 7 (ITGA7), mRNA.	3679 TR(1..4079)CDS(162..3575)	280..286,3914..3921	1.52	0.28	-0.07	-0.01	0.89 No
NM_030802	221249_s_at	FAM117A	Homo sapiens family with sequence similarity 117, member A (FAM117A), mRNA.	81558 TR(1..2398)CDS(82..1443)	1480..1486	0.73	0.66	-0.21	-0.15	0.88 No
NM_152444	230774_at	ZADH1	Homo sapiens zinc binding alcohol dehydrogenase, domain containing 1 (ZADH1), mRNA.	145482 TR(1..2538)CDS(201..1256)	987..994	1.94	0.08	-0.35	0.01	0.88 No
NM_012179	1554423_a_at,201178_at	FBXO7	Homo sapiens F-box protein 7 (FBXO7), transcript variant 1, mRNA.	25793 TR(1..2153)CDS(284..1852)	1040..1046	1.26	0.44	-0.19	-0.07	0.88 No
NM_201629	202085_at	TJP2	Homo sapiens tight junction protein 2 (zona occludens 2) (TJP2), transcript variant 2, mRNA.	9414 TR(1..4177)CDS(201..3332)	3786..3792	1.36	0.38	-0.97	-0.35	0.88 No
NM_024618	1553695_a_at,219680_at	NLRX1	Homo sapiens NLR family member X1 (NLRX1), transcript variant 1, mRNA.	79671 TR(1..3761)CDS(240..3167)	3587..3593,3731..3737	0.95	0.58	-0.74	0.01	0.88 No
NM_018374	222787_s_at,226529_at,218930_s_at	TMEM106B	Homo sapiens transmembrane protein 106B (TMEM106B), mRNA.	54664 TR(1..6456)CDS(323..1147)	4583..4589,6004..6010	0.54	0.69	-1	-1.07	0.88 No
NM_152238	205573_s_at	SNX7	Homo sapiens sorting nexin 7 (SNX7), transcript variant 2, mRNA.	51375 TR(1..1585)CDS(183..1193)	1115..1121	1.31	0.41	-0.56	-0.08	0.88 No
NM_018298	220484_at	MCOLN3	Homo sapiens mucolipin 3 (MCOLN3), mRNA.	55283 TR(1..2813)CDS(94..1755)	1440..1446	2.07	0.03	-1.51	0.28	0.88 No
NM_020453	213238_at	ATP10D	Homo sapiens ATPase, class V, type 10D (ATP10D), mRNA.	57205 TR(1..6550)CDS(165..4445)	3801..3808	2.44	0.01	-1.55	-0.05	0.87 No
NM_024579	219506_at	C1orf54	Homo sapiens chromosome 1 open reading frame 54 (C1orf54), mRNA.	79630 TR(1..517)CDS(24..419)	385..391	1.1	0.51	0.11	0.29	0.87 No
NM_032204	215684_s_at	ASCC2	Homo sapiens activating signal cointegrator 1 complex subunit 2 (ASCC2), mRNA.	84164 TR(1..2799)CDS(106..2379)	2568..2575	0.96	0.57	-0.35	-0.04	0.87 No
NM_004339	200677_at	PTTG1IP	Homo sapiens pituitary tumor-transforming 1 interacting protein (PTTG1IP), mRNA.	754 TR(1..2737)CDS(211..753)	2691..2698	0.88	0.6	-0.69	-0.38	0.87 No
NM_015254	202962_at	KIF13B	Homo sapiens kinesin family member 13B (KIF13B), mRNA.	23303 TR(1..8774)CDS(60..5540)	5936..5942,5546..5552,8464..8470,8309..8315,4662..4668,302..308	1.17	0.48	-0.24	-0.31	0.87 Yes
NM_144598	227423_at	LRRC28	Homo sapiens leucine rich repeat containing 28 (LRRC28), mRNA.	123355 TR(1..1464)CDS(156..1259)	449..455,620..626	0.98	0.56	-0.03	-0.01	0.87 No
NM_000112	224959_at,224963_at,205097_at	SLC26A2	Homo sapiens solute carrier family 26 (sulfate transporter), member 2 (SLC26A2), mRNA.	1836 TR(1..8082)CDS(269..2488)	1198..1204,772..779,7847..7853,6750..6756	1.7	0.08	-1.53	-0.49	0.86 No
NM_022776	222586_s_at,218304_s_at	OSBPL11	Homo sapiens oxysterol binding protein-like 11 (OSBPL11), mRNA.	114885 TR(1..4206)CDS(306..2549)	3859..3866	1.07	0.51	-0.88	-0.49	0.85 No
NM_006275	208804_s_at	SFRS6	Homo sapiens splicing factor, arginine/serine-rich 6 (SFRS6), mRNA.	6431 TR(1..3738)CDS(157..1191)	3597..3603,1679..1685	1.3	0.39	-0.62	-0.04	0.85 No
NM_001497	201883_s_at,216627_s_at	B4GALT1	Homo sapiens UDP-Gal:betaGlcNAc beta 1,4-galactosyltransferase, polypeptide 1 (B4GALT1), mRNA.	2683 TR(1..4080)CDS(73..1269)	2732..2738,3360..3367,3111..3117	0.9	0.58	-0.66	-0.36	0.84 Yes
NM_005808	201904_s_at,201905_s_at,201906_s_at	CTDSPL	Homo sapiens CTD (carboxy-terminal domain, RNA polymerase II, polypeptide A) small phosphatase-like (CTDSPL), transcript variant 2, mRNA.	10217 TR(1..4422)CDS(23..820)	2778..2784,1218..1224,2400..2406	1.08	0.5	-0.78	-0.39	0.84 Yes
NM_013276	219713_at	CARKL	Homo sapiens carbohydrate kinase-like (CARKL), mRNA.	23729 TR(1..3838)CDS(104..1540)	316..322,1745..1751,3226..3233	0.78	0.61	-0.12	-0.5	0.83 No
NM_002198	202531_at	IRF1	Homo sapiens interferon regulatory factor 1 (IRF1), mRNA.	3659 TR(1..2035)CDS(198..1175)	1851..1857	0.84	0.6	-0.33	-0.25	0.83 No
NM_017966	219053_s_at,1560059_at,1560060_s_at	VPS37C	Homo sapiens vacuolar protein sorting 37 homolog C (S. cerevisiae) (VPS37C), mRNA.	55048 TR(1..2711)CDS(61..1128)	2402..2408,2017..2023	0.78	0.61	-0.38	-0.38	0.83 Yes
NM_015554	213552_at	GLCE	Homo sapiens glucuronic acid epimerase (GLCE), mRNA.	26035 TR(1..5043)CDS(229..2082)	3970..3977	1.36	0.33	-0.9	-0.06	0.83 No
NM_007219	204669_s_at,210706_s_at	RNF24	Homo sapiens ring finger protein 24 (RNF24), mRNA.	11237 TR(1..1692)CDS(253..699)	561..568	1.07	0.5	-0.07	0.02	0.83 No
NM_134433	208031_s_at,226872_at	RFX2	Homo sapiens regulatory factor X, 2 (influences HLA class II expression) (RFX2), transcript variant 2, mRNA.	5990 TR(1..3940)CDS(160..2258)	3934..3940,1598..1604	0.87	0.58	-0.03	-0.01	0.83 No
NM_194356	213434_at,207346_at	STX2	Homo sapiens syntaxin 2 (STX2), transcript variant 2, mRNA.	2054 TR(1..3469)CDS(168..1034)	2143..2149	1.07	0.5	-0.71	-0.42	0.82 No

NM_018285	221688_s_at	IMP3	Homo sapiens IMP3, U3 small nucleolar ribonucleoprotein, homolog (yeast) (IMP3), mRNA.	55272 TR(1..1160)CDS(87..641)	859..865	1.19	0.43	-0.04	-0.08	0.82 No
NM_001748	208683_at	CAPN2	Homo sapiens calpain 2, (m/II) large subunit (CAPN2), mRNA.	824 TR(1..3492)CDS(225..2327)	1677..1683,3253..3259	1.18	0.43	-0.5	-0.09	0.81 No
NM_001196	211725_s_at,204493_at,227143_s_at	BID	Homo sapiens BH3 interacting domain death agonist (BID), transcript variant 2, mRNA.	637 TR(1..2214)CDS(171..758)	1035..1041	0.81	0.59	-0.45	-0.19	0.81 No
NM_080670	224626_at	SLC35A4	Homo sapiens solute carrier family 35, member A4 (SLC35A4), mRNA.	113829 TR(1..2676)CDS(729..1703)	1808..1814	0.99	0.52	-0.45	-0.24	0.8 Yes
NM_144720	238600_at	JAKMIP1	Homo sapiens janus kinase and microtubule interacting protein 1 (JAKMIP1), transcript variant 2, mRNA.	152789 TR(1..2593)CDS(487..2367)	2471..2477	0.95	0.54	0.33	-0.01	0.8 No
NM_012216	208384_s_at	MID2	Homo sapiens midline 2 (MID2), transcript variant 1, mRNA.	11043 TR(1..2523)CDS(258..2405)	841..848	1.45	0.23	-0.18	0.19	0.8 No
NM_005981	203226_s_at,203227_s_at	TSPAN31	Homo sapiens tetraspanin 31 (TSPAN31), mRNA.	6302 TR(1..1749)CDS(155..787)	774..780	0.99	0.52	0.1	0.09	0.79 No
NM_033083	226952_at	EAF1	Homo sapiens ELL associated factor 1 (EAF1), mRNA.	85403 TR(1..4503)CDS(224..1030)	3240..3246	1.13	0.45	-0.81	-0.36	0.79 No
NM_018346	218307_at	RSAD1	Homo sapiens radical S-adenosyl methionine domain containing 1 (RSAD1), mRNA.	55316 TR(1..2499)CDS(57..1385)	2339..2345	1.1	0.46	0.07	0.07	0.79 No
NM_133496	239596_at,226217_at	SLC30A7	Homo sapiens solute carrier family 30 (zinc transporter), member 7 (SLC30A7), mRNA.	148867 TR(1..5862)CDS(142..1272)	3537..3543,1870..1876	1.16	0.43	-1.6	-0.56	0.79 No
NM_145869	214783_s_at,206200_s_at	ANXA11	Homo sapiens annexin A11 (ANXA11), transcript variant c, mRNA.	311 TR(1..2731)CDS(484..2001)	2650..2657	0.56	0.65	-0.11	-0.2	0.78 Yes
NM_022736	218109_s_at	MFSD1	Homo sapiens major facilitator superfamily domain containing 1 (MFSD1), mRNA.	64747 TR(1..2161)CDS(31..1428)	713..719	1.37	0.28	-0.93	-0.17	0.78 Yes
NM_007146	202172_at	VEZF1	Homo sapiens vascular endothelial zinc finger 1 (VEZF1), mRNA.	7716 TR(1..4652)CDS(144..1709)	3565..3571	1.22	0.39	-0.5	-0.19	0.78 No
NM_006987	221614_s_at	RPH3AL	Homo sapiens rabphilin 3A-like (without C2 domains) (RPH3AL), mRNA.	9501 TR(1..2610)CDS(287..1234)	1386..1392,84..90	1.51	0.12	-0.06	-0.01	0.78 No
NM_022833	223019_at,233974_s_at	FAM129B	Homo sapiens family with sequence similarity 129, member B (FAM129B), transcript variant 1, mRNA.	64855 TR(1..4007)CDS(203..2443)	2532..2538,3846..3852	0.77	0.59	-0.32	-0.33	0.78 No
NM_002835	202006_at,216915_s_at	PTPN12	Homo sapiens protein tyrosine phosphatase, non-receptor type 12 (PTPN12), mRNA.	5782 TR(1..3161)CDS(30..2372)	2580..2586,2517..2523,3018..3024	1.08	0.47	-0.8	-0.16	0.77 No
NM_014585	223044_at	SLC40A1	Homo sapiens solute carrier family 40 (iron-regulated transporter), member 1 (SLC40A1), mRNA.	30061 TR(1..3457)CDS(428..2143)	2257..2263,3257..3263	1.04	0.48	-1.02	-0.27	0.77 No
NM_002655	205372_at	PLAG1	Homo sapiens pleiomorphic adenoma gene 1 (PLAG1), mRNA.	5324 TR(1..7313)CDS(481..1983)	4977..4984	1.07	0.47	-0.93	-0.56	0.77 No
NM_002593	202465_at	PCOLCE	Homo sapiens procollagen C-endopeptidase enhancer (PCOLCE), mRNA.	5118 TR(1..1510)CDS(91..1440)	1348..1354	1.33	0.3	-0.11	-0.01	0.77 Yes
NM_024034	219668_at	GDAP1L1	Homo sapiens ganglioside-induced differentiation-associated protein 1-like 1 (GDAP1L1), mRNA.	78997 TR(1..2244)CDS(68..1171)	2155..2161,858..864	0.22	0.69	-0.07	-0.41	0.77 No
NM_032488	224329_s_at	CNFN	Homo sapiens cornifelin (CNFN), mRNA.	84518 TR(1..538)CDS(51..389)	188..194	1.51	0.1	0.71	-0.01	0.76 Yes
NM_175063	224727_at	C19orf63	Homo sapiens chromosome 19 open reading frame 63 (C19orf63), transcript variant HSS1, mRNA.	284361 TR(1..2048)CDS(67..831)	1312..1318	0.99	0.5	-0.08	0.02	0.76 No
NM_015470	210879_s_at	RAB11FIP5	Homo sapiens RAB11 family interacting protein 5 (class I) (RAB11FIP5), mRNA.	26056 TR(1..4362)CDS(242..2203)	3108..3114	1.1	0.45	-0.3	-0.07	0.76 No
NM_030794	1569069_s_at,208089_s_at	TDRD3	Homo sapiens tudor domain containing 3 (TDRD3), mRNA.	81550 TR(1..2501)CDS(258..2213)	531..537	1.36	0.27	-0.68	0.14	0.75 No
NM_005443	209043_at	PAPSS1	Homo sapiens 3-phosphoadenosine 5-phosphosulfate synthase 1 (PAPSS1), mRNA.	9061 TR(1..2558)CDS(85..1959)	2105..2111,951..957	1.18	0.4	-0.17	0	0.75 No
NM_005573	203276_at	LMNB1	Homo sapiens lamin B1 (LMNB1), mRNA.	4001 TR(1..2882)CDS(369..2129)	278..284	0.52	0.64	-0.65	-0.82	0.75 No
NM_018439	218637_at,222698_s_at	IMPACT	Homo sapiens Impact homolog (mouse) (IMPACT), mRNA.	55364 TR(1..3680)CDS(40..1002)	1313..1319	1.34	0.28	-0.96	-0.14	0.75 No
NM_001406	205031_at	EFNB3	Homo sapiens ephrin-B3 (EFNB3), mRNA.	1949 TR(1..3236)CDS(398..1420)	2341..2347	0.66	0.61	0.04	-0.29	0.74 No
NM_057178	1552651_a_at,228980_at,1562603_at,1552649_a_at	RFFL	Homo sapiens ring finger and FYVE-like domain containing 1 (RFFL), transcript variant 1, mRNA.	117584 TR(1..4143)CDS(180..1271)	1293..1300,2581..2587	0.67	0.61	-0.14	-0.06	0.74 Yes
NM_152390	229302_at	TMEM178	Homo sapiens transmembrane protein 178 (TMEM178), mRNA.	130733 TR(1..1648)CDS(26..919)	1440..1446	1.27	0.33	-0.24	0	0.74 No
NM_022743	218788_s_at	SMYD3	Homo sapiens SET and MYND domain containing 3 (SMYD3), mRNA.	64754 TR(1..1477)CDS(128..1237)	215..221,1281..1287	1.07	0.45	-0.05	-0.02	0.74 No
NM_005226	231741_at,228176_at	EDG3	Homo sapiens endothelial differentiation, sphingolipid G-protein-coupled receptor, 3 (EDG3), mRNA.	1903 TR(1..4221)CDS(397..1533)	1716..1722	1.22	0.37	-0.89	-0.22	0.74 No
NM_016551	222477_s_at,217974_at	TM7SF3	Homo sapiens transmembrane 7 superfamily member 3 (TM7SF3), mRNA.	51768 TR(1..2527)CDS(38..1750)	1469..1475	1.02	0.48	-0.17	-0.1	0.73 No
NM_198391	219250_s_at,222853_at	FLRT3	Homo sapiens fibronectin leucine rich transmembrane protein 3 (FLRT3), transcript variant 2, mRNA.	23767 TR(1..3977)CDS(468..2417)	3731..3737	0.81	0.56	-0.13	-0.08	0.72 No
NM_004945	202253_s_at	DNM2	Homo sapiens dynamin 2 (DNM2), transcript variant 3, mRNA.	1785 TR(1..3621)CDS(165..2765)	3336..3342	1.14	0.4	-0.64	-0.25	0.71 No
NM_198977	203055_s_at	ARHGEF1	Homo sapiens Rho guanine nucleotide exchange factor (GEF) 1 (ARHGEF1), transcript variant 3, mRNA.	9138 TR(1..3137)CDS(110..2749)	1573..1579,1620..1626,3048..3054	1.41	0.18	-0.61	-0.01	0.71 No
NM_153605	214030_at	DKFZp667G2110	Homo sapiens hypothetical protein DKFZp667G2110 (DKFZp667G2110), mRNA.	131544 TR(1..4802)CDS(65..3133)	4740..4746	1.03	0.46	-1.49	-0.26	0.71 No
NM_133487	205023_at,205024_s_at	RAD51	Homo sapiens RAD51 homolog (RecA homolog, E. coli) (S. cerevisiae) (RAD51), transcript variant 2, mRNA.	5888 TR(1..1976)CDS(268..996)	1269..1276	0.87	0.53	-0.58	-0.36	0.7 No
NM_002114	204512_at	HIVEP1	Homo sapiens human immunodeficiency virus type 1 enhancer binding protein 1 (HIVEP1), mRNA.	3096 TR(1..8891)CDS(180..8336)	8526..8532	1	0.47	-0.77	-0.19	0.7 No
NM_002069	227692_at,209576_at	GNAI1	Homo sapiens guanine nucleotide binding protein (G protein), alpha inhibiting activity polypeptide 1 (GNAI1), mRNA.	2770 TR(1..3318)CDS(338..1402)	3080..3086	1.08	0.43	-0.42	-0.16	0.69 No
NM_021070	219922_s_at	LTBP3	Homo sapiens latent transforming growth factor beta binding protein 3 (LTBP3), mRNA.	4054 TR(1..4064)CDS(1..3771)	2628..2634	1.23	0.33	0.31	0.16	0.69 No
NM_004275	206961_s_at,212872_s_at	MED20	Homo sapiens mediator complex subunit 20 (MED20), mRNA.	9477 TR(1..2478)CDS(122..760)	371..377,973..980	1.39	0.18	-0.76	-0.28	0.69 No
NM_001779	211744_s_at,216942_s_at	CD58	Homo sapiens CD58 molecule (CD58), mRNA.	965 TR(1..1040)CDS(10..762)	154..160	1.14	0.39	-0.5	0.01	0.69 No
NM_024704	219570_at	C20orf23	Homo sapiens chromosome 20 open reading frame 23 (C20orf23), mRNA.	55614 TR(1..5274)CDS(159..4112)	1080..1087,1019..1025,4144..4150	0.52	0.62	-0.17	-0.15	0.69 No
NM_006335	201821_s_at,215171_s_at,201822_at	TIMM17A	Homo sapiens translocase of inner mitochondrial membrane 17 homolog A (yeast) (TIMM17A), mRNA.	10440 TR(1..1645)CDS(8..523)	294..301	1.32	0.25	-0.42	-0.03	0.68 No
NM_145257	225904_at,1553698_a_at,1553697_at	C1orf96	Homo sapiens chromosome 1 open reading frame 96 (C1orf96), mRNA.	126731 TR(1..5174)CDS(125..937)	591..598,2818..2824	1.21	0.34	-0.18	0.02	0.68 No
NM_175839	217074_at,1555680_a_at,210357_s_at	SMOX	Homo sapiens spermine oxidase (SMOX), transcript variant 1, mRNA.	54498 TR(1..2203)CDS(202..1869)	2132..2138	0.85	0.53	0.03	-0.05	0.68 No
NM_005397	201578_at	PODXL	Homo sapiens podocalyxin-like (PODXL), transcript variant 2, mRNA.	5420 TR(1..5911)CDS(259..1839)	2262..2268	1.29	0.27	-0.79	-0.27	0.68 No
NM_015055	209307_at,209306_s_at	SWAP70	Homo sapiens SWAP-70 protein (SWAP70), mRNA.	23075 TR(1..4848)CDS(100..1857)	3759..3765,308..314	1.25	0.31	-1.17	-0.27	0.67 No
NM_017554	224701_at	PARP14	Homo sapiens poly (ADP-ribose) polymerase family, member 14 (PARP14), mRNA.	54625 TR(1..7713)CDS(60..5465)	6586..6592,1840..1847,6298..6304	1.01	0.45	-0.79	-0.1	0.67 No
NM_002305	201105_at	LGALS1	Homo sapiens lectin, galactoside-binding, soluble, 1 (galectin 1) (LGALS1), mRNA.	3956 TR(1..586)CDS(98..505)	145..151	1.43	0.09	0.48	-0.01	0.67 No
NM_017842	218417_s_at,48106_at	FLJ20489	Homo sapiens hypothetical protein FLJ20489 (FLJ20489), mRNA.	55652 TR(1..1950)CDS(483..1202)	1874..1880	0.61	0.59	-0.01	-0.1	0.66 No
NM_001099650	226868_at,232021_at	GLT8D3	Homo sapiens glycosyltransferase 8 domain containing 3 (GLT8D3), transcript variant 2, mRNA.	283464 TR(1..7416)CDS(226..1455)	4639..4646,7300..7306,1956..1962,5835..5841	1.21	0.33	-1.05	-0.44	0.66 No

NM_203352	203370_s_at,214266_s_at	PDLIM7	Homo sapiens PDZ and LIM domain 7 (enigma) (PDLIM7), transcript variant 2, mRNA.	9260 TR(1..1607)CDS(86..1357)	1447..1453	0.67	0.58	0.3	-0.12	0.66 No
NM_017686	219473_at	GDAP2	Homo sapiens ganglioside induced differentiation associated protein 2 (GDAP2), mRNA.	54834 TR(1..2879)CDS(203..1696)	2675..2682	1.01	0.45	-1.11	-0.15	0.66 No
NM_006742	213141_at	PSKH1	Homo sapiens protein serine kinase H1 (PSKH1), mRNA.	5681 TR(1..3460)CDS(131..1405)	2151..2157,1544..1550,3096..3102,1919..1925	1.4	0.13	-0.85	-0.13	0.65 Yes
NM_000901	205259_at	NR3C2	Homo sapiens nuclear receptor subfamily 3, group C, member 2 (NR3C2), mRNA.	4306 TR(1..5747)CDS(215..3169)	4244..4250,3313..3319,5260..5266	1.37	0.17	-1.04	-0.08	0.65 No
NM_000140	203115_at,203116_s_at	FECH	Homo sapiens ferrochelatase (protoporphyrin) (FECH), nuclear gene encoding mitochondrial protein, transcript variant 2, mRNA.	2235 TR(1..3817)CDS(139..1410)	2107..2113	1.11	0.39	-0.91	-0.43	0.64 No
NM_024819	221224_s_at,224522_s_at	DCAKD	Homo sapiens dephospho-CoA kinase domain containing (DCAKD), mRNA.	79877 TR(1..2070)CDS(246..941)	1677..1683,928..934	0.92	0.48	0.21	-0.17	0.63 Yes
NM_012413	205174_s_at	QPCT	Homo sapiens glutaminyl-peptide cyclotransferase (glutaminyl cyclase) (QPCT), mRNA.	25797 TR(1..1719)CDS(123..1208)	637..644	1.05	0.41	-0.17	-0.12	0.63 Yes
NM_024493	211773_s_at	ZKSCAN3	Homo sapiens zinc finger with KRAB and SCAN domains 3 (ZKSCAN3), mRNA.	80317 TR(1..2242)CDS(149..1765)	2185..2191	0.97	0.46	0.1	0.3	0.63 No
NM_014953	214194_at,234660_s_at,222607_s_at	DIS3	Homo sapiens DIS3 mitotic control homolog (S. cerevisiae) (DIS3), mRNA.	22894 TR(1..7320)CDS(37..2913)	5316..5322,3844..3850,7116..7122	1.28	0.25	-1.14	-0.1	0.63 No
NM_032557	223288_at,223289_s_at	USP38	Homo sapiens ubiquitin specific peptidase 38 (USP38), mRNA.	84640 TR(1..4684)CDS(507..3635)	4395..4401	1.04	0.42	-1.18	-0.36	0.62 No
NM_147157	214484_s_at,201692_at	OPRS1	Homo sapiens opioid receptor, sigma 1 (OPRS1), transcript variant 2, mRNA.	10280 TR(1..1580)CDS(75..653)	1178..1184,353..359	0.95	0.46	0.12	0.03	0.62 No
NM_005596	209290_s_at,213032_at,211467_s_at,213033_s_at,209289_at,213029_at	NFIB	Homo sapiens nuclear factor I/B (NFIB), mRNA.	4781 TR(1..8159)CDS(436..1698)	3181..3187,2122..2128	1.36	0.14	-1.42	-0.36	0.61 No
NM_012339	218693_at	TSPAN15	Homo sapiens tetraspanin 15 (TSPAN15), mRNA.	23555 TR(1..1726)CDS(126..1010)	295..301,1336..1342,1500..1506	1.08	0.39	-0.09	-0.09	0.61 Yes
NM_001003962	200001_at	CAPNS1	Homo sapiens calpain, small subunit 1 (CAPNS1), transcript variant 2, mRNA.	826 TR(1..1489)CDS(155..961)	876..882,1401..1407,741..747	1.07	0.39	-0.19	-0.04	0.6 No
NM_001099642	222811_at	FLJ11171	Homo sapiens hypothetical protein FLJ11171 (FLJ11171), transcript variant 2, mRNA.	55783 TR(1..4040)CDS(407..2719)	2098..2104	1.35	0.14	-0.81	0.01	0.6 No
NM_021649	228234_at,239431_at	TICAM2	Homo sapiens toll-like receptor adaptor molecule 2 (TICAM2), mRNA.	353376 TR(1..3453)CDS(848..1555)	1426..1432	1.36	0.12	-0.89	-0.2	0.6 No
NM_018075	218910_at	TMEM16K	Homo sapiens transmembrane protein 16K (TMEM16K), mRNA.	55129 TR(1..2743)CDS(172..2154)	1770..1776	1.36	0.13	-0.17	0.05	0.59 No
NM_153371	227569_at	LNK2	Homo sapiens ligand of numb-protein X 2 (LNK2), mRNA.	222484 TR(1..4804)CDS(310..2382)	4712..4718	1.29	0.22	-0.79	-0.19	0.59 No
NM_016224	223028_s_at,223027_at	SNX9	Homo sapiens sorting nexin 9 (SNX9), mRNA.	51429 TR(1..4200)CDS(174..1961)	3290..3297	1.12	0.36	-0.68	-0.26	0.59 No
NM_175871	227352_at	C19orf39	Homo sapiens chromosome 19 open reading frame 39 (C19orf39), mRNA.	126074 TR(1..1661)CDS(38..727)	382..388,1360..1366	0.88	0.48	0.29	0.13	0.58 No
NM_000523	236681_at,207398_at,207397_s_at	HOXD13	Homo sapiens homeobox D13 (HOXD13), mRNA.	3239 TR(1..2341)CDS(88..1119)	928..935	1.06	0.39	-0.38	-0.1	0.58 No
NM_198507	226752_at	TMEM157	Homo sapiens transmembrane protein 157 (TMEM157), mRNA.	345757 TR(1..1260)CDS(121..693)	698..704	0.86	0.49	-0.36	-0.34	0.58 No
NM_016524	205613_at	SYT17	Homo sapiens synaptotagmin XVII (SYT17), mRNA.	51760 TR(1..1912)CDS(332..1756)	1214..1220	0.49	0.59	0.04	-0.13	0.58 No
NM_013388	217861_s_at	PREB	Homo sapiens prolactin regulatory element binding (PREB), mRNA.	10113 TR(1..2211)CDS(254..1507)	1485..1491	0.96	0.44	-0.08	-0.01	0.57 Yes
NM_001105214	209517_s_at	ASH2L	Homo sapiens ash2 (absent, small, or homeotic)-like (Drosophila) (ASH2L), transcript variant 2, mRNA.	9070 TR(1..2662)CDS(404..2008)	625..632	1.15	0.33	-0.11	0.02	0.57 No
NM_183075	226402_at,226393_at	CYP2U1	Homo sapiens cytochrome P450, family 2, subfamily U, polypeptide 1 (CYP2U1), mRNA.	113612 TR(1..4760)CDS(84..1718)	4539..4545,3397..3403,3627..3633,2103..2109	1.12	0.34	-1.08	-0.27	0.57 Yes
NM_173797	225761_at,238706_at	PAPD4	Homo sapiens PAP associated domain containing 4 (PAPD4), mRNA.	167153 TR(1..2755)CDS(300..1754)	2270..2276,618..624	1.3	0.17	-0.61	-0.06	0.56 No
NM_005245	201579_at	FAT	Homo sapiens FAT tumor suppressor homolog 1 (Drosophila) (FAT), mRNA.	2195 TR(1..14773)CDS(189..13955)	12310..12316	1.06	0.38	-0.7	-0.06	0.56 No
NM_138782	1555153_s_at,228220_at	FCHO2	Homo sapiens FCH domain only 2 (FCHO2), mRNA.	115548 TR(1..4909)CDS(26..2458)	3215..3221	1.02	0.4	-0.91	-0.39	0.54 No
NM_199320	235024_at,218517_at,225820_at	PHF17	Homo sapiens PHD finger protein 17 (PHF17), transcript variant L, mRNA.	79960 TR(1..5771)CDS(281..2809)	5724..5730,1904..1910,4954..4960	1.14	0.32	-0.61	-0.17	0.54 No
NM_170714	204346_s_at	RASSF1	Homo sapiens Ras association (RalGDS/AF-6) domain family 1 (RASSF1), transcript variant D, mRNA.	11186 TR(1..1979)CDS(132..1166)	1705..1712	0.97	0.42	0.04	0.01	0.53 No
NM_173552	228079_at,226464_at	C3orf58	Homo sapiens chromosome 3 open reading frame 58 (C3orf58), mRNA.	205428 TR(1..4091)CDS(263..1555)	4013..4019	0.85	0.47	-0.49	-0.17	0.53 No
NM_020962	227870_at	NOPE	Homo sapiens neighbor of Punc E11 (NOPE), mRNA.	57722 TR(1..6485)CDS(211..3963)	6259..6265	0.84	0.48	-0.7	-0.49	0.53 Yes
NM_001039933	205297_s_at	CD79B	Homo sapiens CD79b molecule, immunoglobulin-associated beta (CD79B), transcript variant 3, mRNA.	974 TR(1..1303)CDS(84..776)	1221..1227	0.09	0.63	-0.05	-0.11	0.52 No
NM_023925	1555608_at,218456_at	CAPRN2	Homo sapiens caprin family member 2 (CAPRN2), transcript variant 2, mRNA.	65981 TR(1..4351)CDS(752..3985)	1133..1139	1.18	0.27	-0.93	-0.09	0.52 No
NM_000254	203774_at	MTR	Homo sapiens 5-methyltetrahydrofolate-homocysteine methyltransferase (MTR), mRNA.	4548 TR(1..7122)CDS(287..4084)	4142..4149	1.1	0.34	-1.29	-0.31	0.52 No
NM_014824	203620_s_at	FCHSD2	Homo sapiens FCH and double SH3 domains 2 (FCHSD2), mRNA.	9873 TR(1..4341)CDS(240..2294)	2466..2472	1	0.4	-1.08	-0.35	0.52 No
NM_170662	209682_at	CBLB	Homo sapiens Cas-Br-M (murine) ecotropic retroviral transforming sequence b (CBLB), mRNA.	868 TR(1..3976)CDS(323..3271)	2063..2069,919..925,1508..1514	1.27	0.17	0.05	0.02	0.51 No
NM_052932	211967_at	TMEM123	Homo sapiens transmembrane protein 123 (TMEM123), mRNA.	114908 TR(1..3605)CDS(422..1048)	593..599	1.24	0.21	-0.93	-0.14	0.51 Yes
NM_014881	209804_at	DCLRE1A	Homo sapiens DNA cross-link repair 1A (PSO2 homolog, S. cerevisiae) (DCLRE1A), mRNA.	9937 TR(1..4270)CDS(667..3789)	1853..1859	1.35	0.02	-1.23	0.06	0.51 No
NM_001343	201280_s_at,201278_at,201279_s_at	DAB2	Homo sapiens disabled homolog 2, mitogen-responsive phosphoprotein (Drosophila) (DAB2), mRNA.	1601 TR(1..4597)CDS(532..2844)	2570..2576,2445..2451,2970..2976	1.3	0.12	-0.78	-0.16	0.51 No
NM_021908	207871_s_at,207524_at	ST7	Homo sapiens suppression of tumorigenicity 7 (ST7), transcript variant b, mRNA.	7982 TR(1..2970)CDS(215..1972)	332..338,2852..2858	0.44	0.58	-0.21	-0.04	0.51 No
NM_021101	222549_at,218182_s_at	CLDN1	Homo sapiens claudin 1 (CLDN1), mRNA.	9076 TR(1..3445)CDS(221..856)	2762..2768,2311..2317,2243..2249,1358..1364,3122..3128,462..468	1.15	0.29	-0.55	-0.01	0.5 No
NM_001039140	218081_at	C20orf27	Homo sapiens chromosome 20 open reading frame 27 (C20orf27), mRNA.	54976 TR(1..1376)CDS(230..829)	253..259	0.94	0.42	-0.03	-0.06	0.5 Yes
NM_014109	222740_at,235266_at,218782_s_at	ATAD2	Homo sapiens ATPase family, AAA domain containing 2 (ATAD2), mRNA.	29028 TR(1..4916)CDS(91..4263)	3267..3273	1.19	0.26	-0.66	-0.22	0.5 No
NM_003618	218311_at	MAP4K3	Homo sapiens mitogen-activated protein kinase kinase kinase kinase 3 (MAP4K3), mRNA.	8491 TR(1..4141)CDS(92..2776)	1558..1564	0.8	0.48	-0.1	-0.13	0.5 No
NM_007111	242939_at	TFDP1	Homo sapiens transcription factor Dp-1 (TFDP1), mRNA.	7027 TR(1..2385)CDS(213..1445)	725..731	0.92	0.43	0.09	0.05	0.5 No