

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
NM_022823	218843_at	FNDC4	Homo sapiens fibronectin type III domain containing 4 (FNDC4), mRNA.	64838	TR(1..1649)CDS(367..1071)	1523..1530	3.73	1.77	-1.91	-0.39	1
NM_003919	204688_at	SGCE	Homo sapiens sarcoglycan, epsilon (SGCE), transcript variant 2, mRNA.	8910	TR(1..1709)CDS(112..1425)	1495..1501	3.09	1.56	-1.02	-0.27	1
NM_006982	206837_at	ALX1	Homo sapiens ALX homeobox 1 (ALX1), mRNA.	8092	TR(1..1320)CDS(5..985)	916..923	2.99	1.93	-0.19	-0.33	1
NM_019024	233642_s_at	HEATR5B	Homo sapiens HEAT repeat containing 5B (HEATR5B), mRNA.	54497	TR(1..6792)CDS(97..6312)	5827..5834,4309..4315	3.28	1.51	-0.92	-0.23	1
NM_018366	223431_at	CNO	Homo sapiens cappuccino homolog (mouse) (CNO), mRNA.	55330	TR(1..1546)CDS(96..749)	1062..1069,925..932	2.89	1.51	-1.2	-0.41	1
NM_032436	226194_at	C13orf8	Homo sapiens chromosome 13 open reading frame 8 (C13orf8), mRNA.	283489	TR(1..3782)CDS(283..2721)	1756..1762,3587..3594,1725..1731,3395..3402	2.75	1.72	-1.38	-0.34	1
NM_031450	221534_at	C11orf68	Homo sapiens chromosome 11 open reading frame 68 (C11orf68), mRNA.	83638	TR(1..1568)CDS(153..908)	967..973	3.07	1.35	-0.72	-0.06	1
NM_033318	225795_at,225794_s_at	C22orf32	Homo sapiens chromosome 22 open reading frame 32 (C22orf32), mRNA.	91689	TR(1..1581)CDS(75..398)	1170..1176	2.47	1.73	-0.05	-0.02	1
NM_000345	204466_s_at	SNCA	Homo sapiens synuclein, alpha (non A4 component of amyloid precursor) (SNCA), transcript variant NACP140, mRNA.	6622	TR(1..1543)CDS(47..469)	589..596	2.62	1.28	-0.72	-0.23	1
NM_052838	227552_at	SEPT1	Homo sapiens septin 1 (SEPT1), mRNA.	1731	TR(1..1634)CDS(188..1291)	1525..1531	2.36	1.36	-0.02	0	1
NM_002880	201244_s_at	RAF1	Homo sapiens v-raf-1 murine leukemia viral oncogene homolog 1 (RAF1), mRNA.	5894	TR(1..3245)CDS(394..2340)	2992..2999,3014..3021,2202..2209	2.37	1.28	-1.34	-0.43	1
NM_019103	218752_at	ZMAT5	Homo sapiens zinc finger, matrin type 5 (ZMAT5), transcript variant 1, mRNA.	55954	TR(1..1051)CDS(252..764)	657..664	2.3	1.29	-0.04	0.02	1
NM_006468	209382_at,210573_s_at	POLR3C	Homo sapiens polymerase (RNA) III (DNA directed) polypeptide C (62kD) (POLR3C), mRNA.	10623	TR(1..1888)CDS(162..1766)	1025..1032	2.82	1.22	-0.18	-0.09	1
NM_181313	203977_at,37278_at	TAZ	Homo sapiens tafazzin (cardiomyopathy, dilated 3A (X-linked); endocardial fibroelastosis 2; Barth syndrome) (TAZ), transcript variant 4, mRNA.	6901	TR(1..1772)CDS(304..1050)	1493..1499	3.35	1.21	-0.36	0.08	1
NM_032790	226531_at	ORAI1	Homo sapiens ORAI calcium release-activated calcium modulator 1 (ORAI1), mRNA.	84876	TR(1..1497)CDS(174..1079)	1274..1280	2.23	1.48	-1.16	-0.32	1
NM_001014764	221255_s_at	TMEM93	Homo sapiens transmembrane protein 93 (TMEM93), transcript variant 1, mRNA.	83460	TR(1..805)CDS(267..599)	49..55,18..24	2.22	1.6	-0.13	-0.04	1
NM_152293	226240_at	MGC21874	Homo sapiens transcriptional adaptor 2 (ADA2 homolog, yeast)-beta (MGC21874), mRNA.	93624	TR(1..4312)CDS(152..1414)	2690..2697	2.16	1.16	-0.91	-0.02	1
NM_006353	209787_s_at,209786_at	HMGN4	Homo sapiens high mobility group nucleosomal binding domain 4 (HMGN4), mRNA.	10473	TR(1..1980)CDS(239..511)	1882..1889	2.62	1.12	-1.01	-0.27	1
NM_015476	212055_at,213617_s_at	C18orf10	Homo sapiens chromosome 18 open reading frame 10 (C18orf10), mRNA.	25941	TR(1..2079)CDS(429..1331)	2053..2059,746..753	2.43	1.11	-1.03	-0.19	1
NM_015680	207511_s_at,200070_at	C2orf24	Homo sapiens chromosome 2 open reading frame 24 (C2orf24), mRNA.	27013	TR(1..2080)CDS(138..1370)	1903..1910	2.15	1.11	-0.54	-0.13	1
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)	963..969,1702..1708	2.48	1.09	-0.25	-0.1	1
NM_198174	232116_at	GRHL3	Homo sapiens grainyhead-like 3 (Drosophila) (GRHL3), transcript variant 3, mRNA.	57822	TR(1..2528)CDS(154..1683)	2089..2095	1.98	1.74	-0.59	0	1
NM_018356	1552660_a_at	C5orf22	Homo sapiens chromosome 5 open reading frame 22 (C5orf22), mRNA.	55322	TR(1..2719)CDS(115..1443)	1517..1524,2557..2564,1666..1672	2.9	1.08	-1.48	-0.66	1
NM_130386	221019_s_at	COLEC12	Homo sapiens collectin sub-family member 12 (COLEC12), mRNA.	81035	TR(1..3089)CDS(172..2400)	2678..2685	2.23	1.08	-1.4	-0.48	1
NM_024093	219662_at	C2orf49	Homo sapiens chromosome 2 open reading frame 49 (C2orf49), mRNA.	79074	TR(1..902)CDS(33..731)	494..501	2.21	1.08	-0.26	-0.31	1
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)	932..938,2759..2765	2.83	1.04	-0.56	-0.49	1
XM_001128716	203330_s_at	STX5	PREDICTED: Homo sapiens syntaxin 5 (STX5), mRNA.	6811	TR(1..1495)CDS(1..888)	874..880	2.29	1.04	-0.67	-0.16	1
NM_001451	205935_at	FOXF1	Homo sapiens forkhead box F1 (FOXF1), mRNA.	2294	TR(1..2579)CDS(44..1183)	1103..1109	2.2	1.04	-0.04	-0.1	1
NM_002555	204981_at	SLC22A18	Homo sapiens solute carrier family 22 (organic cation transporter), member 18 (SLC22A18), transcript variant 1, mRNA.	5002	TR(1..1549)CDS(205..1479)	1186..1193,1007..1013	2.2	1.03	-0.12	-0.04	1
NM_005165	202022_at	ALDOC	Homo sapiens aldolase C, fructose-bisphosphate (ALDOC), mRNA.	230	TR(1..1665)CDS(146..1240)	1357..1363,1495..1501,381..387	2.23	1.02	-0.7	0.05	1
NM_017632	218929_at	CDKN2AIP	Homo sapiens CDKN2A interacting protein (CDKN2AIP), mRNA.	55602	TR(1..2391)CDS(163..1905)	1080..1087,418..424	2.67	1.02	-0.23	-0.17	1
NM_024587	219462_at,1554077_a_at	TMEM53	Homo sapiens transmembrane protein 53 (TMEM53), mRNA.	79639	TR(1..1614)CDS(37..870)	359..365,617..623	1.96	1.03	0.36	0.08	1
NM_005697	218143_s_at	SCAMP2	Homo sapiens secretory carrier membrane protein 2 (SCAMP2), mRNA.	10066	TR(1..1313)CDS(75..1064)	912..918	1.99	1.02	-0.29	-0.08	1
NM_003009	201194_at,1555851_s_at	SEPW1	Homo sapiens selenoprotein W, 1 (SEPW1), mRNA.	6415	TR(1..893)CDS(202..465)	723..729,649..655	1.88	1.04	-0.52	-0.08	1
NM_007121	218215_s_at	NR1H2	Homo sapiens nuclear receptor subfamily 1, group H, member 2 (NR1H2), mRNA.	7376	TR(1..2087)CDS(277..1662)	1840..1847	2.49	1.01	-1.89	-0.46	1
NM_006223	204572_s_at	PIN4	Homo sapiens protein (peptidylprolyl cis/trans isomerase) NIMA-interacting, 4 (parvulin) (PIN4), mRNA.	5303	TR(1..1088)CDS(36..506)	958..964	2.36	1	0.06	-0.06	1
NM_001031703	229863_s_at,223277_at	TMEM103	Homo sapiens transmembrane protein 103 (TMEM103), mRNA.	54859	TR(1..1374)CDS(120..920)	399..406	2.34	0.99	0.08	0.05	1
NM_152520	1555800_at,1555801_s_at,229019_at	ZNF533	Homo sapiens zinc finger protein 533 (ZNF533), mRNA.	151126	TR(1..3154)CDS(440..1855)	1761..1767	2.9	0.98	-1.95	-0.45	1
NM_019896	1553587_a_at,225844_at	POLE4	Homo sapiens polymerase (DNA-directed), epsilon 4 (p12 subunit) (POLE4), mRNA.	56655	TR(1..710)CDS(33..386)	558..565,578..585	1.72	1.23	-1.26	-0.74	1
NM_004969	217496_s_at	IDE	Homo sapiens insulin-degrading enzyme (IDE), mRNA.	3416	TR(1..3737)CDS(77..3136)	3180..3187	1.71	1.12	-1.06	-0.27	1
NM_003000	202675_at	SDHB	Homo sapiens succinate dehydrogenase complex, subunit B, iron sulfur (lp) (SDHB), mRNA.	6390	TR(1..1161)CDS(152..994)	538..544	2.07	0.95	-0.11	-0.11	1
NM_016326	223451_s_at,221058_s_at,219161_s_at	CKLF	Homo sapiens chemokine-like factor (CKLF), transcript variant 3, mRNA.	51192	TR(1..434)CDS(148..351)	402..408	1.64	1.15	0.32	-0.08	1
NM_153045	221865_at	C9orf91	Homo sapiens chromosome 9 open reading frame 91 (C9orf91), mRNA.	203197	TR(1..4413)CDS(218..1246)	4175..4181	2.51	0.94	-2.15	-0.44	1
NM_206914	213861_s_at	FAM119B	Homo sapiens family with sequence similarity 119, member B (FAM119B), transcript variant 2, mRNA.	25895	TR(1..2856)CDS(126..575)	2175..2181,679..685	1.6	1.11	-1.09	-0.7	1
NM_001031628	209679_s_at	LOC57228	Homo sapiens small trans-membrane and glycosylated protein (LOC57228), transcript variant 1, mRNA.	57228	TR(1..1087)CDS(202..495)	331..337,903..909	1.86	0.92	-0.45	-0.01	1
NM_003761	202546_at	VAMP8	Homo sapiens vesicle-associated membrane protein 8 (endobrevin) (VAMP8), mRNA.	8673	TR(1..702)CDS(54..356)	437..443	1.56	1.08	-0.57	-0.32	1
NM_020979	205367_at	SH2B2	Homo sapiens SH2B adaptor protein 2 (SH2B2), mRNA.	10603	TR(1..2116)CDS(134..2032)	1837..1843	2.16	0.91	-0.4	0.23	1
NM_021734	223222_at	SLC25A19	Homo sapiens solute carrier family 25 (mitochondrial thiamine pyrophosphate carrier), member 19 (SLC25A19), mRNA.	60386	TR(1..1651)CDS(231..1193)	1212..1218,1155..1162	1.85	0.91	-0.29	-0.14	1
NM_024334	222418_s_at,217795_s_at	TMEM43	Homo sapiens transmembrane protein 43 (TMEM43), mRNA.	79188	TR(1..3254)CDS(150..1352)	2626..2632,1531..1537	2.79	0.89	-1.73	-0.31	1
NM_024608	219396_s_at	NEIL1	Homo sapiens nei endonuclease VIII-like 1 (E. coli) (NEIL1), mRNA.	79661	TR(1..1828)CDS(428..1600)	653..659,1751..1757	1.6	0.92	-0.42	-0.17	1
NM_022915	222555_s_at	MRPL44	Homo sapiens mitochondrial ribosomal protein L44 (MRPL44), nuclear gene encoding mitochondrial protein, mRNA.	65080	TR(1..1728)CDS(57..1055)	32..39	1.76	0.87	-0.06	-0.12	1
NM_001532	1553540_a_at,204717_s_at	SLC29A2	Homo sapiens solute carrier family 29 (nucleoside transporters), member 2 (SLC29A2), mRNA.	3177	TR(1..2529)CDS(230..1600)	2217..2223,1085..1092,1148..1154	1.58	0.89	-0.67	-0.03	1

NM_006825	200999_s_at,200998_s_at	CKAP4	Homo sapiens cytoskeleton-associated protein 4 (CKAP4), mRNA.	10970 TR(1..2913)/CDS(85..1893)	2357..2364	2.14	0.86	-1.23	-0.44	1
NM_012425	201980_s_at	RSU1	Homo sapiens Ras suppressor protein 1 (RSU1), transcript variant 1, mRNA.	6251 TR(1..3769)/CDS(144..977)	1391..1397	1.81	0.86	-1.26	-0.39	1
NM_025161	221800_s_at,225322_s_at	C17orf70	Homo sapiens chromosome 17 open reading frame 70 (C17orf70), mRNA.	80233 TR(1..3843)/CDS(696..2888)	2520..2526,173..179	1.7	0.86	0	-0.01	1
NM_019556	1557455_s_at,218853_s_at	MOSPD1	Homo sapiens motile sperm domain containing 1 (MOSPD1), mRNA.	56180 TR(1..2358)/CDS(188..829)	2095..2102,674..680,1616..1622	1.95	0.84	-0.6	-0.52	1
NM_018019	218372_at	MED9	Homo sapiens mediator complex subunit 9 (MED9), mRNA.	55090 TR(1..2222)/CDS(57..497)	1581..1587	1.51	0.88	0.11	-0.23	1
NM_002695	213887_s_at,217854_s_at	POLR2E	Homo sapiens polymerase (RNA) II (DNA directed) polypeptide E, 25kDa (POLR2E), mRNA.	5434 TR(1..1238)/CDS(23..655)	1030..1036	1.95	0.83	-0.35	-0.01	1
NM_032889	212861_at	MFSD5	Homo sapiens major facilitator superfamily domain containing 5 (MFSD5), mRNA.	84975 TR(1..1760)/CDS(192..1544)	990..996	1.92	0.82	-0.56	0.08	1
NM_021240	231800_s_at	DMRT3	Homo sapiens doublesex and mab-3 related transcription factor 3 (DMRT3), mRNA.	58524 TR(1..2184)/CDS(39..1457)	1010..1017	1.8	0.82	-0.44	-0.2	1
NM_032714	224469_s_at	C14orf151	Homo sapiens chromosome 14 open reading frame 151 (C14orf151), mRNA.	84800 TR(1..1704)/CDS(129..833)	1546..1553	2.2	0.81	-0.93	-0.04	1
NM_014319	218604_at	LEM3	Homo sapiens LEM domain containing 3 (LEM3), mRNA.	23592 TR(1..4785)/CDS(7..2742)	2751..2757,2805..2811,4377..4383	1.66	0.82	-0.79	-0.54	1
NM_004059	206037_at	CCBL1	Homo sapiens cysteine conjugate-beta lyase; cytoplasmic (glutamine transaminase K, kynurenine aminotransferase) (CCBL1), mRNA.	883 TR(1..1925)/CDS(186..1454)	1014..1021	2.35	0.81	-0.14	0.28	1
NM_000745	206533_at	CHRNA5	Homo sapiens cholinergic receptor, nicotinic, alpha 5 (CHRNA5), mRNA.	1138 TR(1..2468)/CDS(163..1569)	1042..1048	2.32	0.81	-0.8	-0.04	1
NM_014573	214283_at	TMEM97	Homo sapiens transmembrane protein 97 (TMEM97), mRNA.	27346 TR(1..2585)/CDS(146..676)	1714..1721,841..847	2.13	0.8	-0.53	-0.2	1
NM_014328	206949_s_at	RUSC1	Homo sapiens RUN and SH3 domain containing 1 (RUSC1), transcript variant 4, mRNA.	23623 TR(1..2078)/CDS(212..1513)	1773..1779,1701..1707	2.33	0.8	-1.11	-0.18	1
NM_016525	221490_at,46270_at	UBAP1	Homo sapiens ubiquitin associated protein 1 (UBAP1), mRNA.	51271 TR(1..2757)/CDS(236..1744)	2204..2210,979..986	2.47	0.79	-1.02	-0.2	1
NM_018683	200867_at,200868_s_at,211678_s_at	ZNF313	Homo sapiens zinc finger protein 313 (ZNF313), mRNA.	55905 TR(1..2478)/CDS(37..723)	919..925	2.28	0.78	-0.77	-0.12	1
NM_017996	219641_at	DET1	Homo sapiens de-etiolated homolog 1 (Arabidopsis) (DET1), mRNA.	55070 TR(1..2315)/CDS(162..1847)	330..336,1874..1881,915..922,1150..1156	1.98	0.77	-0.39	-0.29	1
NM_014871	203117_s_at	USP52	Homo sapiens ubiquitin specific peptidase 52 (USP52), mRNA.	9924 TR(1..4509)/CDS(220..3816)	4381..4388	2.13	0.77	-1.02	-0.32	1
NM_152285	226405_s_at	ARRDC1	Homo sapiens arrestin domain containing 1 (ARRDC1), mRNA.	92714 TR(1..1621)/CDS(75..1376)	1068..1074	1.59	0.78	-0.27	0.01	1
NM_024032	219879_s_at	C17orf53	Homo sapiens chromosome 17 open reading frame 53 (C17orf53), mRNA.	78995 TR(1..2727)/CDS(186..2129)	1223..1230	1.15	0.78	-0.01	-0.24	1
NM_001012506	228122_at	CCDC66	Homo sapiens coiled-coil domain containing 66 (CCDC66), mRNA.	285331 TR(1..3098)/CDS(413..2908)	2943..2950	1.13	0.98	-0.04	-0.11	1
NM_003914	205899_at	CCNA1	Homo sapiens cyclin A1 (CCNA1), mRNA.	8900 TR(1..1743)/CDS(130..1527)	229..236	1.7	0.74	0.2	-0.18	1
NM_001003794	225102_at,211026_s_at	MGLL	Homo sapiens monoglyceride lipase (MGLL), transcript variant 2, mRNA.	11343 TR(1..4192)/CDS(146..1057)	989..996,491..497	1.95	0.73	-0.53	-0.02	1
NM_021727	204257_at,216080_s_at	FADS3	Homo sapiens fatty acid desaturase 3 (FADS3), mRNA.	3995 TR(1..1807)/CDS(154..1491)	836..842,871..877,1620..1626	1.59	0.74	-0.19	-0.18	1
NM_017918	218802_at	CCDC109B	Homo sapiens coiled-coil domain containing 109B (CCDC109B), mRNA.	55013 TR(1..1236)/CDS(342..1088)	937..943	1.64	0.73	-0.16	-0.07	1
NM_018413	219634_at	CHST11	Homo sapiens carbohydrate (chondroitin 4) sulfotransferase 11 (CHST11), mRNA.	50515 TR(1..1686)/CDS(296..1354)	1248..1254	1.54	0.74	-0.4	0.04	1
NM_001098	200793_s_at	ACO2	Homo sapiens aconitase 2, mitochondrial (ACO2), nuclear gene encoding mitochondrial protein, mRNA.	50 TR(1..2744)/CDS(23..2365)	2556..2562	1.96	0.72	-0.9	-0.12	1
NM_001010972	200808_s_at	ZYX	Homo sapiens zyxin (ZYX), transcript variant 2, mRNA.	7791 TR(1..2322)/CDS(156..1874)	2099..2106	2.05	0.71	-0.54	0.02	1
NM_001008216	202528_at	GALE	Homo sapiens UDP-galactose-4-epimerase (GALE), transcript variant 2, mRNA.	2582 TR(1..1585)/CDS(170..1216)	1411..1417	1.39	0.74	-0.13	-0.01	1
NM_001217	209726_at	CA11	Homo sapiens carbonic anhydrase XI (CA11), mRNA.	770 TR(1..1686)/CDS(563..1549)	1401..1407	1.7	0.71	-0.02	0.22	1
NM_030919	225687_at	FAM83D	Homo sapiens family with sequence similarity 83, member D (FAM83D), mRNA.	81610 TR(1..2445)/CDS(42..1889)	1915..1921,1559..1565,1545..1551,1574..1580	1.86	0.69	-0.65	-0.21	1
NM_001032291	201896_s_at	PSRC1	Homo sapiens proline/serine-rich coiled-coil 1 (PSRC1), transcript variant 3, mRNA.	84722 TR(1..1738)/CDS(123..1124)	849..855	2.35	0.69	-0.86	-0.11	1
NM_014773	201977_s_at,201978_s_at	KIAA0141	Homo sapiens KIAA0141 (KIAA0141), mRNA.	9812 TR(1..3056)/CDS(99..1646)	814..820,317..323	2.28	0.69	-0.01	-0.24	1
NM_001031690	205368_at	FAM131B	Homo sapiens family with sequence similarity 131, member B (FAM131B), mRNA.	9715 TR(1..4395)/CDS(234..1232)	1602..1609,3955..3962,2058..2064,2995..3002,662..669	1.83	0.68	-0.03	0.01	1
NM_017825	223097_at	ADPRHL2	Homo sapiens ADP-ribosylhydrolase like 2 (ADPRHL2), mRNA.	54936 TR(1..1691)/CDS(21..1112)	1095..1102,635..642	1.95	0.68	-0.38	-0.01	1
NM_005528	223371_s_at,228622_s_at,206782_s_at	DNAJC4	Homo sapiens DnaJ (Hsp40) homolog, subfamily C, member 4 (DNAJC4), mRNA.	3338 TR(1..1355)/CDS(761..1168)	270..276	1.08	0.84	-0.01	-0.14	1
NM_001535	202098_s_at	PRMT2	Homo sapiens protein arginine methyltransferase 2 (PRMT2), transcript variant 2, mRNA.	3275 TR(1..2140)/CDS(206..1507)	2062..2069,1487..1493	1.7	0.67	-0.72	-0.22	1
NM_014969	203855_at	WDR47	Homo sapiens WD repeat domain 47 (WDR47), mRNA.	22911 TR(1..4256)/CDS(262..3024)	1011..1017,3204..3211	1.19	0.77	-0.52	-0.55	1
NM_005865	208165_s_at	PRSS16	Homo sapiens protease, serine, 16 (thymus) (PRSS16), mRNA.	10279 TR(1..2722)/CDS(13..1557)	2447..2453,1493..1499,2555..2561	2.64	0.66	-0.81	-0.03	1
NM_017865	220661_s_at	ZNF692	Homo sapiens zinc finger protein 692 (ZNF692), mRNA.	55657 TR(1..1986)/CDS(168..1727)	69..76,1893..1899	3.25	0.66	-1.07	0.07	1
XM_001128702	238567_at	SGPP2	PREDICTED: Homo sapiens sphingosine-1-phosphate phosphatase 2 (SGPP2), mRNA.	130367 TR(1..1190)/CDS(1..1017)	132..138,704..710	1.56	0.67	-0.02	0.11	1
NM_014285	214507_s_at,209527_at	EXOSC2	Homo sapiens exosome component 2 (EXOSC2), mRNA.	23404 TR(1..2001)/CDS(3..884)	1191..1198	2.07	0.66	-0.67	-0.34	1
NM_013319	219131_at	UBIAD1	Homo sapiens UbiA prenyltransferase domain containing 1 (UBIAD1), mRNA.	29914 TR(1..1520)/CDS(333..1349)	738..744	1.6	0.65	-0.35	-0.14	1
NM_006385	205437_at	ZNF211	Homo sapiens zinc finger protein 211 (ZNF211), transcript variant 1, mRNA.	10520 TR(1..2469)/CDS(132..1865)	2224..2230	1.96	0.64	0.05	-0.01	1
NM_138411	229289_at	LOC112703	Homo sapiens hypothetical protein BC004941 (LOC112703), mRNA.	112703 TR(1..1329)/CDS(359..1054)	1115..1122	1.39	0.68	-0.04	0.21	1
NM_145019	1553239_at	FAM124A	Homo sapiens family with sequence similarity 124A (FAM124A), mRNA.	220108 TR(1..2104)/CDS(132..1880)	1697..1704	0.83	1.03	0.06	-0.01	1
NM_018226	218301_at	RNPEPL1	Homo sapiens arginyl aminopeptidase (aminopeptidase B)-like 1 (RNPEPL1), mRNA.	57140 TR(1..2948)/CDS(594..2078)	849..855,633..640	0.99	0.81	-0.05	-0.53	1
NM_005184	200623_s_at	CALM3	Homo sapiens calmodulin 3 (phosphorylase kinase, delta) (CALM3), mRNA.	808 TR(1..2277)/CDS(181..630)	857..863,869..876	1.72	0.64	-0.96	-0.05	1
NM_080760	205471_s_at,228915_at,205472_s_at	DACH1	Homo sapiens dachshund homolog 1 (Drosophila) (DACH1), transcript variant 2, mRNA.	1602 TR(1..4796)/CDS(424..2100)	3737..3743,1668..1675,3927..3933	1.53	0.65	-1.02	-0.95	1
NM_170696	207015_s_at,207016_s_at	ALDH1A2	Homo sapiens aldehyde dehydrogenase 1 family, member A2 (ALDH1A2), transcript variant 2, mRNA.	8854 TR(1..3284)/CDS(59..1501)	1918..1924,2810..2816,2822..2828	1.44	0.66	-1.34	-0.17	1
NM_015329	216926_s_at,227465_at,212505_s_at	KIAA0892	Homo sapiens KIAA0892 (KIAA0892), mRNA.	23383 TR(1..4853)/CDS(40..1881)	875..881,3490..3496,1448..1454,482..488	2.05	0.62	-0.45	-0.09	0.99
NM_006646	204042_at	WASF3	Homo sapiens WAS protein family, member 3 (WASF3), mRNA.	10810 TR(1..4851)/CDS(227..1735)	3913..3919,2131..2138,910..916	1.84	0.62	-1.81	-0.54	0.99
NM_014169	218572_at,218571_s_at	CHMP4A	Homo sapiens chromatin modifying protein 4A (CHMP4A), mRNA.	29082 TR(1..1505)/CDS(412..1209)	1223..1229	1.46	0.64	0.04	-0.09	0.99

NM_139212	211597_s_at	HOPX	Homo sapiens HOP homeobox (HOPX), transcript variant 3, mRNA.	84525 TR(1..1116)CDS(158..379)	138..144	1.02	0.77	0.02	0	0.99
NM_002149	205462_s_at,212552_at	HPCAL1	Homo sapiens hippocalcin-like 1 (HPCAL1), transcript variant 1, mRNA.	3241 TR(1..1769)CDS(375..956)	576..582	2.01	0.6	-0.15	0.08	0.99
NM_002408	211061_s_at,203102_s_at,203101_s_at	MGAT2	Homo sapiens mannosyl (alpha-1,6-)-glycoprotein beta-1,2-N-acetylglucosaminyltransferase (MGAT2), mRNA.	4247 TR(1..2728)CDS(499..1842)	1568..1575	1.89	0.6	-0.44	-0.09	0.99
NM_002576	1565772_at,209615_s_at,226507_at	PAK1	Homo sapiens p21/Cdc42/Rac1-activated kinase 1 (STE20 homolog, yeast) (PAK1), mRNA.	5058 TR(1..3264)CDS(359..1996)	2281..2287	1.73	0.59	-0.67	-0.21	0.99
NM_001029859	229873_at	KCTD21	Homo sapiens potassium channel tetramerisation domain containing 21 (KCTD21), mRNA.	283219 TR(1..3396)CDS(76..858)	487..493	0.64	1.08	0.02	-0.33	0.99
NM_145806	225307_at	ZNF511	Homo sapiens zinc finger protein 511 (ZNF511), mRNA.	118472 TR(1..1098)CDS(26..784)	423..429	1.37	0.63	-0.1	0	0.99
XM_379194	1557754_at	LOC401068	PREDICTED: Homo sapiens hypothetical LOC401068 (LOC401068), mRNA.	401068 TR(1..2060)CDS(1..432)	659..665	1.29	0.65	0.12	0.17	0.99
NM_015915	223340_at	SPG3A	Homo sapiens spastic paraplegia 3A (autosomal dominant) (SPG3A), transcript variant 1, mRNA.	51062 TR(1..2271)CDS(235..1911)	1556..1562	1.44	0.61	-0.61	-0.02	0.99
NM_001039569	1555731_a_at,1555733_s_at	AP1S3	Homo sapiens adaptor-related protein complex 1, sigma 3 subunit (AP1S3), mRNA.	130340 TR(1..4005)CDS(153..617)	3864..3870	1.22	0.67	-0.66	-0.08	0.99
NM_015062	203737_s_at	PPRC1	Homo sapiens peroxisome proliferator-activated receptor gamma, coactivator-related 1 (PPRC1), mRNA.	23082 TR(1..5354)CDS(40..5034)	642..648,2741..2748,4509..4515,4380..4386	1.41	0.62	-0.81	-0.24	0.99
NM_181762	201898_s_at,201899_s_at	UBE2A	Homo sapiens ubiquitin-conjugating enzyme E2A (RAD6 homolog) (UBE2A), transcript variant 2, mRNA.	7319 TR(1..1709)CDS(177..545)	392..399	1.3	0.65	-0.39	-0.12	0.99
NM_014413	217736_s_at	EIF2AK1	Homo sapiens eukaryotic translation initiation factor 2-alpha kinase 1 (EIF2AK1), mRNA.	27102 TR(1..2813)CDS(63..1955)	2235..2241,2349..2355	1.61	0.59	-0.98	-0.06	0.99
NM_001099432	220488_s_at	BCAS3	Homo sapiens breast carcinoma amplified sequence 3 (BCAS3), transcript variant 1, mRNA.	54828 TR(1..3626)CDS(110..2896)	1241..1248	1.42	0.61	-0.15	-0.01	0.99
NM_000154	204374_s_at	GALK1	Homo sapiens galactokinase 1 (GALK1), mRNA.	2584 TR(1..1361)CDS(64..1242)	1316..1322	1.16	0.68	0.05	0.29	0.99
NM_017653	223171_at	DYM	Homo sapiens dymeclin (DYM), mRNA.	54808 TR(1..2644)CDS(366..2375)	985..992,925..931,603..609	2.27	0.57	-0.81	-0.08	0.99
NM_016166	217864_s_at	PIAS1	Homo sapiens protein inhibitor of activated STAT, 1 (PIAS1), mRNA.	8554 TR(1..2309)CDS(97..2052)	1586..1593,1945..1951	1.79	0.57	-0.54	-0.15	0.99
NM_006395	218673_s_at,224025_s_at	ATG7	Homo sapiens ATG7 autophagy related 7 homolog (S. cerevisiae) (ATG7), mRNA.	10533 TR(1..2396)CDS(26..2137)	681..687,486..493,1436..1442,1973..1979	2.13	0.57	-0.41	-0.02	0.99
NM_145265	226515_at	CCDC127	Homo sapiens coiled-coil domain containing 127 (CCDC127), mRNA.	133957 TR(1..1437)CDS(101..883)	954..960	1.51	0.59	-0.29	-0.15	0.99
NM_012267	202415_s_at	HSPBP1	Homo sapiens hsp70-interacting protein (HSPBP1), mRNA.	23640 TR(1..1559)CDS(166..1245)	1473..1479	1.55	0.58	0	0.02	0.99
NM_013339	219649_at	ALG6	Homo sapiens asparagine-linked glycosylation 6 homolog (S. cerevisiae, alpha-1,3-glucosyltransferase) (ALG6), mRNA.	29929 TR(1..2003)CDS(269..1792)	1552..1558,1229..1235	1.7	0.57	-0.13	0.04	0.99
NM_194255	209776_s_at,211576_s_at,209777_s_at	SLC19A1	Homo sapiens solute carrier family 19 (folate transporter), member 1 (SLC19A1), mRNA.	6573 TR(1..2839)CDS(120..1895)	1167..1173,1858..1865	1.67	0.57	-0.48	0.04	0.99
NM_017819	223267_at	RG9MTD1	Homo sapiens RNA (guanine-9-)-methyltransferase domain containing 1 (RG9MTD1), mRNA.	54931 TR(1..1631)CDS(149..1360)	268..275	2.07	0.56	0.25	0.01	0.99
NM_024075	218132_s_at	TSEN34	Homo sapiens tRNA splicing endonuclease 34 homolog (S. cerevisiae) (TSEN34), transcript variant 1, mRNA.	79042 TR(1..1368)CDS(110..1042)	604..610	1.7	0.56	-0.4	-0.07	0.99
NM_181900	213820_s_at	STARD5	Homo sapiens STAR-related lipid transfer (START) domain containing 5 (STARD5), mRNA.	80765 TR(1..1344)CDS(85..726)	570..576,608..614	3.48	0.55	-0.22	0.04	0.99
NM_178148	224716_at	SLC35B2	Homo sapiens solute carrier family 35, member B2 (SLC35B2), mRNA.	347734 TR(1..2057)CDS(126..1424)	1609..1615	1.8	0.55	0.03	0.08	0.99
NM_001839	201445_at	CNN3	Homo sapiens calponin 3, acidic (CNN3), mRNA.	1266 TR(1..2045)CDS(252..1241)	1323..1329	2.25	0.55	-0.97	-0.17	0.99
NM_004900	206632_s_at	APOBEC3B	Homo sapiens apolipoprotein B mRNA editing enzyme, catalytic polypeptide-like 3B (APOBEC3B), mRNA.	9582 TR(1..1536)CDS(55..1203)	1510..1516,1266..1272	0.78	0.8	-0.36	-0.64	0.99
NM_032808	227933_at	LINGO1	Homo sapiens leucine rich repeat and Ig domain containing 1 (LINGO1), mRNA.	84894 TR(1..2932)CDS(53..1915)	1478..1484,2345..2351,1112..1119	1.51	0.57	-0.51	0.01	0.99
NM_021179	206721_at	C1orf114	Homo sapiens chromosome 1 open reading frame 114 (C1orf114), mRNA.	57821 TR(1..1866)CDS(169..1695)	943..949	0.95	0.74	-0.18	-0.26	0.98
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)	2648..2654	1.51	0.57	-0.7	0.03	0.98
NM_175039	220937_s_at,223285_s_at	ST6GALNAC4	Homo sapiens ST6 (alpha-N-acetylneuraminy-2,3-beta-galactosyl-1, 3)-N-acetylgalactosaminide alpha-2,6-sialyltransferase 4 (ST6GALNAC4), transcript variant 1, mRNA.	27090 TR(1..1717)CDS(277..1185)	860..866	1.64	0.55	0.1	-0.02	0.98
NM_001733	212067_s_at	C1R	Homo sapiens complement component 1, r subcomponent (C1R), mRNA.	715 TR(1..2526)CDS(60..2177)	2422..2428,234..241	0.99	0.72	-0.03	-0.53	0.98
NM_015472	202134_s_at	WWTR1	Homo sapiens WW domain containing transcription regulator 1 (WWTR1), mRNA.	25937 TR(1..2015)CDS(235..1437)	1322..1329	1.97	0.54	-0.47	-0.01	0.98
NM_014388	220251_at,214193_s_at,204699_s_at	C1orf107	Homo sapiens chromosome 1 open reading frame 107 (C1orf107), mRNA.	27042 TR(1..8463)CDS(77..2347)	6080..6086,2681..2687,1808..1815	1.51	0.55	-0.75	-0.15	0.98
NM_003877	203372_s_at	SOCS2	Homo sapiens suppressor of cytokine signaling 2 (SOCS2), mRNA.	8835 TR(1..2210)CDS(591..1187)	2111..2117	1.41	0.56	-0.58	-0.35	0.98
NM_018379	221856_s_at	FAM63A	Homo sapiens family with sequence similarity 63, member A (FAM63A), transcript variant 1, mRNA.	55793 TR(1..4043)CDS(838..2247)	522..528,2279..2285	0.37	1.1	0.43	-0.27	0.98
NM_032989	1861_at,209364_at	BAD	Homo sapiens BCL2-antagonist of cell death (BAD), transcript variant 2, mRNA.	572 TR(1..956)CDS(71..577)	518..525	1.33	0.58	-0.09	0.11	0.98
NM_001571	202621_at	IRF3	Homo sapiens interferon regulatory factor 3 (IRF3), mRNA.	3661 TR(1..1606)CDS(246..1529)	403..409,888..895	1.39	0.57	0.32	0.05	0.98
NM_032718	224483_s_at	MFSD9	Homo sapiens major facilitator superfamily domain containing 9 (MFSD9), mRNA.	84804 TR(1..2682)CDS(45..1469)	371..377	1.01	0.69	-0.11	-0.01	0.98
NM_006371	226656_at,227138_at,201380_at,1554464_a_at,1555889_a_at	CRTAP	Homo sapiens cartilage associated protein (CRTAP), mRNA.	10491 TR(1..6622)CDS(92..1297)	4905..4912,1884..1890,1863..1869	2.15	0.52	-1.3	-0.14	0.98
NM_194318	227100_at,227083_at	B3GALTL	Homo sapiens beta 1,3-galactosyltransferase-like (B3GALTL), mRNA.	145173 TR(1..4221)CDS(111..1607)	4149..4156,1429..1435	1.67	0.52	-0.83	-0.3	0.98
NM_015168	213390_at	C19orf7	Homo sapiens chromosome 19 open reading frame 7 (C19orf7), mRNA.	23211 TR(1..6131)CDS(39..3950)	6030..6037	0.99	0.69	-0.45	-0.23	0.98
NM_033276	227678_at	XRCC6BP1	Homo sapiens XRCC6 binding protein 1 (XRCC6BP1), mRNA.	91419 TR(1..1182)CDS(41..781)	1095..1101,119..125	1.51	0.53	-0.79	-0.09	0.97
NM_024789	219745_at	TMEM180	Homo sapiens transmembrane protein 180 (TMEM180), mRNA.	79847 TR(1..2834)CDS(220..1773)	1048..1055,1520..1526,313..319	1.5	0.53	-0.46	-0.04	0.97
NM_005506	201647_s_at,224983_at,201646_at	SCARB2	Homo sapiens scavenger receptor class B, member 2 (SCARB2), mRNA.	950 TR(1..4748)CDS(340..1776)	2338..2344,1141..1147,2355..2362,421..428	2.11	0.5	-1.05	-0.21	0.97
NM_021931	218579_s_at,234728_s_at	DHX35	Homo sapiens DEAH (Asp-Glu-Ala-His) box polypeptide 35 (DHX35), mRNA.	60625 TR(1..3316)CDS(12..2123)	3289..3295,3012..3018	2.68	0.49	-0.23	0.09	0.97
NM_001755	206788_s_at,202370_s_at	CBFB	Homo sapiens core-binding factor, beta subunit (CBFB), transcript variant 2, mRNA.	865 TR(1..3181)CDS(262..810)	1153..1159,2014..2020	1.64	0.5	-0.17	0	0.97
NM_001001655	225625_at	ALKBH2	Homo sapiens alkB, alkylation repair homolog 2 (E. coli) (ALKBH2), mRNA.	121642 TR(1..1051)CDS(196..981)	905..911,781..787	1.49	0.52	-0.03	0.02	0.97
NM_001009184	212090_at	GRINA	Homo sapiens glutamate receptor, ionotropic, N-methyl D-aspartate-associated protein 1 (glutamate binding) (GRINA), transcript variant 2, mRNA.	2907 TR(1..1874)CDS(167..1282)	14..20,1640..1646,521..527	1.57	0.5	-0.26	0.09	0.97
NM_016019	217842_at,220099_s_at	LUC7L2	Homo sapiens LUC7-like 2 (S. cerevisiae) (LUC7L2), mRNA.	51631 TR(1..2725)CDS(375..1553)	175..182,1881..1887	2.1	0.49	-0.93	-0.32	0.97
NM_006855	204017_at,207265_s_at,207264_at	KDEL3	Homo sapiens KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 3 (KDEL3), transcript variant 1, mRNA.	11015 TR(1..1705)CDS(157..801)	782..789	1.27	0.57	-0.35	0.01	0.97
NM_016422	226106_at	RNF141	Homo sapiens ring finger protein 141 (RNF141), mRNA.	50862 TR(1..4084)CDS(154..846)	1013..1020	1.31	0.56	-0.84	-0.4	0.97
NM_144580	1558693_s_at,225401_at	C1orf85	Homo sapiens chromosome 1 open reading frame 85 (C1orf85), mRNA.	112770 TR(1..1603)CDS(14..1234)	803..810	0.73	0.75	-0.16	-0.14	0.97

NM_001012300	202556_s_at	MCRS1	Homo sapiens microspherule protein 1 (MCRS1), transcript variant 2, mRNA.	10445 TR(1..1785)CDS(1..1428)	1360..1366	1.4	0.53	-0.28	0.05	0.97
NM_014746	204040_at	RNF144A	Homo sapiens ring finger protein 144A (RNF144A), mRNA.	9781 TR(1..5740)CDS(443..1321)	5317..5324	1.06	0.64	-1.05	-0.42	0.97
NM_001008222	222451_s_at	ZDHHHC9	Homo sapiens zinc finger, DHHC-type containing 9 (ZDHHHC9), transcript variant 2, mRNA.	51114 TR(1..2913)CDS(370..1464)	2643..2649,2517..2523,159..165,226..233	1.74	0.48	-1.08	-0.11	0.96
NM_138290	215321_at	RUNDC3B	Homo sapiens RUN domain containing 3B (RUNDC3B), mRNA.	154661 TR(1..2241)CDS(410..1831)	263..270	1.79	0.48	-1.22	-0.19	0.96
NM_032344	224464_s_at	NUDT22	Homo sapiens nudix (nucleoside diphosphate linked moiety X)-type motif 22 (NUDT22), mRNA.	84304 TR(1..1125)CDS(157..1068)	559..565	2.08	0.47	-0.13	0.24	0.96
NM_006062	209516_at	SMYD5	Homo sapiens SMYD family member 5 (SMYD5), mRNA.	10322 TR(1..2585)CDS(30..1286)	2217..2224	1.33	0.54	-0.78	-0.22	0.96
NM_005567	200923_at	LGALS3BP	Homo sapiens lectin, galactoside-binding, soluble, 3 binding protein (LGALS3BP), mRNA.	3959 TR(1..2254)CDS(180..1937)	1588..1594	1.77	0.47	-0.31	0.05	0.96
NM_005573	203276_at	LMNB1	Homo sapiens lamin B1 (LMNB1), mRNA.	4001 TR(1..2882)CDS(369..2129)	2141..2147	0.84	0.71	-1.01	-1.05	0.96
NM_004899	205550_s_at	BRE	Homo sapiens brain and reproductive organ-expressed (TNFRSF1A modulator) (BRE), transcript variant 1, mRNA.	9577 TR(1..1893)CDS(178..1425)	1819..1825	1.29	0.55	-0.02	0.01	0.96
NM_006509	205205_at	RELB	Homo sapiens v-rel reticuloendotheliosis viral oncogene homolog B, nuclear factor of kappa light polypeptide gene enhancer in B-cells 3 (avian) (RELB), mRNA.	5971 TR(1..2287)CDS(127..1866)	2239..2245,1261..1267	2.1	0.46	-0.34	-0.01	0.96
NM_000263	204360_s_at	NAGLU	Homo sapiens N-acetylglucosaminidase, alpha- (Sanfilippo disease IIIB) (NAGLU), mRNA.	4669 TR(1..2798)CDS(341..2572)	2441..2448,1398..1404	1.34	0.53	-0.14	-0.01	0.96
NM_001035513	238056_at	SDHC	Homo sapiens succinate dehydrogenase complex, subunit C, integral membrane protein, 15kDa (SDHC), nuclear gene encoding mitochondrial protein, transcript variant 4, mRNA.	6391 TR(1..2699)CDS(31..381)	2248..2255,232..238	1.04	0.64	-0.72	-0.58	0.96
NM_016016	223649_s_at	SLC25A39	Homo sapiens solute carrier family 25, member 39 (SLC25A39), mRNA.	51629 TR(1..1556)CDS(118..1173)	1133..1139,1414..1420	1.46	0.49	-0.31	0.02	0.96
XM_939866	228953_at	WHDC1	PREDICTED: Homo sapiens WAS protein homology region 2 domain containing 1, transcript variant 6 (WHDC1), mRNA.	123720 TR(1..4261)CDS(507..2936)	4190..4196,1619..1625,1219..1225	0.94	0.67	-0.5	-0.11	0.96
NM_018728	218966_at	MYO5C	Homo sapiens myosin VC (MYO5C), mRNA.	55930 TR(1..6946)CDS(138..5366)	5788..5795,5639..5645	1.45	0.49	-0.08	-0.08	0.96
NM_130459	227972_at	TOR2A	Homo sapiens torsin family 2, member A (TOR2A), transcript variant 2, mRNA.	27433 TR(1..2509)CDS(56..817)	384..390	1.34	0.52	0	0.03	0.96
NM_024610	219284_at	HSPBAP1	Homo sapiens HSPB (heat shock 27kDa) associated protein 1 (HSPBAP1), mRNA.	79663 TR(1..1958)CDS(124..1590)	1868..1874	1.18	0.57	-0.58	-0.07	0.95
NM_020832	234924_s_at	ZNF687	Homo sapiens zinc finger protein 687 (ZNF687), mRNA.	57592 TR(1..4598)CDS(99..3812)	986..992	1.86	0.44	-0.5	0.28	0.95
NM_018704	231822_at	CTTNBP2NL	Homo sapiens CTTNBP2 N-terminal like (CTTNBP2NL), mRNA.	55917 TR(1..4318)CDS(169..2088)	2026..2032,1387..1393	1.6	0.45	-1.23	-0.45	0.95
NM_005729	201489_at,201490_s_at	PP1F	Homo sapiens peptidylprolyl isomerase F (cyclophilin F) (PP1F), nuclear gene encoding mitochondrial protein, mRNA.	10105 TR(1..2213)CDS(86..709)	1103..1109	1.48	0.46	-0.62	-0.13	0.94
NM_153611	224735_at	CYBASC3	Homo sapiens cytochrome b, ascorbate dependent 3 (CYBASC3), mRNA.	220002 TR(1..2994)CDS(637..1365)	1469..1475,1009..1015,1048..1054,2113..2119	1.78	0.43	-0.52	0.15	0.94
NM_002208	205055_at	ITGAE	Homo sapiens integrin, alpha E (antigen CD103, human mucosal lymphocyte antigen 1; alpha polypeptide) (ITGAE), mRNA.	3682 TR(1..3878)CDS(100..3639)	3442..3448,3068..3074,2540..2546,107..113	1.34	0.49	-0.01	-0.01	0.94
NM_001080493	229732_at	HSZFP36	Homo sapiens ZFP-36 for a zinc finger protein (HSZFP36), mRNA.	55552 TR(1..2414)CDS(127..1959)	2190..2196	1.02	0.61	-0.35	-0.03	0.94
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)	3886..3893,1731..1737	0.94	0.64	-0.44	-0.49	0.93
NM_006084	203882_at	IRF9	Homo sapiens interferon regulatory factor 9 (IRF9), mRNA.	10379 TR(1..1699)CDS(128..1309)	1325..1331	0.06	0.82	0.06	-0.01	0.93
NM_032806	225689_at	C3orf39	Homo sapiens chromosome 3 open reading frame 39 (C3orf39), mRNA.	84892 TR(1..2556)CDS(344..2086)	627..633	1.36	0.47	-0.06	0.11	0.93
NM_003730	217984_at	RNASET2	Homo sapiens ribonuclease T2 (RNASET2), mRNA.	8635 TR(1..1250)CDS(408..1178)	334..340	1.22	0.53	-0.05	-0.17	0.93
NM_021971	219920_s_at	GMPPB	Homo sapiens GDP-mannose pyrophosphorylase B (GMPPB), transcript variant 2, mRNA.	29925 TR(1..1583)CDS(226..1308)	814..820	0.51	0.75	0.09	-0.24	0.93
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)	2037..2043	1.73	0.41	-0.32	-0.08	0.93
NM_001360	201790_s_at,201791_s_at	DHCR7	Homo sapiens 7-dehydrocholesterol reductase (DHCR7), mRNA.	1717 TR(1..2665)CDS(274..1701)	2057..2063	1.31	0.48	-0.45	0.07	0.93
NM_000280	235795_at,205646_s_at	PAX6	Homo sapiens paired box 6 (PAX6), transcript variant 1, mRNA.	5080 TR(1..2816)CDS(513..1781)	2462..2468	1.94	0.4	-0.59	-0.03	0.93
NM_005388	204449_at,204448_s_at	PDCL	Homo sapiens phosducin-like (PDCL), mRNA.	5082 TR(1..2774)CDS(97..1002)	1553..1559	1.21	0.52	-0.26	-0.06	0.92
NM_001659	211622_s_at,200734_s_at,200011_s_at	ARF3	Homo sapiens ADP-ribosylation factor 3 (ARF3), mRNA.	377 TR(1..3595)CDS(312..857)	225..232	1.22	0.5	0.04	0.17	0.92
NM_181829	218915_at,238618_at,204991_s_at,211092_s_at,217150_s_at,211017_s_at,211091_s_at,210767_at	NF2	Homo sapiens neurofibromin 2 (bilateral acoustic neuroma) (NF2), transcript variant 6, mRNA.	4771 TR(1..5943)CDS(424..2073)	2380..2386,2911..2917,4052..4058	1.25	0.51	-0.72	-0.14	0.91
NM_024704	219570_at	C20orf23	Homo sapiens chromosome 20 open reading frame 23 (C20orf23), mRNA.	55614 TR(1..5274)CDS(159..4112)	4723..4730,3450..3456,4552..4558	1.64	0.39	-0.7	-0.28	0.91
NM_020895	224807_at	GRAMD1A	Homo sapiens GRAM domain containing 1A (GRAMD1A), mRNA.	57655 TR(1..2715)CDS(148..2322)	1270..1277	1.21	0.51	-0.29	0.04	0.91
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)	1471..1477,1507..1513	1.3	0.47	0.04	0.09	0.91
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)	892..899,280..286,55..61	1.89	0.37	-0.52	-0.05	0.91
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)	1297..1304,180..187	1.45	0.41	-0.27	0.01	0.9
NM_153000	225016_at	APCDD1	Homo sapiens adenomatosis polyposis coli down-regulated 1 (APCDD1), mRNA.	147495 TR(1..2572)CDS(354..1898)	2304..2310,1294..1300	1.4	0.43	-1.27	-0.05	0.9
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)	2129..2136	1.08	0.55	-0.6	-0.35	0.9
NM_014876	201751_at	JOSD1	Homo sapiens Josephin domain containing 1 (JOSD1), mRNA.	9929 TR(1..3205)CDS(300..908)	429..436	1.17	0.51	-0.34	-0.13	0.9
NM_198563	213851_at	TMEM110	Homo sapiens transmembrane protein 110 (TMEM110), mRNA.	375346 TR(1..1614)CDS(82..966)	1569..1575	1.39	0.42	-0.08	-0.01	0.9
NM_012160	209943_at	FBXL4	Homo sapiens F-box and leucine-rich repeat protein 4 (FBXL4), mRNA.	26235 TR(1..2777)CDS(359..2224)	23..29	1.2	0.5	0.02	-0.06	0.9
NM_003239	209747_at	TGFB3	Homo sapiens transforming growth factor, beta 3 (TGFB3), mRNA.	7043 TR(1..2574)CDS(254..1492)	762..768,2317..2323	0.06	0.78	-0.03	-0.01	0.89
NM_173039	229526_at	AQP11	Homo sapiens aquaporin 11 (AQP11), mRNA.	282679 TR(1..1561)CDS(359..1174)	885..891	0.09	0.78	0.26	0.01	0.89
NM_058237	220673_s_at,220672_at,233002_at	KIAA1622	Homo sapiens KIAA1622 (KIAA1622), transcript variant 1, mRNA.	57718 TR(1..3868)CDS(155..2776)	1522..1528	1.66	0.35	-0.03	-0.01	0.89
NM_002435	202472_at	MPI	Homo sapiens mannose phosphate isomerase (MPI), mRNA.	4351 TR(1..1771)CDS(6..1277)	1505..1511	1.34	0.43	-0.2	-0.04	0.89
NM_199133	225670_at	LOC134145	Homo sapiens hypothetical protein LOC134145 (LOC134145), mRNA.	134145 TR(1..1865)CDS(30..731)	765..771	1.17	0.5	-0.89	-0.42	0.88
NM_004779	202163_s_at,202164_s_at,202162_s_at	CNOT8	Homo sapiens CCR4-NOT transcription complex, subunit 8 (CNOT8), mRNA.	9337 TR(1..2550)CDS(252..1130)	1339..1345,1614..1620	1.51	0.37	-0.27	-0.27	0.88
NM_018727	219632_s_at	TRPV1	Homo sapiens transient receptor potential cation channel, subfamily V, member 1 (TRPV1), transcript variant 2, mRNA.	7442 TR(1..4074)CDS(159..2678)	3947..3953	0.22	0.76	0.26	-0.07	0.88
NM_173491	241599_at,242112_at	LSM11	Homo sapiens LSM11, U7 small nuclear RNA associated (LSM11), mRNA.	134353 TR(1..2561)CDS(5..1087)	1367..1373,1744..1750	1.2	0.48	-1.33	-0.14	0.88
NM_001018108	226692_at	SERF2	Homo sapiens small EDRK-rich factor 2 (SERF2), mRNA.	10169 TR(1..2611)CDS(99..278)	1363..1369,998..1004	1.42	0.39	-0.8	-0.01	0.88

NM_000676	205891_at	ADORA2B	Homo sapiens adenosine A2b receptor (ADORA2B), mRNA.	136 TR(1..1885)CDS(333..1331)	1346..1352	0.82	0.55	-0.06	-0.19	0.71
NM_024602	218632_at	HECTD3	Homo sapiens HECT domain containing 3 (HECTD3), mRNA.	79654 TR(1..3411)CDS(724..2457)	2555..2562,2935..2941	0.9	0.52	-0.42	-0.23	0.71
NM_152341	212858_at	PAQR4	Homo sapiens progesterin and adipoQ receptor family member IV (PAQR4), mRNA.	124222 TR(1..2701)CDS(335..1156)	1733..1739,1313..1319,1556..1562	1.2	0.37	-0.77	-0.08	0.7
NM_001348	203891_s_at,203890_s_at	DAPK3	Homo sapiens death-associated protein kinase 3 (DAPK3), mRNA.	1613 TR(1..2105)CDS(94..1458)	1952..1959	0.82	0.55	-0.06	-0.11	0.7
NM_025187	223440_at	C16orf70	Homo sapiens chromosome 16 open reading frame 70 (C16orf70), mRNA.	80262 TR(1..2943)CDS(165..1433)	2651..2657,1274..1280,541..547	0.8	0.55	-0.55	-0.28	0.69
NM_033647	1552788_a_at,1552787_at	HELB	Homo sapiens helicase (DNA) B (HELB), mRNA.	92797 TR(1..3368)CDS(29..3292)	3211..3218,1079..1085,615..621	1.97	0.13	-1.72	-0.97	0.69
NM_025000	231921_at,220172_at	C2orf37	Homo sapiens chromosome 2 open reading frame 37 (C2orf37), mRNA.	80067 TR(1..5744)CDS(230..1792)	4292..4298,3903..3909	1.54	0.18	-1.2	-0.04	0.69
NM_016462	223106_at	TMEM14C	Homo sapiens transmembrane protein 14C (TMEM14C), mRNA.	51522 TR(1..1011)CDS(182..520)	568..574	0.95	0.49	-0.13	-0.03	0.69
NM_020147	219596_at	THAP10	Homo sapiens THAP domain containing 10 (THAP10), mRNA.	56906 TR(1..2100)CDS(143..916)	1971..1977,1843..1849	1.09	0.42	-0.31	-0.13	0.68
NM_005789	209853_s_at,200988_s_at	PSME3	Homo sapiens proteasome (prosome, macropain) activator subunit 3 (PA28 gamma; Ki) (PSME3), transcript variant 1, mRNA.	10197 TR(1..3189)CDS(227..991)	2322..2329,3159..3165,2404..2410,2169..2175	1.26	0.32	-0.13	-0.24	0.68
NM_018444	218273_s_at,222572_at	PPM2C	Homo sapiens protein phosphatase 2C, magnesium-dependent, catalytic subunit (PPM2C), nuclear gene encoding mitochondrial protein, mRNA.	54704 TR(1..4215)CDS(191..1804)	1704..1710	0.4	0.65	-0.27	-0.57	0.67
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)	1426..1433	1.13	0.4	-0.35	-0.32	0.67
NM_001013842	227263_at	C8orf58	Homo sapiens chromosome 8 open reading frame 58 (C8orf58), mRNA.	541565 TR(1..2075)CDS(113..1210)	1260..1266	0.76	0.56	-0.05	-0.01	0.67
NM_194285	235440_at	SPTY2D1	Homo sapiens SPT2, Suppressor of Ty, domain containing 1 (S. cerevisiae) (SPTY2D1), mRNA.	144108 TR(1..5752)CDS(217..2274)	970..976,1843..1849,4466..4473	1.05	0.44	-0.65	-0.19	0.67
NM_018082	219459_at	POLR3B	Homo sapiens polymerase (RNA) III (DNA directed) polypeptide B (POLR3B), mRNA.	55703 TR(1..4237)CDS(170..3571)	3507..3513,3472..3478	1.35	0.26	-0.49	-0.02	0.67
NM_001099225	225607_at	CCDC43	Homo sapiens coiled-coil domain containing 43 (CCDC43), transcript variant 2, mRNA.	124808 TR(1..2088)CDS(45..509)	946..952	1.41	0.22	-0.88	-0.18	0.67
NM_017575	214940_s_at	SMG6	Homo sapiens Smg-6 homolog, nonsense mediated mRNA decay factor (C. elegans) (SMG6), mRNA.	23293 TR(1..5968)CDS(56..4315)	4437..4443,3091..3097,2644..2650	0.19	0.67	-0.04	-0.01	0.67
XM_001132511	221011_s_at	LBH	PREDICTED: Homo sapiens hypothetical protein DKFZp566J091 (LBH), mRNA.	81606 TR(1..3682)CDS(1004..1270)	3016..3022,3193..3199,2195..2202	0.79	0.55	-0.75	-0.34	0.67
NM_144770	232549_at	RBM11	Homo sapiens RNA binding motif protein 11 (RBM11), mRNA.	54033 TR(1..1952)CDS(382..885)	675..681,549..555	2.84	0.08	-0.78	0	0.67
NM_018999	209378_s_at,209379_s_at,1554132_a_at	KIAA1128	Homo sapiens KIAA1128 (KIAA1128), mRNA.	54462 TR(1..7676)CDS(195..2699)	3398..3404	1.19	0.36	-0.3	-0.41	0.67
NM_005540	213643_s_at,213804_at	INPP5B	Homo sapiens inositol polyphosphate-5-phosphatase, 75kDa (INPP5B), nuclear gene encoding mitochondrial protein, mRNA.	3633 TR(1..4469)CDS(95..2836)	4380..4387,2642..2649	1.13	0.39	-0.68	-0.2	0.66
NM_001229	203984_s_at	CASP9	Homo sapiens caspase 9, apoptosis-related cysteine peptidase (CASP9), transcript variant alpha, mRNA.	842 TR(1..2034)CDS(96..1346)	1754..1760	1.02	0.45	-0.57	-0.18	0.66
NM_003318	204822_at	TTK	Homo sapiens TTK protein kinase (TTK), mRNA.	7272 TR(1..2984)CDS(75..2648)	2168..2174	1.17	0.36	0.08	-0.12	0.66
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)	2356..2362	0.18	0.67	0.16	-0.51	0.66
NM_015117	213445_at	ZC3H3	Homo sapiens zinc finger CCH-type containing 3 (ZC3H3), mRNA.	23144 TR(1..3302)CDS(30..2876)	3059..3065	1.84	0.07	0.08	0.02	0.66
NM_004161	207791_s_at	RAB1A	Homo sapiens RAB1A, member RAS oncogene family (RAB1A), mRNA.	5861 TR(1..2496)CDS(241..858)	1888..1894	0.81	0.53	-0.79	-0.23	0.66
XM_944210	228832_at	FLJ20021	PREDICTED: Homo sapiens hypothetical LOC90024 (FLJ20021), mRNA.	90024 TR(1..1138)CDS(1..789)	345..351	0.63	0.59	-0.05	-0.29	0.65
NM_178013	230087_at	PRIMA1	Homo sapiens proline rich membrane anchor 1 (PRIMA1), mRNA.	145270 TR(1..3650)CDS(43..504)	1551..1557,1542..1548,650..656,2662..2668	1.98	0.04	-1.82	0.02	0.65
NM_002951	208689_s_at	RPN2	Homo sapiens ribophorin II (RPN2), mRNA.	6185 TR(1..2509)CDS(289..2184)	1538..1544,313..319	1.64	0.09	-0.35	-0.26	0.65
NM_022825	219483_s_at	PORCN	Homo sapiens porcupine homolog (Drosophila) (PORCN), transcript variant A, mRNA.	64840 TR(1..1849)CDS(159..1511)	447..453,1345..1351,358..364	2.79	0	-0.42	1.44	0.65
NM_018698	209629_s_at,209628_at	NXT2	Homo sapiens nuclear transport factor 2-like export factor 2 (NXT2), mRNA.	55916 TR(1..2692)CDS(103..696)	1948..1954,1807..1814,1799..1806	1.19	0.34	-0.73	-0.14	0.64
NM_002500	1556057_s_at,206282_at	NEUROD1	Homo sapiens neurogenic differentiation 1 (NEUROD1), mRNA.	4760 TR(1..2641)CDS(237..1307)	951..958	1.47	0.16	-0.1	0.07	0.64
NM_153831	207821_s_at,208820_at	PTK2	Homo sapiens PTK2 protein tyrosine kinase 2 (PTK2), transcript variant 1, mRNA.	5747 TR(1..4453)CDS(231..3389)	132..138,4264..4270	1.57	0.1	-1.08	-0.03	0.64
NM_015431	215047_at	TRIM58	Homo sapiens tripartite motif-containing 58 (TRIM58), mRNA.	25893 TR(1..5158)CDS(49..1509)	1846..1852,1754..1760,3373..3379	1.73	0.01	-2.01	0.01	0.64
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)	2767..2773	0.66	0.57	-1.1	-0.92	0.64
NM_033450	213485_s_at	ABCC10	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 10 (ABCC10), mRNA.	89845 TR(1..5118)CDS(360..4754)	462..468,372..378,1014..1021,4133..4139,482..488	1.29	0.27	-0.73	-0.01	0.63
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)	552..559	1.16	0.36	-0.1	0.01	0.63
NM_022484	234726_s_at,218962_s_at	TMEM168	Homo sapiens transmembrane protein 168 (TMEM168), mRNA.	64418 TR(1..3951)CDS(393..2486)	3144..3150,3263..3269,1588..1594,3249..3255	1.5	0.13	-1.05	-0.12	0.63
NM_002971	203408_s_at	SATB1	Homo sapiens SATB homeobox 1 (SATB1), mRNA.	6304 TR(1..3782)CDS(246..2537)	894..900,1915..1921,1142..1148,3550..3556,1358..1364	1.27	0.28	-0.99	-0.05	0.63
NM_000237	203548_s_at,203549_s_at	LPL	Homo sapiens lipoprotein lipase (LPL), mRNA.	4023 TR(1..3747)CDS(371..1798)	1388..1395	1.64	0.02	-0.23	-0.11	0.63
NM_194330	218528_s_at,222660_s_at	RNF38	Homo sapiens ring finger protein 38 (RNF38), transcript variant 5, mRNA.	152006 TR(1..4999)CDS(339..1637)	244..250,1043..1049	1.47	0.13	-0.49	-0.07	0.61
NM_018438	231769_at	FBXO6	Homo sapiens F-box protein 6 (FBXO6), mRNA.	26270 TR(1..1535)CDS(136..1017)	827..833	0.16	0.66	0.25	0	0.61
NM_005993	211052_s_at,201759_at,229192_s_at	TBCD	Homo sapiens tubulin folding cofactor D (TBCD), mRNA.	6904 TR(1..4449)CDS(131..3709)	1934..1940	1.31	0.24	-0.56	-0.01	0.61
NM_004827	209735_at	ABCG2	Homo sapiens ATP-binding cassette, sub-family G (WHITE), member 2 (ABCG2), mRNA.	9429 TR(1..4445)CDS(494..2461)	496..503	1.57	0.01	-0.26	0.18	0.61
NM_014738	202650_s_at	KIAA0195	Homo sapiens KIAA0195 (KIAA0195), mRNA.	9772 TR(1..5472)CDS(276..4346)	4496..4502	1.12	0.37	-0.36	0.18	0.61
NM_001012969	239711_at	ADAL	Homo sapiens adenosine deaminase-like (ADAL), mRNA.	161823 TR(1..1622)CDS(421..1224)	1212..1219	0.63	0.57	-0.03	-0.01	0.59
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)	1788..1794,1550..1556,2695..2701	1.29	0.24	-0.3	-0.05	0.59
NM_144703	231200_at,219653_at,1553304_at	LSM14B	Homo sapiens LSM14B, SCD6 homolog B (S. cerevisiae) (LSM14B), mRNA.	149986 TR(1..2618)CDS(207..1364)	497..503	1.35	0.19	-0.16	-0.15	0.59
NM_004267	203921_at	CHST2	Homo sapiens carbohydrate (N-acetylglucosamine-6-O) sulfotransferase 2 (CHST2), mRNA.	9435 TR(1..3046)CDS(890..2482)	1577..1584	0.67	0.55	-0.02	-0.01	0.58
NM_144973	228551_at,243613_at,215058_at,238917_s_at	MGC24039	Homo sapiens hypothetical protein MGC24039 (MGC24039), mRNA.	160518 TR(1..9409)CDS(187..4011)	6019..6025	1.02	0.41	-0.42	-0.11	0.57
NM_002061	203925_at,236140_at	GLCLM	Homo sapiens glutamate-cysteine ligase, modifier subunit (GLCLM), mRNA.	2730 TR(1..3074)CDS(294..1118)		1.24	0.27	-0.47	-0.16	0.57
NM_015997	218914_at	C1orf66	Homo sapiens chromosome 1 open reading frame 66 (C1orf66), mRNA.	51093 TR(1..2258)CDS(620..2047)	448..454	1.2	0.3	-0.03	-0.01	0.57
NM_144679	227668_at	C17orf56	Homo sapiens chromosome 17 open reading frame 56 (C17orf56), mRNA.	146705 TR(1..2249)CDS(23..1600)	1187..1193	1.39	0.13	-0.2	0.14	0.56

NM_002390	207880_at,239837_at	ADAM11	Homo sapiens ADAM metalloproteinase domain 11 (ADAM11), mRNA.	4185 TR(1..4402)CDS(1..2310)	2407..2414	1.05	0.39	-0.48	-0.06	0.56
NM_013236	208833_s_at,208832_at	ATXN10	Homo sapiens ataxin 10 (ATXN10), mRNA.	25814 TR(1..2963)CDS(258..1685)	394..400	1.45	0.05	-0.17	-0.07	0.56
NM_004622	201513_at,201515_s_at,201504_s_at	TSN	Homo sapiens translin (TSN), mRNA.	7247 TR(1..3408)CDS(236..922)	2148..2154,1266..1272	1.2	0.29	-0.51	-0.15	0.56
NM_014908	204488_at	DOLK	Homo sapiens dolichol kinase (DOLK), mRNA.	22845 TR(1..2267)CDS(431..2047)	1203..1209,1302..1308,638..645	0.99	0.42	-0.19	0	0.56
NM_033386	221779_at,55081_at	MICALL1	Homo sapiens MICAL-like 1 (MICALL1), mRNA.	85377 TR(1..3882)CDS(85..2676)	3134..3140,3060..3066	1.13	0.34	-0.77	-0.08	0.55
NM_005497	243502_at,228563_at,208460_at,228776_at	GJA7	Homo sapiens gap junction protein, alpha 7, 45kDa (GJA7), transcript variant 1, mRNA.	10052 TR(1..7644)CDS(275..1465)	1701..1708,362..368	1.2	0.28	-0.8	-0.29	0.55
NM_007204	223331_s_at,224315_at	DDX20	Homo sapiens DEAD (Asp-Glu-Ala-Asp) box polypeptide 20 (DDX20), mRNA.	11218 TR(1..3449)CDS(294..2768)	168..174,2516..2522,2027..2033	1.31	0.19	-0.63	-0.12	0.54
NM_006038	204434_at,204433_s_at	SPATA2	Homo sapiens spermatogenesis associated 2 (SPATA2), mRNA.	9825 TR(1..4058)CDS(255..1817)	2806..2812,3425..3431,2674..2681,4028..4035	0.89	0.46	-0.93	-0.33	0.54
NM_001042437	203217_s_at	ST3GAL5	Homo sapiens ST3 beta-galactoside alpha-2,3-sialyltransferase 5 (ST3GAL5), transcript variant 2, mRNA.	8869 TR(1..2262)CDS(54..1241)	1914..1920	1.1	0.35	-0.6	-0.16	0.54
NM_033143	210310_s_at	FGF5	Homo sapiens fibroblast growth factor 5 (FGF5), transcript variant 2, mRNA.	2250 TR(1..5295)CDS(238..609)	399..405	1.43	0	-0.01	0	0.53
NM_002648	209193_at	PIM1	Homo sapiens pim-1 oncogene (PIM1), mRNA.	5292 TR(1..2684)CDS(364..1305)	1971..1977	0.76	0.51	-0.19	-0.37	0.53
NM_004199	202733_at	P4HA2	Homo sapiens procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), alpha polypeptide II (P4HA2), transcript variant 1, mRNA.	8974 TR(1..2588)CDS(565..2172)	1213..1220,1966..1972	0.95	0.43	0.11	-0.17	0.53
NM_020335	226029_at	VANGL2	Homo sapiens vang-like 2 (van gogh, Drosophila) (VANGL2), mRNA.	57216 TR(1..5415)CDS(487..2052)	2805..2811,2794..2800	0.8	0.49	-1.18	-0.14	0.53
NM_018180	218198_at	DHX32	Homo sapiens DEAH (Asp-Glu-Ala-His) box polypeptide 32 (DHX32), mRNA.	55760 TR(1..3070)CDS(492..2723)	2488..2494	1.21	0.26	-0.08	0.19	0.52
NM_020830	224800_at,233559_s_at	WDFY1	Homo sapiens WD repeat and FYVE domain containing 1 (WDFY1), mRNA.	57590 TR(1..4607)CDS(52..1284)	2343..2350	1.3	0.18	-0.2	-0.77	0.52
NM_015102	213471_at	NPHP4	Homo sapiens nephronophthisis 4 (NPHP4), mRNA.	261734 TR(1..5020)CDS(267..4547)	1336..1342,1245..1251,3663..3669	0.78	0.5	0.05	0.03	0.52
NM_002556	201800_s_at,201799_s_at	OSBP	Homo sapiens oxysterol binding protein (OSBP), mRNA.	5007 TR(1..5083)CDS(481..2904)	1942..1948,283..289	1.4	0.02	-0.77	-0.04	0.52
NM_016456	222988_s_at,222987_s_at	TMEM9	Homo sapiens transmembrane protein 9 (TMEM9), mRNA.	252839 TR(1..1592)CDS(120..671)	768..774	0.9	0.45	-0.11	-0.04	0.51
NM_014016	202797_at	SACM1L	Homo sapiens SAC1 suppressor of actin mutations 1-like (yeast) (SACM1L), mRNA.	22908 TR(1..3550)CDS(16..1779)	990..996	1.07	0.35	-0.96	-0.26	0.51
NM_016376	219868_s_at,224900_at	ANKFY1	Homo sapiens ankyrin repeat and FYVE domain containing 1 (ANKFY1), transcript variant 1, mRNA.	51479 TR(1..7959)CDS(39..3551)	332..339,3914..3920,3649..3655,738..744	1.28	0.19	-0.92	0.01	0.51