

Supplemental Table 2									
Site ID	Intron	Poly(A) site	Type	NM/KG	Inum	Region	Gene ID	Gene Symbol	Gene Annotation
Hs.335003.1.21	chr16:87882436:87880096	87880574	b	NM_013275	7	CDS	29123	ANKRD11	ankyrin repeat domain 11
Hs.445534.1.19	chr17:7988793:7987920	7988099	b	NM_002616	18	CDS	5187	PER1	period homolog 1 (Drosophila)
Hs.464912.1.22	chr18:31860860:31827262	31859562	b	NM_018170	6	CDS	55197	P15RS	hypothetical protein FLJ10656
Hs.497575.1.21	chr1:203016770:203020367	203017904	b	NM_015326	17	CDS	23380	SRGAP2	SLIT-ROBO Rho GTPase activating protein 2
Hs.527061.1.38	chr4:3469652:3478150	3470865	b	NM_198227	15	CDS	6002	RGS12	regulator of G-protein signalling 12
Hs.104672.1.3	chr3:101049828:101034796	101049470	c	NM_182909	5	3UTR	11259	DOC1	downregulated in ovarian cancer 1
Hs.104672.1.4	chr3:101049828:101034796	101049551	c	NM_182909	5	3UTR	11259	DOC1	downregulated in ovarian cancer 1
Hs.106019.1.66	chr6:30692233:30685712	30691583	c	NM_002714	2	5UTR	5514	PPP1R10	protein phosphatase 1, regulatory subunit 10
Hs.113876.1.37	chr4:1911391:1920029	1911499	c	NM_133333	12	3UTR	7468	WHSC1	Wolf-Hirschhorn syndrome candidate 1
Hs.113876.1.47	chr4:1911524:1920029	1917777	c	NM_133332	12	3UTR	7468	WHSC1	Wolf-Hirschhorn syndrome candidate 1
Hs.118738.1.3	chr3:51411205:51409596	51410323	c	NM_014703	24	3UTR	9730	VprBP	Vpr-binding protein
Hs.12420.1.21	chr4:130140986:130146615	130142683	c	NM_199320	8	CDS	79960	PHF17	PHD finger protein 17
Hs.128316.1.40	chr17:76704213:76704624	76704620	c	NM_017451	14	CDS	10458	BAIAP2	BAI1-associated protein 2
Hs.12967.1.101	chr6:152847043:152843959	152846778	c	NM_033071	34	CDS	23345	SYNE1	spectrin repeat containing, nuclear envelope 1
Hs.130690.1.16	chr13:99341693:99355524	99341868	c	NM_138280	8	3UTR	171425	CLYBL	citrate lyase beta like
Hs.13305.1.12	chr3:77749615:77754067	77750021	c	NM_002942	22	CDS	6092	ROBO2	roundabout, axon guidance receptor, homolog 2 (Drosophila)
Hs.133512.1.13	chr7:138203036:138197297	138202341	c	NM_020119	9	CDS	56829	ZC3HAV1	zinc finger CCCH-type, antiviral 1
Hs.134859.1.7	chr16:78190182:78185952	78187879	c	NM_005360	1	CDS	4094	MAF	v-maf musculoaponeurotic fibrosarcoma oncogene homolog (avian)
Hs.136885.1.2	chr13:25687610:25604590	25684910	c	NM_183045	5	CDS	6049	RNF6	ring finger protein (C3H2C3 type) 6
Hs.137183.1.21	chr11:116894390:116892665	116894372	c	NM_020693	7	CDS	57453	DSCAML1	Down syndrome cell adhesion molecule like 1
Hs.138207.1.37	chr2:27193924:27202012	27194051	c	NM_021831	11	CDS	60509	FLJ21839	hypothetical protein FLJ21839
Hs.138207.1.47	chr2:27204226:27204610	27204354	c	NM_021831	15	3UTR	60509	FLJ21839	hypothetical protein FLJ21839
Hs.159014.1.54	chr6:4008353:4010033	4008397	c	NM_176800	19	3UTR	8899	PRPF4B	PRP4 pre-mRNA processing factor 4 homolog B (yeast)
Hs.161220.1.8	chr13:31875044:31874039	31874973	c	AL049786	5	3UTR	90634	CG018	hypothetical gene CG018
Hs.165607.1.9	chr11:82320540:82346103	82323349	c	AK058145	7	CDS	220042	FLJ25416	hypothetical protein FLJ25416
Hs.172772.1.10	chr16:2761995:2761607	2761947	c	NM_207013	4	CDS	6923	TCEB2	transcription elongation factor B (SIII), polypeptide 2 (18kDa, elongin B)
Hs.178347.1.23	chr17:1347979:1346848	1347685	c	NM_016532	8	CDS	51763	SKIP	skeletal muscle and kidney enriched inositol phosphatase
Hs.178499.1.10	chr11:129608412:129606821	129608141	c	NM_014155	8	3UTR	29068	BTBD15	BTB (POZ) domain containing 15
Hs.179615.1.3	chr9:4590652:4586531	4590614	c	NM_017985	12	3UTR	55064	C9orf68	chromosome 9 open reading frame 68
Hs.184720.1.20	chr2:73712207:73738182	73712359	c	NM_015120	16	CDS	7840	ALMS1	Alstrom syndrome 1
Hs.188553.1.9	chr16:24467804:24472334	24468223	c	NM_006910	3	CDS	5930	RBBP6	retinoblastoma binding protein 6
Hs.196054.1.29	chr7:18479340:18540442	18481706	c	NM_058176	12	CDS	9734	HDAC9	histone deacetylase 9
Hs.201712.1.8	chr16:73044569:73043531	73044189	c	NM_012201	26	CDS	2734	GLG1	golgi apparatus protein 1
Hs.201921.1.20	chr1:112846188:112838791	112846146	c	NM_017744	6	CDS	54879	ST7L	suppression of tumorigenicity 7 like
Hs.202238.1.32	chr8:95592588:95592057	95592151	c	NM_015496	13	CDS	25962	DKFZP434I116	DKFZP434I116 protein
Hs.203514.1.10	chr20:367193:366243	367129	c	NM_144628	8	3UTR	128637	TBC1D20	TBC1 domain family, member 20
Hs.203514.1.5	chr20:366083:364615	364930	c	NM_144628	9	3UTR	128637	TBC1D20	TBC1 domain family, member 20
Hs.2076.1.7	chr8:146046736:146037933	146046657	c	NM_014066	2	3UTR	28991	COMMD5	COMM domain containing 5
Hs.21899.1.4	chrX:48518270:48516991	48517138	c	NM_005660	4	CDS	7355	SLC35A2	solute carrier family 35 (UDP-galactose transporter), member A2
Hs.229830.1.13	chr12:121836045:121836672	121836648	c	NM_032573	12	3UTR	84660	TSP-NY	testis-specific protein TSP-NY
Hs.24808.1.10	chr1:115033804:115031123	115033770	c	NM_025073	3	CDS	80143	FLJ21168	hypothetical protein FLJ21168
Hs.274337.1.10	chr14:44635712:44635839	44635819	c	NM_017922	4	CDS	55015	PRPF39	PRP39 pre-mRNA processing factor 39 homolog (yeast)
Hs.279257.1.11	chr6:150215667:150223865	150215779	c	NM_005389	7	3UTR	5110	PCMT1	protein-L-isoaspartate (D-aspartate) O-methyltransferase
Hs.282855.1.2	chr15:54173122:54167000	54170027	c	NM_022841	9	CDS	64864	RFXDC2	regulatory factor X domain containing 2
Hs.285313.1.4	chr10:3811538:3808596	3811212	c	NM_001008490	4	3UTR	1316	KLF6	Kruppel-like factor 6
Hs.287629.1.14	chr19:63202987:63191902	63200792	c	NM_025027	4	CDS	80095	ZNF606	zinc finger protein 606
Hs.291587.1.12	chr6:157348824:157497908	157348879	c	NM_017519	4	CDS	57492	ARID1B	AT rich interactive domain 1B (SWI1-like)

Hs.293884.1.2	chr2:240790461:240789436	240790237	c	NM_138336	3	CDS	150678	MYEOV2	myeloma overexpressed 2
Hs.301756.1.10	chr1:43519809:43519317	43519692	c	NM_052877	7	CDS	112950	MED8	mediator of RNA polymerase II transcription, subunit 8 homolog (yeast)
Hs.301756.1.9	chr1:43519809:43519317	43519544	c	NM_052877	7	CDS	112950	MED8	mediator of RNA polymerase II transcription, subunit 8 homolog (yeast)
Hs.304362.1.18	chr8:133892556:133896063	133893743	c	NM_016018	9	CDS	51105	PHF20L1	PHD finger protein 20-like 1
Hs.304362.1.19	chr8:133892556:133896063	133894621	c	NM_016018	9	CDS	51105	PHF20L1	PHD finger protein 20-like 1
Hs.305985.2.26	chr12:8983686:8984438	8984154	c	BX640639	17	3UTR	NA	NA	NA
Hs.309767.1.21	chr7:23645203:23717096	23645372	c	AK057430	4	CDS	56164	STK31	serine/threonine kinase 31
Hs.310536.1.31	chrX:71570759:71498140	71570456	c	NM_018486	4	CDS	55869	HDAC8	histone deacetylase 8
Hs.314338.1.15	chr21:39490293:39488548	39489921	c	AB080587	41	CDS	54014	BRWD1	bromodomain and WD repeat domain containing 1
Hs.314338.1.41	chr21:39592227:39590160	39590237	c	NM_018963	5	CDS	54014	BRWD1	bromodomain and WD repeat domain containing 1
Hs.32148.1.8	chr15:99630491:99629374	99629921	c	NM_203472	6	3UTR	55829	SELS	selenoprotein S
Hs.321653.1.24	chr1:158013377:158025339	158013609	c	NM_032174	10	3UTR	84134	TOMM40L	translocase of outer mitochondrial membrane 40 homolog-like (yeast)
Hs.326420.1.6	chr17:42308914:42317398	42312925	c	BC064534	4	CDS	7484	WNT9B	wingless-type MMTV integration site family, member 9B
Hs.343571.1.45	chr19:59349390:59350858	59350048	c	NM_014516	17	CDS	4849	CNOT3	CCR4-NOT transcription complex, subunit 3
Hs.346354.1.1	chr15:67459404:67464069	67459759	c	NM_017705	4	CDS	54852	PAQR5	progesterin and adipoQ receptor family member V
Hs.348921.1.7	chr6:64453772:64459585	64454184	c	NM_015153	3	CDS	23469	PHF3	PHD finger protein 3
Hs.350321.1.35	chr10:42942177:42943565	42942958	c	NM_020975	19	CDS	5979	RET	ret proto-oncogene (multiple endocrine neoplasia and medullary thyroid carcinoma 1, Hirschsprung disease)
Hs.351327.1.16	chr15:73419615:73420222	73419663	c	NM_017828	7	3UTR	54939	COMMD4	COMM domain containing 4
Hs.351593.1.6	chr4:155864294:155863591	155864040	c	NM_000508	5	CDS	2243	FGA	fibrinogen alpha chain
Hs.368305.1.5	chr4:80198866:80196214	80196275	c	BC031256	8	3UTR	152559	PAQR3	progesterin and adipoQ receptor family member III
Hs.368853.1.2	chr5:1514954:1514513	1514545	c	AK090444	12	3UTR	79888	FLJ12443	hypothetical protein FLJ12443
Hs.370549.1.8	chr2:60601067:60591453	60595980	c	NM_138559	4	CDS	53335	BCL11A	B-cell CLL/lymphoma 11A (zinc finger protein)
Hs.374421.1.31	chr9:78111174:78115310	78111535	c	BC091515	3	CDS	84131	C9orf81	chromosome 9 open reading frame 81
Hs.374503.1.8	chr15:76966747:76970385	76966747	c	NM_006791	5	CDS	10933	MORF4L1	mortality factor 4 like 1
Hs.380389.1.3	chr9:71711268:71707078	71709626	c	NM_016014	4	CDS	51104	C9orf77	chromosome 9 open reading frame 77
Hs.380774.1.46	chrX:40963423:40973339	40963640	c	NM_024005	17	3UTR	1654	DDX3X	DEAD (Asp-Glu-Ala-Asp) box polypeptide 3, X-linked
Hs.380774.1.50	chrX:40963423:40973339	40965711	c	NM_024005	17	3UTR	1654	DDX3X	DEAD (Asp-Glu-Ala-Asp) box polypeptide 3, X-linked
Hs.381219.1.8	chr3:23935742:23938092	23937338	c	BC081565	4	CDS	6138	RPL15	ribosomal protein L15
Hs.381870.1.5	chr11:65390891:65390760	65390872	c	NM_016938	11	3UTR	30008	EFEMP2	EGF-containing fibulin-like extracellular matrix protein 2
Hs.388245.1.2	chr4:174822893:174821259	174822391	c	NM_021973	2	3UTR	9464	HAND2	heart and neural crest derivatives expressed 2
Hs.388245.1.5	chr4:174822893:174821259	174822772	c	NM_021973	2	3UTR	9464	HAND2	heart and neural crest derivatives expressed 2
Hs.410596.1.20	chr17:59020332:59025296	59020546	c	NM_001003725	7	3UTR	10238	WDR68	WD repeat domain 68
Hs.412767.1.2	chr17:54431523:54415065	54430345	c	NM_001005207	24	CDS	4591	TRIM37	tripartite motif-containing 37
Hs.415846.1.3	chr21:46381326:46380837	46381121	c	NM_006657	14	3UTR	10841	FTCD	formiminotransferase cyclodeaminase
Hs.42644.1.16	chr10:131867675:131868366	131867859	c	NM_006541	11	3UTR	10539	TXNL2	thioredoxin-like 2
Hs.430551.1.35	chr15:88817247:88817950	88817314	c	NM_003870	19	CDS	8826	IQGAP1	IQ motif containing GTPase activating protein 1
Hs.435933.1.8	chr6:169927963:169923453	169924764	c	NM_018288	9	CDS	55274	PHF10	PHD finger protein 10
Hs.436680.1.11	chr17:58977039:58979894	58977820	c	NM_173092	14	3UTR	81033	KCNH6	potassium voltage-gated channel, subfamily H (eag-related), member 6
Hs.438823.1.6	chr7:150348791:150342842	150348243	c	NM_005692	15	CDS	10061	ABCF2	ATP-binding cassette, sub-family F (GCN20), member 2
Hs.440092.1.1	chr22:48916369:48917824	48916394	c	AF398511	2	5UTR	56666	PANX2	pannexin 2
Hs.440961.1.50	chr5:96134348:96134727	96134696	c	NM_173061	18	3UTR	831	CAST	calpastatin
Hs.445503.1.10	chr3:12199873:12203868	12202226	c	NM_133625	11	CDS	6854	SYN2	synapsin II
Hs.445893.1.19	chr1:32177414:32180024	32178568	c	AK091346	10	3UTR	10657	KHDRBS1	KH domain containing, RNA binding, signal transduction associated 1
Hs.446091.1.15	chr6:160139813:160144902	160140859	c	NM_004906	6	CDS	9589	WTAP	Wilms tumor 1 associated protein
Hs.446091.1.8	chr6:160139813:160144902	160139988	c	NM_004906	6	CDS	9589	WTAP	Wilms tumor 1 associated protein
Hs.446438.1.22	chr17:69954733:69956556	69955163	c	AK000249	4	CDS	55890	GPRC5C	G protein-coupled receptor, family C, group 5, member C
Hs.450230.1.11	chr7:45727643:45726674	45727083	c	NM_000598	4	3UTR	3486	IGFBP3	insulin-like growth factor binding protein 3

Hs.463928.1.26	chr17:7046945:7041096	7046761	c	NM_001365	10	CDS	1742	DLG4	discs, large homolog 4 (Drosophila)
Hs.464243.1.6	chr17:74303993:74295336	74303329	c	NM_025090	20	3UTR	57602	USP36	ubiquitin specific protease 36
Hs.467338.1.8	chr2:233461117:233459029	233460745	c	NM_002242	2	CDS	3769	KCNJ13	potassium inwardly-rectifying channel, subfamily J, member 13
Hs.470369.1.49	chr2:160086866:160082683	160086866	c	NM_013450	11	CDS	29994	BAZ2B	bromodomain adjacent to zinc finger domain, 2B
Hs.470544.1.10	chr2:170296742:170312790	170296748	c	NM_004792	9	CDS	9360	PPIG	peptidyl-prolyl isomerase G (cyclophilin G)
Hs.472854.1.17	chr20:43902827:43904078	43903773	c	NM_152897	4	CDS	90203	C20orf161	chromosome 20 open reading frame 161
Hs.473179.1.10	chr20:56255783:56254715	56255518	c	NM_018498	8	CDS	55370	PPP4R1L	protein phosphatase 4, regulatory subunit 1-like
Hs.474018.1.18	chr21:45466451:45469889	45466925	c	NM_015834	12	CDS	104	ADARB1	adenosine deaminase, RNA-specific, B1 (RED1 homolog rat)
Hs.475300.1.13	chr3:4330446:4332895	4331086	c	NM_006515	2	CDS	6419	SETMAR	SET domain and mariner transposase fusion gene
Hs.479298.1.16	chr4:20405290:20430456	20405440	c	AK098692	9	3UTR	133015	MGC29898	hypothetical protein MGC29898
Hs.479298.1.22	chr4:20405290:20430456	20406241	c	AK098692	9	3UTR	133015	MGC29898	hypothetical protein MGC29898
Hs.480615.1.13	chr4:120310786:120336158	120312487	c	NM_133477	4	CDS	171024	SYNPO2	synaptopodin 2
Hs.481927.1.60	chr5:37099838:37100385	37100337	c	NM_133433	46	CDS	25836	NIPBL	Nipped-B homolog (Drosophila)
Hs.483200.1.11	chr5:122365675:122371275	122365734	c	BC069012	6	3UTR	28966	SNX24	sorting nexin 24
Hs.484782.1.17	chr1:10457380:10456026	10457227	c	NM_004401	5	CDS	1676	DFFA	DNA fragmentation factor, 45kDa, alpha polypeptide
Hs.485628.1.26	chr6:57126805:57133840	57126892	c	NM_015555	12	CDS	26036	ZNF451	zinc finger protein 451
Hs.487925.1.46	chr1:142392120:142381634	142391742	c	NM_001002811	17	CDS	9659	PDE4DIP	phosphodiesterase 4D interacting protein (myomegalin)
Hs.488051.1.10	chr20:18071564:18073876	18071679	c	NM_020536	2	CDS	57325	CSRP2BP	CSRP2 binding protein
Hs.489132.1.8	chr7:93268314:93267874	93268189	c	BC012595	4	CDS	10282	BET1	BET1 homolog (S. cerevisiae)
Hs.494326.1.28	chr1:166088106:166096789	166089197	c	NM_003666	7	3UTR	8548	BLZF1	basic leucine zipper nuclear factor 1 (JEM-1)
Hs.494538.1.23	chr9:95323785:95322426	95323666	c	NM_000264	5	CDS	5727	PTCH	patched homolog (Drosophila)
Hs.498292.1.3	chr1:239989079:239977794	239989064	c	NM_181690	13	3UTR	10000	AKT3	v-akt murine thymoma viral oncogene homolog 3 (protein kinase B, gamma)
Hs.498740.1.2	chr10:13529271:13521504	13523094	c	NM_152751	8	CDS	222389	C10orf30	chromosome 10 open reading frame 30
Hs.5025.1.16	chr10:21142873:21141876	21142675	c	NM_006393	23	CDS	10529	NEBL	nebulin
Hs.50499.1.5	chr5:95020076:95018565	95019892	c	NM_031952	5	3UTR	83890	SPATA9	spermatogenesis associated 9
Hs.506216.1.17	chr12:80336218:80302151	80336105	c	NM_003625	7	CDS	8499	PPFIA2	protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (liprin), alpha 2
Hs.50841.1.23	chr15:55764165:55785423	55764850	c	NM_152451	13	3UTR	145781	Gcom1	GRIN1A combined protein
Hs.511956.1.13	chr2:28974987:28984366	28975316	c	NM_182756	7	CDS	245711	SPDY1	speedy homolog 1 (Drosophila)
Hs.513225.1.38	chr16:254909:258775	256115	c	AK057165	13	CDS	83986	C16orf9	chromosome 16 open reading frame 9
Hs.513265.1.2	chr16:1817868:1829263	1818909	c	BC063017	1	CDS	81889	FAHD1	fumarylacetoacetate hydrolase domain containing 1
Hs.514505.1.5	chr17:72226409:72224400	72226132	c	AB011157	6	CDS	23210	PTDSR	phosphatidylserine receptor
Hs.519672.1.67	chr5:149756599:149758109	149756599	c	NM_000356	23	CDS	6949	TCOF1	Treacher Collins-Franceschetti syndrome 1
Hs.520313.1.6	chr6:109796817:109796222	109796277	c	D14043	6	CDS	8763	CD164	CD164 antigen, sialomucin
Hs.520313.1.7	chr6:109796817:109796222	109796545	c	D14043	6	CDS	8763	CD164	CD164 antigen, sialomucin
Hs.520721.1.1	chr7:26929203:26934777	26929523	c	AK056230	1	CDS	NA	NA	NA
Hs.522684.1.41	chrX:70252331:70252041	70252206	c	NM_005096	7	CDS	9203	ZNF261	zinc finger protein 261
Hs.523036.1.19	chr1:19400729:19403022	19401084	c	NM_017765	9	3UTR	54896	PQLC2	PQ loop repeat containing 2
Hs.524496.1.9	chr1:36455904:36456417	36455920	c	NM_024676	9	3UTR	79729	C1orf113	chromosome 1 open reading frame 113
Hs.524750.1.52	chr11:3690369:3684434	3690021	c	NM_016320	20	CDS	4928	NUP98	nucleoporin 98kDa
Hs.526728.1.22	chr2:17806051:17804805	17806043	c	NM_024624	18	CDS	79677	SMC6L1	SMC6 structural maintenance of chromosomes 6-like 1 (yeast)
Hs.526972.1.3	chr6:76075275:76062387	76074526	c	AL832009	6	CDS	27145	FILIP1	filamin A interacting protein 1
Hs.530199.1.3	chr8:101604009:101592044	101602176	c	BC050396	5	CDS	157567	ANKRD46	ankyrin repeat domain 46
Hs.532091.1.22	chr5:77441786:77432606	77441786	c	NM_003664	20	CDS	8546	AP3B1	adaptor-related protein complex 3, beta 1 subunit
Hs.533582.1.18	chr1:100676840:100695827	100677037	c	NM_003672	15	CDS	8556	CDC14A	CDC14 cell division cycle 14 homolog A (S. cerevisiae)
Hs.535711.1.33	chr3:42226482:42235989	42229117	c	AB028965	14	CDS	22906	OIP106	OGT(O-GlcNAc transferase)-interacting protein 106 kDa
Hs.54609.1.26	chr22:36537233:36537435	36537397	c	AK123190	9	3UTR	23464	GCAT	glycine C-acetyltransferase (2-amino-3-ketobutyrate coenzyme A ligase)
Hs.546242.1.5	chr11:14946947:14945977	14946623	c	M64486	3	CDS	796	CALCA	calcitonin/calcitonin-related polypeptide, alpha

Hs.546255.1.4	chr4:155883653:155883112	155883333	c	NM_000509	9	CDS	2266	FGG	fibrinogen gamma chain
Hs.546361.1.49	chr3:132202887:132217693	132203319	c	NM_001001486	27	CDS	27032	ATP2C1	ATPase, Ca++ transporting, type 2C, member 1
Hs.546361.1.52	chr3:132202887:132217693	132203537	c	NM_001001486	27	CDS	27032	ATP2C1	ATPase, Ca++ transporting, type 2C, member 1
Hs.546361.1.57	chr3:132202887:132217693	132204676	c	NM_001001486	27	CDS	27032	ATP2C1	ATPase, Ca++ transporting, type 2C, member 1
Hs.549101.1.6	chr6:123715382:123700514	123712324	c	NM_006073	20	CDS	10345	TRDN	triadin
Hs.549166.1.6	chr13:19119432:19120561	19119432	c	NM_017520	3	CDS	54737	HSMPP8	M-phase phosphoprotein, mpp8
Hs.549177.1.3	chr10:69775464:69771052	69773281	c	NM_017987	18	3UTR	55680	RUFY2	RUN and FYVE domain containing 2
Hs.549204.1.15	chr6:32228726:32227931	32228559	c	BC013201	1	5UTR	80863	PRRT1	proline-rich transmembrane protein 1
Hs.549248.1.6	chr16:22002691:22004029	22003465	c	NM_173501	4	3UTR	146174	C16orf52	chromosome 16 open reading frame 52
Hs.550527.1.23	chr8:22507291:22507987	22507752	c	NM_176871	9	CDS	64236	PDLIM2	PDZ and LIM domain 2 (mystique)
Hs.550558.1.6	chr16:29824782:29824900	29824872	c	NM_198907	3	3UTR	253982	LOC253982	hypothetical protein LOC253982
Hs.554733.1.2	chr22:49297739:49297655	49297661	c	NM_152253	26	3UTR	1120	CHKB	choline kinase beta
Hs.68714.1.15	chr17:53437282:53436362	53436555	c	BC033785	3	3UTR	6426	SFRS1	splicing factor, arginine/serine-rich 1 (splicing factor 2, alternate splicing factor)
Hs.7155.1.2	chr2:6940617:6931408	6939049	c	NM_207315	5	3UTR	129607	LOC129607	hypothetical protein LOC129607
Hs.724.1.25	chr17:35499113:35502915	35502406	c	BC035137	9	CDS	7067	THRA	thyroid hormone receptor, alpha (erythroblastic leukemia viral (v-erb-a) oncogene homolog, avian)
Hs.72964.1.1	chr15:21362328:21406863	21364260	c	BC070293	1	CDS	7681	MKRN3	makorin, ring finger protein, 3
Hs.73965.1.6	chr17:72242737:72242240	72242514	c	BC066958	4	3UTR	6427	SFRS2	splicing factor, arginine/serine-rich 2
Hs.79769.1.8	chr5:141222980:141217221	141222403	c	NM_032420	3	CDS	5097	PCDH1	protocadherin 1 (cadherin-like 1)
Hs.89029.1.16	chr10:45455269:45442560	45455165	c	NM_174890	6	CDS	93550	ANUBL1	AN1, ubiquitin-like, homolog (Xenopus laevis)
Hs.91816.1.6	chr11:113775920:113767598	113775824	c	BC071695	1	CDS	54494	FLJ20010	hypothetical protein FLJ20010
Hs.9873.1.27	chr2:8876933:8876759	8876931	c	NM_020738	16	CDS	57498	KIDINS220	kinase D-interacting substance of 220 kDa
Hs.99491.1.21	chr11:64264995:64264212	64264917	c	NM_005825	6	CDS	10235	RASGRP2	RAS guanyl releasing protein 2 (calcium and DAG-regulated)
Hs.101774.1.9	chr20:16299230:16285098	16295488	s	NM_024704	22	CDS	55614	C20orf23	chromosome 20 open reading frame 23
Hs.102336.1.20	chr22:43451580:43502873	43454097	s	AL355192	7	CDS	55615	PRR5	proline rich protein 5
Hs.102929.1.4	chr3:48691000:48676572	48686285	s	AJ242655	12	CDS	51517	NCKIPSD	NCK interacting protein with SH3 domain
Hs.115474.1.19	chr13:33307355:33438223	33308514	s	NM_181558	8	3UTR	5983	RFC3	replication factor C (activator 1) 3, 38kDa
Hs.122926.1.3	chr5:142642325:142639170	142639323	s	NM_001020825	8	CDS	2908	NR3C1	nuclear receptor subfamily 3, group C, member 1 (glucocorticoid receptor)
Hs.125898.1.10	chr20:56907436:56911980	56908860	s	NM_000516	3	CDS	2778	GNAS	GNAS complex locus
Hs.128316.1.30	chr17:76696905:76704164	76698907	s	NM_017451	13	CDS	10458	BAIAP2	BAI1-associated protein 2
Hs.128856.1.9	chr8:27572976:27590004	27586454	s	NM_182826	5	CDS	51435	SCARA3	scavenger receptor class A, member 3
Hs.131055.1.4	chr8:21947733:21947966	21947753	s	NM_182795	6	CDS	10361	NPM2	nucleophosmin/nucleoplasmin, 2
Hs.134602.1.247	chr2:179443356:179432155	179435556	s	NM_003319	44	CDS	7273	TTN	titin
Hs.137556.1.18	chr10:85963085:85968814	85966258	s	BC038799	16	CDS	92211	PCDH21	protocadherin 21
Hs.14155.1.2	chr21:44474364:44467390	44471157	s	AF199028	6	CDS	23308	ICOSLG	inducible T-cell co-stimulator ligand
Hs.145150.1.8	chr2:230867758:230863181	230866698	s	NM_004509	14	CDS	3431	SP110	SP110 nuclear body protein
Hs.147062.1.16	chr16:56548745:56541946	56546080	s	NM_001297	12	CDS	1258	CNGB1	cyclic nucleotide gated channel beta 1
Hs.14945.1.5	chr5:131323127:131311255	131317497	s	AF099740	19	CDS	23305	ACSL6	acyl-CoA synthetase long-chain family member 6
Hs.149849.1.3	chr10:90064202:90024770	90034704	s	NM_018363	6	CDS	55328	C10orf59	chromosome 10 open reading frame 59
Hs.154276.1.17	chr21:29623886:29645825	29640179	s	NM_001011545	4	CDS	571	BACH1	BTB and CNC homology 1, basic leucine zipper transcription factor 1
Hs.154276.1.18	chr21:29623886:29645825	29640335	s	NM_001011545	4	CDS	571	BACH1	BTB and CNC homology 1, basic leucine zipper transcription factor 1
Hs.158560.1.42	chrX:70463675:70531412	70468876	s	AJ555149	10	CDS	6872	TAF1	TAF1 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 250kDa
Hs.158665.1.8	chr8:143854012:143843561	143849622	s	NM_023946	3	CDS	66004	LYNX1	Ly6/neurotoxin 1
Hs.16229.1.26	chr10:90672174:90722918	90673224	s	AB037794	10	CDS	57559	AMSH-LP	associated molecule with the SH3 domain of STAM (AMSH) like protein
Hs.172587.1.18	chr14:68004721:68130954	68014569	s	NM_133509	10	CDS	5890	RAD51L1	RAD51-like 1 (S. cerevisiae)
Hs.179646.1.12	chr15:36027190:36046117	36030913	s	AK093126	11	CDS	145942	MGC35118	hypothetical protein MGC35118

Hs.188634.1.28	chr15:62216185:62217293	62217258	s	AK128179	14	CDS	6642	SNX1	sorting nexin 1
Hs.192619.1.22	chr10:116611084:116649238	116614257	s	BC037207	16	CDS	57700	KIAA1600	KIAA1600
Hs.192619.1.23	chr10:116611084:116649238	116614466	s	BC037207	16	CDS	57700	KIAA1600	KIAA1600
Hs.21016.1.33	chr14:30425270:30433869	30429096	s	BC007230	10	CDS	1690	COCH	coagulation factor C homolog, cochlin (Limulus polyphemus)
Hs.21016.1.36	chr14:30425270:30433869	30429575	s	BC007230	10	CDS	1690	COCH	coagulation factor C homolog, cochlin (Limulus polyphemus)
Hs.227948.1.4	chr18:59475044:59456338	59473414	s	AF428135	6	CDS	6317	SERPINB3	serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 3
Hs.232416.1.4	chr17:43987891:43984994	43986646	s	NM_002146	2	5UTR	3213	HOXB3	homeo box B3
Hs.233495.1.10	chr7:29886989:29942121	29891692	s	BC053990	13	CDS	84725	PLEKHA8	pleckstrin homology domain containing, family A (phosphoinositide binding specific) member 8
Hs.254335.1.17	chr15:62282333:62259707	62260432	s	NM_001011664	10	CDS	53944	CSNK1G1	casein kinase 1, gamma 1
Hs.256747.1.5	chr9:35800243:35799510	35799814	s	NM_172312	6	CDS	26206	SPAG8	sperm associated antigen 8
Hs.269099.1.7	chr1:78126860:78125259	78126111	s	BC017247	19	CDS	8880	FUBP1	far upstream element (FUSE) binding protein 1
Hs.272759.1.6	chr12:122010204:122007000	122009726	s	NM_020845	8	CDS	57605	PITPNM2	phosphatidylinositol transfer protein, membrane-associated 2
Hs.28326.1.9	chr10:101466094:101463227	101464064	s	NM_004376	8	CDS	1355	COX15	COX15 homolog, cytochrome c oxidase assembly protein (yeast)
Hs.285354.1.2	chr14:64630178:64542756	64611601	s	NM_197957	3	CDS	4149	MAX	MYC associated factor X
Hs.288772.1.32	chr11:112740993:112744118	112742319	s	NM_017868	20	CDS	54970	TTC12	tetratricopeptide repeat domain 12
Hs.291363.1.21	chr22:27445784:27445466	27445507	s	AY551301	2	CDS	11200	CHEK2	CHK2 checkpoint homolog (S. pombe)
Hs.298469.1.58	chr17:58928079:58938148	58928713	s	NM_152831	13	CDS	1636	ACE	angiotensin I converting enzyme (peptidyl-dipeptidase A) 1
Hs.301048.1.25	chr18:12974190:12976860	12975718	s	NM_001013437	8	CDS	81929	SEH1L	SEH1-like (S. cerevisiae)
Hs.302047.1.2	chr3:156685846:156576281	156680372	s	AK022610	13	CDS	23007	PLCL3	phospholipase C-like 3
Hs.303197.1.5	chr16:30807681:30753971	30806618	s	BC058863	5	CDS	9274	BCL7C	B-cell CLL/lymphoma 7C
Hs.308122.1.5	chr14:92482428:92474625	92475962	s	BC003622	10	CDS	3705	ITPK1	inositol 1,3,4-triphosphate 5/6 kinase
Hs.3192.1.2	chr10:72314914:72312249	72313274	s	NM_001001939	3	CDS	5092	PCBD1	6-pyruvoyl-tetrahydropterin synthase/dimerization cofactor of hepatocyte nuclear factor 1 alpha (TCF1)
Hs.32099.1.14	chr8:382955569:38292718	38293092	s	NM_023034	9	CDS	54904	WHSC1L1	Wolf-Hirschhorn syndrome candidate 1-like 1
Hs.32099.1.15	chr8:382955569:38292718	38293474	s	NM_023034	9	CDS	54904	WHSC1L1	Wolf-Hirschhorn syndrome candidate 1-like 1
Hs.32099.1.17	chr8:382955569:38292718	38294237	s	NM_023034	9	CDS	54904	WHSC1L1	Wolf-Hirschhorn syndrome candidate 1-like 1
Hs.325667.1.2	chr13:51858163:51833190	51849303	s	AY358149	4	CDS	55901	THSD1	thrombospondin, type I, domain containing 1
Hs.32567.1.5	chr5:69358099:69373106	69364172	s	NM_021967	2	CDS	8293	SERF1A	small EDRK-rich factor 1A (telomeric)
Hs.332138.1.28	chr6:32238378:32240333	32239431	s	NM_138934	8	CDS	9374	PPT2	palmitoyl-protein thioesterase 2
Hs.332422.1.65	chr8:62756080:62743389	62750373	s	NM_020164	4	CDS	444	ASPH	aspartate beta-hydroxylase
Hs.356231.1.6	chr19:6395161:6389833	6391077	s	BC001656	10	CDS	79085	SLC25A23	solute carrier family 25 (mitochondrial carrier; phosphate carrier), member 23
Hs.357128.1.4	chr6:39985556:39980017	39981281	s	BC036839	9	CDS	4337	MOCS1	molybdenum cofactor synthesis 1
Hs.36587.1.20	chr2:241825847:241829185	241826574	s	NM_002712	8	CDS	5510	PPP1R7	protein phosphatase 1, regulatory subunit 7
Hs.368296.1.36	chr11:47139318:47142219	47140371	s	NM_001003678	8	CDS	79096	MGC4707	MGC4707 protein
Hs.370292.1.14	chr10:127512506:127519813	127515110	s	NM_016567	6	CDS	56647	BCCIP	BRCA2 and CDKN1A interacting protein
Hs.370510.1.7	chr11:114554573:114545557	114552229	s	BC047021	8	CDS	23705	IGSF4	immunoglobulin superfamily, member 4
Hs.370510.1.8	chr11:114554573:114545557	114552335	s	BC047021	8	CDS	23705	IGSF4	immunoglobulin superfamily, member 4
Hs.378532.1.47	chr6:135402403:135365674	135398696	s	NM_006620	4	CDS	10767	HBS1L	HBS1-like (S. cerevisiae)
Hs.380271.1.21	chr3:9773501:9782492	9774091	s	NM_016821	6	CDS	4968	OGG1	8-oxoguanine DNA glycosylase
Hs.380774.1.51	chrX:40963423:40973339	40965781	s	NM_024005	17	3UTR	1654	DDX3X	DEAD (Asp-Glu-Ala-Asp) box polypeptide 3, X-linked
Hs.389678.1.22	chr17:76018227:76024527	76018735	s	NM_173627	9	CDS	284131	FLJ35220	hypothetical protein FLJ35220
Hs.397705.1.5	chr1:6819537:7085629	6866346	s	NM_015215	3	CDS	23261	CAMTA1	calmodulin binding transcription activator 1
Hs.406062.1.3	chr19:5847463:5844302	5845688	s	AK127692	3	5UTR	126328	NDUFA11	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 11, 14.7kDa
Hs.424126.1.20	chr15:41872574:41879984	41873533	s	AK000438	2	CDS	10169	SERF2	small EDRK-rich factor 2
Hs.425769.1.3	chr1:58683752:58651662	58658412	s	AK091101	6	CDS	115209	OMA1	OMA1 homolog, zinc metalloproteinase (S. cerevisiae)

Hs.431861.1.24	chr19:51585230:51586247	51585943	s	BC000750	12	CDS	5536	PPP5C	protein phosphatase 5, catalytic subunit
Hs.433307.1.11	chr19:46576265:46608381	46581822	s	AK127598	2	CDS	593	BCKDHA	branched chain keto acid dehydrogenase E1, alpha polypeptide (maple syrup urine disease)
Hs.436896.1.6	chr10:79409904:79399820	79406754	s	BC014399	7	CDS	11128	POLR3A	polymerase (RNA) III (DNA directed) polypeptide A, 155kDa
Hs.438906.1.2	chr22:30408695:30397559	30401887	s	NM_173566	5	CDS	253143	MGC50372	hypothetical protein MGC50372
Hs.441550.1.5	chr20:25230854:25223667	25228850	s	NM_015600	13	CDS	26090	C20orf22	chromosome 20 open reading frame 22
Hs.442993.1.3	chr5:176667387:176666803	176666817	s	AK057034	5	CDS	83463	MXD3	MAX dimerization protein 3
Hs.44385.1.12	chr2:48810471:48871682	48818391	s	NM_172196	8	CDS	11036	ALF	TFIIA-alpha/beta-like factor
Hs.444451.1.25	chr2:173907486:173919923	173917378	s	NM_016653	11	CDS	51776	ZAK	sterile alpha motif and leucine zipper containing kinase AZK
Hs.444986.1.4	chr12:94372423:94382122	94379488	s	AK091730	2	CDS	10988	METAP2	methionyl aminopeptidase 2
Hs.449585.1.66	chr22:21566390:21573065	21568166	s	BC028090	4	CDS	3538	IGLC2	immunoglobulin lambda constant 2 (Kern-Oz- marker)
Hs.459790.1.46	chr9:77226558:77247705	77229657	s	NM_001018037	67	CDS	23230	VPS13A	vacuolar protein sorting 13A (yeast)
Hs.464243.1.9	chr17:74306076:74295336	74304745	s	AK022913	21	UTR	57602	USP36	ubiquitin specific protease 36
Hs.466871.1.7	chr19:48848216:48842541	48844575	s	NM_001005376	6	CDS	5329	PLAUR	plasminogen activator, urokinase receptor
Hs.467868.1.16	chr2:24298856:24307085	24304155	s	NM_198553	6	CDS	375190	FLJ30851	FLJ30851 protein
Hs.473317.1.15	chr20:62369808:62389475	62377818	s	BC032332	5	CDS	55251	C20orf36	chromosome 20 open reading frame 36
Hs.473819.1.7	chr21:38684786:38661441	38676732	s	AY204741	11	CDS	2078	ERG	v-ets erythroblastosis virus E26 oncogene like (avian)
Hs.474833.1.16	chr22:37019290:37015041	37016712	s	NM_001894	7	CDS	1454	CSNK1E	casein kinase 1, epsilon
Hs.479491.1.5	chr1:68315483:68276458	68303100	s	NM_001002292	11	CDS	79971	C1orf139	chromosome 1 open reading frame 139
Hs.479491.1.7	chr1:68315483:68276458	68303202	s	NM_001002292	11	CDS	79971	C1orf139	chromosome 1 open reading frame 139
Hs.480311.1.30	chr4:95864774:95896332	95866453	s	NM_001011513	7	CDS	10611	PDLIM5	PDZ and LIM domain 5
Hs.480712.1.22	chr4:129386074:129393329	129387054	s	NM_018078	9	CDS	55132	LARP2	La ribonucleoprotein domain family, member 2
Hs.483561.1.3	chr5:139554422:139603073	139588679	s	NM_032412	2	CDS	84418	ORF1-FL49	putative nuclear protein ORF1-FL49
Hs.485102.1.23	chr6:32056558:32089496	32057202	s	NM_004197	7	CDS	8859	STK19	serine/threonine kinase 19
Hs.487925.1.41	chr1:142381544:142375374	142379637	s	NM_014644	22	CDS	9659	PDE4DIP	phosphodiesterase 4D interacting protein (myomegalin)
Hs.49075.1.12	chrX:86726081:86730473	86729078	s	NM_057162	10	CDS	56062	KLHL4	kelch-like 4 (Drosophila)
Hs.491232.1.26	chr8:22331294:22347348	22336133	s	BC015770	8	CDS	23516	SLC39A14	solute carrier family 39 (zinc transporter), member 14
Hs.494312.1.34	chr9:84596555:84711711	84660172	s	NM_001018064	14	CDS	4915	NTRK2	neurotrophic tyrosine kinase, receptor, type 2
Hs.494312.1.37	chr9:84711901:84778630	84722365	s	NM_001018064	15	CDS	4915	NTRK2	neurotrophic tyrosine kinase, receptor, type 2
Hs.495391.1.32	chr1:171658344:171670070	171658976	s	AB007940	4	CDS	9910	RABGAP1L	RAB GTPase activating protein 1-like
Hs.497332.1.6	chr1:197349223:197325833	197344910	s	NM_031306	7	CDS	83479	DDX59	DEAD (Asp-Glu-Ala-Asp) box polypeptide 59
Hs.497417.1.4	chr14:74200354:74197397	74197715	s	NM_014821	19	CDS	9870	KIAA0317	KIAA0317
Hs.500741.1.9	chr10:99629995:99615462	99629397	s	NM_018058	14	CDS	55118	CRTAC1	cartilage acidic protein 1
Hs.501114.1.4	chr18:42168297:42267150	42171606	s	NM_152470	1	CDS	147341	C18orf23	chromosome 18 open reading frame 23
Hs.502.1.7	chr6:32905154:32898074	32904338	s	NM_018833	11	CDS	6891	TAP2	transporter 2, ATP-binding cassette, sub-family B (MDR/TAP)
Hs.502829.1.12	chr11:64290947:64289567	64289829	s	NM_201998	12	CDS	7536	SF1	splicing factor 1
Hs.503357.1.4	chr11:77258512:77306754	77261046	s	BC016854	3	CDS	28971	PTD015	PTD015 protein
Hs.503743.1.18	chr11:105286482:105294647	105289046	s	NM_000829	9	CDS	2893	GRIA4	glutamate receptor, ionotropic, AMPA 4
Hs.503878.1.41	chr11:112636256:112646119	112641123	s	NM_000615	15	CDS	4684	NCAM1	neural cell adhesion molecule 1
Hs.504538.1.2	chr12:3793463:3786531	3788288	s	BC031073	7	CDS	57097	PARP11	poly (ADP-ribose) polymerase family, member 11
Hs.504538.1.4	chr12:3793463:3786531	3791398	s	BC031073	7	CDS	57097	PARP11	poly (ADP-ribose) polymerase family, member 11
Hs.506458.1.6	chr12:97647600:97631881	97640220	s	NM_020140	10	CDS	56899	EB-1	E2a-Pbx1-associated protein

Hs.506458.1.7	chr12:97647600:97631881	97640462	s	NM_020140	10	CDS	56899	EB-1	E2a-Pbx1-associated protein
Hs.507537.1.25	chr13:24803697:24809073	24808482	s	NM_001008564	12	CDS	9818	NUPL1	nucleoporin like 1
Hs.509200.1.37	chr14:38887933:38925839	38889378	s	AF338234	23	CDS	4253	CTAGE5	CTAGE family, member 5
Hs.509200.1.38	chr14:38887933:38925839	38889603	s	AF338234	23	CDS	4253	CTAGE5	CTAGE family, member 5
Hs.509200.1.39	chr14:38887933:38925839	38889645	s	AF338234	23	CDS	4253	CTAGE5	CTAGE family, member 5
Hs.511951.1.3	chr12:121941618:121934238	121938426	s	AB112582	11	CDS	23457	ABC9	ATP-binding cassette, sub-family B (MDR/TAP), member 9
Hs.512690.1.11	chr2:85800234:85797637	85798003	s	NM_198843	10	3UTR	6439	SFTPB	surfactant, pulmonary-associated protein B
Hs.514227.1.10	chr17:40343508:40341044	40342570	s	NM_002055	7	CDS	2670	GFAP	glial fibrillary acidic protein
Hs.514846.1.13	chr18:42654893:42649311	42652312	s	NM_004671	12	CDS	9063	PIAS2	protein inhibitor of activated STAT, 2
Hs.517262.1.36	chr21:33853952:33861332	33854838	s	NM_003103	7	CDS	6651	SON	SON DNA binding protein
Hs.517262.1.5	chr21:33840556:33851331	33843745	s	AK127947	2	CDS	6651	SON	SON DNA binding protein
Hs.517949.1.64	chr3:47994358:47944845	47989580	s	NM_002375	3	CDS	4134	MAP4	microtubule-associated protein 4
Hs.518265.1.8	chr3:134788265:134789437	134788812	s	NM_017548	4	CDS	55573	H41	hypothetical protein H41
Hs.518595.1.14	chr4:4537727:4538582	4538345	s	NM_014392	5	3UTR	27065	D4S234E	DNA segment on chromosome 4 (unique) 234 expressed sequence
Hs.519452.1.23	chr5:170765013:170767308	170766335	s	NM_002520	9	CDS	4869	NPM1	nucleophosmin (nucleolar phosphoprotein B23, numatrin)
Hs.520146.1.16	chr6:43596765:43604964	43597162	s	NM_004875	8	CDS	9533	POLR1C	polymerase (RNA) I polypeptide C, 30kDa
Hs.520955.1.2	chr7:73911468:73886835	73900951	s	NM_148842	10	CDS	81554	WBSCR16	Williams-Beuren syndrome chromosome region 16
Hs.523772.1.22	chr1:175795010:175798619	175796787	s	AJ299441	5	CDS	64222	TOR3A	torsin family 3, member A
Hs.524690.1.20	chr1:39887818:39898453	39888468	s	NM_203456	9	CDS	10450	PPIE	peptidylprolyl isomerase E (cyclophilin E)
Hs.524809.1.18	chr12:121297915:121288573	121292195	s	NM_002956	20	CDS	6249	RSN	restin (Reed-Steinberg cell-expressed intermediate filament-associated protein)
Hs.532901.1.18	chr2:38143011:38205775	38156336	s	NM_144713	10	CDS	151393	FAM82A	family with sequence similarity 82, member A
Hs.532901.1.19	chr2:38143011:38205775	38156535	s	NM_144713	10	CDS	151393	FAM82A	family with sequence similarity 82, member A
Hs.533017.1.10	chr5:132124392:132115773	132121734	s	AF440762	9	CDS	23176	SEPT8	septin 8
Hs.533017.1.8	chr5:132124392:132115773	132119600	s	AF440762	9	CDS	23176	SEPT8	septin 8
Hs.533030.1.24	chr22:36479772:36486176	36480442	s	NM_007032	7	CDS	11078	HRIHFB2122	Tara-like protein
Hs.533600.1.19	chr11:65377006:65380175	65377744	s	NM_152760	10	3UTR	254122	FLJ30934	hypothetical protein FLJ30934
Hs.534847.1.59	chr6:32072817:32105636	32078421	s	NM_001002029	29	CDS	432395	C4B	complement component 4B, telomeric
Hs.546423.1.5	chr15:81012245:80616166	81009006	s	NM_030594	11	CDS	64506	CPEB1	cytoplasmic polyadenylation element binding protein 1
Hs.549094.1.7	chr9:97929278:97918833	97926190	s	AK097480	8	3UTR	9830	TRIM14	tripartite motif-containing 14
Hs.549105.1.2	chr4:974745:963123	971275	s	NM_134425	2	CDS	10861	SLC26A1	solute carrier family 26 (sulfate transporter), member 1
Hs.549169.1.19	chr6:34947681:34955763	34953269	s	NM_017754	20	CDS	54887	C6orf107	chromosome 6 open reading frame 107
Hs.549177.1.18	chr10:69809186:69806736	69806956	s	NM_017987	12	CDS	55680	RUFY2	RUN and FYVE domain containing 2
Hs.549181.1.2	chr21:32600922:32608732	32606462	s	NM_206898	4	CDS	56246	MRAP	melanocortin 2 receptor accessory protein
Hs.549306.1.26	chr6:117973454:117997475	117977653	s	NM_173674	14	CDS	285761	DCBLD1	discoidin, CUB and LCCL domain containing 1
Hs.549317.1.2	chr1:16471144:16468548	16470444	s	AK128106	15	3UTR	374955	SPATA21	spermatogenesis associated 21
Hs.80305.1.2	chr10:98978893:98914638	98971928	s	AY336750	11	CDS	84986	ARHGAP19	Rho GTPase activating protein 19
Hs.82201.1.10	chr16:56755458:56749759	56753925	s	NM_001896	11	3UTR	1459	CSNK2A2	casein kinase 2, alpha prime polypeptide
Hs.82201.1.8	chr16:56755458:56749759	56750986	s	NM_001896	11	3UTR	1459	CSNK2A2	casein kinase 2, alpha prime polypeptide
Hs.87306.1.2	chr4:5685056:5663012	5682238	s	AY152403	21	CDS	132884	EVC2	Ellis van Creveld syndrome 2 (limbin)
Hs.9573.1.11	chr6:30656276:30658821	30658238	s	NM_001090	8	CDS	23	ABCF1	ATP-binding cassette, sub-family F (GCN20), member 1
Hs.97858.1.54	chr1:10369698:10372662	10370969	s	AB017133	47	CDS	23095	KIF1B	kinesin family member 1B
Hs.98510.1.26	chrX:117360340:117372769	117365780	s	AK127556	19	CDS	54521	WDR44	WD repeat domain 44
Hs.9933.1.16	chr1:6626822:6628345	6627888	s	NM_138350	4	CDS	90326	THAP3	THAP domain containing, apoptosis associated protein 3