

Fold change										Annotations												
ProbeName	[103T] vs [103N]	[104T] vs [104N]	[109T] vs [109N]	[110T] vs [110N]	[119T] vs [119N]	[120T] vs [120N]	type	seqname	GeneSymbol	source	RNAlength	chrom	strand	txStart	txEnd	Xhyb	probeSeq	EntrezID	unigene	GO	ProteinAcce	product
ASHG19A3A029250	3.0334985			-2.4515448		8.88348	coding	NM_031266	HNRNPAB	RefSeq_codi	1837	chr5	+	177631507	177638183		TCAGAGTC3182		Hs.591731	GO:0003688NP_112556		Heterogeneous nuclear ribonucleoprotein A/B
ASHG19A3A005217	5.2145667	2.1745803				3.1623528	coding	NM_021644	HNRNP3	RefSeq_codi	2382	chr10	+	70091767	70102952		GGTTATTA3189		Hs.643472	GO:0003723NP_067676		Heterogeneous nuclear ribonucleoprotein H3 (2H9)
ASHG19A3A016647		3.0594022			2.4225197	5.413126	coding	NM_194247	HNRNP3	RefSeq_codi	5808	chr2	+	178077421	178088685		GAATGT220988		Hs.516539	GO:0003723NP_919223		Heterogeneous nuclear ribonucleoprotein A3
ASHG19A3A001508		-2.0576186				-2.2072835	coding	NM_001077	HNRNPC	RefSeq_codi	3226	chr14	-	21677297	21737638		AACACCC3183		Hs.508848	GO:0003723NP_112604		Heterogeneous nuclear ribonucleoprotein C (C1/C2)
ASHG19A3A003913						-2.5210278	coding	NM_001533	HNRNPL	RefSeq_codi	2129	chr19	-	39327028	39340617		gcgggggggc3191		Hs.644906	GO:0003723NP_001524		Heterogeneous nuclear ribonucleoprotein L
ASHG19A3A005448						-4.5613346	coding	NM_031262	HNRNPK	RefSeq_codi	2935	chr9	-	86582998	86595569		ATGCACAC3190		Hs.522257	GO:0003677NP_112553		Heterogeneous nuclear ribonucleoprotein K
ASHG19A3A041305						-2.266011	coding	NM_001032	HNRNP2	RefSeq_codi	2315	chrX	+	100663190	100669115		CATGTTTG6173		Hs.432485	GO:0005515NP_066357		Ribosomal protein L36a
ASHG19A3A012167						-2.5504644	coding	NM_005968	HNRNPM	RefSeq_codi	2553	chr19	+	8509802	8553994		TGATTAC14670		Hs.465808	GO:0003723NP_112480		Heterogeneous nuclear ribonucleoprotein M
ASHG19A3A011973		-5.739483				-2.3587754	coding	NM_002819	PTBP1	RefSeq_codi	3340	chr19	+	797391	812327		GGTATAA15725		Hs.172550	GO:0003723NP_787041		Poly(pyrimidine tract binding protein 1
ASHG19A3A003535		-2.5619657				-2.3441238	coding	NM_001170	SRPK3	RefSeq_codi	1900	chrX	+	153046455	153051186		GGCAGCG26576		Hs.104865	GO:0005524NP_055185		SRSF protein kinase 3
ASHG19A3A033341						-2.0918567	coding	NM_182692	SRPK2	RefSeq_codi	3671	chr7	-	104756823	105029341		CGTCGCC6733		Hs.285197	GO:0005524NP_872634		SRSF protein kinase 2
ASHG19A3A022042		2.430689				-9.527317	coding	NM_004593	TRA2B	RefSeq_codi	4290	chr3	-	185632359	185655924		CTCTATAA6434		Hs.531322	GO:0003723NP_004584		Transformer 2 beta homolog (Drosophila)
ASHG19A3A043221		2.7555702				2.080928	coding	NM_005066	SFPQ	RefSeq_codi	3073	chr1	-	35649202	35658743		AAAGGCA 6421		Hs.355934	GO:0003677NP_005057		Splicing factor proline/glutamine-rich
ASHG19A3A005770						2.4541771	coding	NM_130438	DYRK1A	RefSeq_codi	5088	chr21	+	38792600	38887678		ACCTCCA1859		Hs.368240	GO:0005524NP_569122		Dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 1A
ASHG19A3A017619		2.0076218				3.0076554	coding	NM_004902	RBM39	RefSeq_codi	2791	chr20	+	34291531	34330193		GGGTTAC19584		Hs.282901	GO:0003723NP_909122		RNA binding motif protein 39
ASHG19A3A013371						4.3147182	coding	NM_016047	SF3B14	RefSeq_codi	783	chr2	-	24290454	24299314		CTATCGGC51639		Hs.177861	GO:0003723NP_057131		Splicing factor 3B, 14 kDa subunit
ASHG19A3A001360		2.8628976					coding	NM_001040	U2AF1L4	RefSeq_codi	792	chr19	-	36233429	36236336		CATGCTG1199746		Hs.351558	GO:0003723NP_659424		U2 small nuclear RNA auxiliary factor 1-like 4
ASHG19A3A004544		2.7370296				-2.2739232	coding	NM_006758	U2AF1	RefSeq_codi	971	chr21	-	44513065	44527688		TCGGTTGC7307		Hs.365116	GO:0003723NP_006749		U2 small nuclear RNA auxiliary factor 1
ASHG19A3A013081						2.433374	coding	NM_001012	U2AF2	RefSeq_codi	3136	chr19	+	56165415	56186081		CTACTCTC11338		Hs.528007	GO:0070742NP_009210		U2 small nuclear RNA auxiliary factor 2
ASHG19A3A030640		4.224404	3.651587	2.7329814		2.056424	coding	NM_004698	PRPF3	RefSeq_codi	2480	chr1	+	150293927	150325703		AAATTTG9129		Hs.11776	GO:0005515NP_004689		PRP3 pre-mRNA processing factor 3 homolog (S. cerevisiae)
ASHG19A3A039127		2.1522481				-2.5694952	coding	NM_004697	PRPF4	RefSeq_codi	2756	chr9	+	116037973	116055054		AGGAGCA.9128		Hs.530284	GO:0005515NP_004688		PRP4 pre-mRNA processing factor 4 homolog (yeast)
ASHG19A3A045239						-2.023461	coding	NM_014502	PRPF19	RefSeq_codi	2342	chr11	-	60658020	60674061		TTGACAA/27339		Hs.502705	GO:0003677NP_055317		PRP19/PSO4 pre-mRNA processing factor 19 homolog (S. cerevisiae)
ASHG19A3A013003		2.7194836					coding	NM_015629	PRPF31	RefSeq_codi	2201	chr19	+	54618789	54635150		TAAGTAC126121		Hs.515598	GO:0003723NP_056444		PRP31 pre-mRNA processing factor 31 homolog (S. cerevisiae)
ASHG19A3A054764		2.7828064	3.0183098	2.0387087		4.2496533	coding	NM_012426	SF3B3	RefSeq_codi	9736	chr16	+	70557690	70611570		AAGACCC23450		Hs.514435	GO:0003676NP_036558		Splicing factor 3b, subunit 3, 130kDa
ASHG19A3A043766		2.15882					coding	NM_006802	SF3A3	RefSeq_codi	2855	chr1	-	38422653	38455761		GAGACCC10946		Hs.77897	GO:0046872NP_006793		Splicing factor 3a, subunit 3, 60kDa
ASHG19A3A004721						-3.1859674	coding	NM_014281	PUF60	RefSeq_codi	1853	chr8	-	144898548	144911537		GCAGCAG22827		Hs.521924	GO:0003677NP_510965		Poly-U binding splicing factor 60kDa
ASHG19A3A037139		2.0154915	2.1092439	2.1071062		2.1713371	coding	NM_021144	PSIP1	RefSeq_codi	1690	chr9	-	15470643	15510287		CTGAGAA111168		Hs.726445	GO:0003677NP_150091		PC4 and SFRS1 interacting protein 1
ASHG19A3A012901		-2.9223475					coding	NM_021228	SCAF1	RefSeq_codi	4323	chr19	+	50145381	50161905		GCCATAC158506		Hs.103521	GO:0003723NP_067051		SR-related CTD-associated factor 1
ASHG19A3A045058		2.3553817					coding	NM_001033	CSF3	RefSeq_codi	655	chr11	-	33162977	33183037		GGTCTTTT1479		Hs.44402	GO:0003723NP_001317		Cleavage stimulation factor, 3' pre-RNA, subunit 3, 77kDa
ASHG19A3A042801						2.0473845	coding	NM_015235	CSF2T	RefSeq_codi	4127	chr10	-	53455246	53459355		CATTTTGA23283		Hs.591358	GO:0003723NP_056050		Cleavage stimulation factor, 3' pre-RNA, subunit 2, 64kDa, tau variant
ASHG19A3A004821		3.4057038	2.4724438			2.0964541	coding	NM_015484	SYF2	RefSeq_codi	1777	chr1	-	25548766	25559013		AAATCCC25949		Hs.20013	GO:0008380NP_997053		SYF2 homolog, RNA splicing factor (S. cerevisiae)