

symbol	chr	gene	regionStart	regionEnd	Zscore	validated	OnArray	MiDAS-significant	SplicingIndex	primer-R1	Primer-S2	primer-3
NASP	1	ENSG00000132780	45845580	45846596	10,2	yes	yes	no	1,1	CCAGCATATCCCAGGCAAG	CAGATGAAAGAGGGTGAAGAAA	GTAGAAAATAATGATAACATAGATGAAA
NRAS	1	ENSG00000009307	115081615	115081707	-3,59	no	yes	no	-0,39	AATGGCAACCTGCAAGACTT	GAACAGCGCACACAACCTTG	GTAAGTCAGATAAAACACTTCTTG
RCC1	1	ENSG00000180198	28729622	28729714	25,08	yes	yes	no	-0,014	CCTCAGCCTGCACAACATC	CAGAAAACCCGACCAGTCT	AAGCAAGAAGGTGAAGGTCT
EIF2A	3	ENSG00000144895	151752833	151752902	-3,52	yes	yes	no	0,05	TTCAAGGCAAACCTGCCTTCA	AGCACAGTGTTCGAAGGGA	GCCGCTCTTGACAGGGA
MDC1	6	ENSG00000137337	30783322	30784113	-2,61	yes	yes	yes	-0,55	AGATCTGGACCTACAAGCTAC	ATGGTTCATCTAGGGTACCTG	TGCAATTCAGGAGGCCTG
QKI	6	ENSG00000112531	163876004	163876147	-2	yes	no	no		GGCCCAGTAATGATCCTTGG	GAAATTATTGGTACCTGCAGCA	CAATGAGGGATAAAAAAAGGCA
WNK2	9	ENSG00000165238	95036911	95037068	3,37	yes	no	no		TGTGTCTCTGCGAAGAAG	CGCAAGGTCACCTGTGGT	TTGCCAAAAGTGTGATAGGGT
LARP5	10	ENSG00000107929	866807	866917	-3,2	yes	no	1,1		GCACTGATGTGGACTTGATTG	CTTTAAATAGTGCTTCTACTTCTTC	CTCGAAGGTATTTGTAAGCTTC
MKI67	10	ENSG00000148773	129803182	129804261	5,2	yes	yes	yes	1,3	ATGTAATAACTATTATTGATCGTTCC	GAATCTTGAGCTTTCATCAG	GGTATTCCTCACTCTCATCAG
PPP3CB	10	ENSG00000107758	74908263	74908387	-2,8	no	yes	no	0,26	GGATGGGATACCCAGGGTTG	CATGGATGTCACCACACACTG	GAACCCATAAATAAGACACACTG
PKM2	15	ENSG00000067225	70282417	70282583	-5,65	no	yes	no	-1,2	TCAAGAAGCCCCGCCCC	CTGCCTCACGGGCAATCAG	TCAGCCTCACGAGCTATCAG
NP_006653.1	16	ENSG00000080603	30652716	30652830	-4,14	no	yes	no	-0,17	TCATCTTCTGCCCGACACA	ACAGCCTATTTCAAACAGCAG	GGGATGTCCACATATATAGCAG
EIF4A1	17	ENSG00000161960	7418588	7418720	-3,3	yes	yes	no	-1,9	GTATCTGCTGAGCCAATTCTC	CATTCTACCTTGTATCAAGGGTT	GAAGGCGTCATCGAGGTT
SUPT4H1	17	ENSG00000108375	53779889	53779944	-3,17	yes	no	no		CATCTGCGGGCCTGTTTG	CTCATCATCGCAATGATTCCA	CACCTGGCTTAAAGTTACCCA
ILF3	19	ENSG00000129351	10652022	10652119	2,14	yes	yes	no	0,53	CTTGGTGGGATCTGAACGG	CACAGAGTGCGCAGCAC	GCATCGTGATGCCAGCAC
UPF1	19	ENSG00000005007	18819553	18819642	4,8	no	yes	no	0,1	GATGAAGCCGAGGAGGAAG	GGAAATACTTCTGGCAGCCA	ATACACGCCTGCAGCCA