

Table S5. Primers used for conventional PCR.

Gene	Primer	Sequence (5' → 3')	Position	Accession number	Cycles
ADAR2	S	CACCTCTCTACACCCTCAACAAG	2245-2267	gi:75709170	30
	AS	CGTAGTAAGTGGGAGGGAACCTT	2457-2479		
ASCC3L1	S	GTGTCTGCTCTGAGAAACAGTGC	4011-4033	gi:40217846	25
	AS	CTGTCAGGAGTACCACCTTCTTG	4292-4314		
GAPDH	S	ACTTTGTCAAGCTCATTTC	1019-1038	gi:83641890	25
	AS	CACAGGGTACTTTATTGATG	1284-1303		
MARS	S	CTCCAGCACTATCACCAGCTACT	2088-2110	gi:14043021	20
	AS	GTGTCCTGCTGGTAAGGTACACA	2380-2402		
MRPL3	S	GCCTATACATCCTTGGGAAACCAG	419-441	gi:21265090	25
	AS	GTGAGCAGCATAAAGAGGAGTGC	733-755		
PABPC1	S	CAGGCTCACCTCACTAACCAGTA	1627-1649	gi:56676313	20
	AS	GAACCTGTGAAGAAGCTGGTCTC	1902-1924		
RAE1	S	ATGACTGGGAGCTGGGATAAGAC	798-820	gi:62739174	25
	AS	AGCAACTCTCCCCTCGATACTTC	1066-1088		
RNPS1	S	TAGGGCTCCTTACCTACCAAAC	320-342	gi:18379335	25
	AS	AAGGACTGGAGAGCCAGAAGAG	608-630		
SNRPB	S	CAGAAAGGGAAGAGAAGCGAGTC	339-361	gi:38150006	20
	AS	GTACCTCTTCTTGTTGGGGTCAT	587-609		
SNRPC	S	CTACTCCATTCTCTGCTCCTCCT	206-228	gi:4507126	25
	AS	CCTCCCCTCTATCCTTATCTGTC	487-509		
SNRPE	S	TAGATCGCGGATTCAGGTG	129-147	gi:61098048	25
	AS	AGCATGATCCGACCCAGTT	266-284		

Sense (S) and antisense (AS) primer sequences are shown. Their position in the transcript, the accession number of the transcript, and the number of cycles used in each PCR are also shown.