

Species	Product length	Forward Primer				Reverse Primer				RNAfold		RNAcofold			RNA duplex		RNAcofold	
		Seq	length	Tm	GC	Seq	length	Tm	GC	for	rev	for	rev	for/rev	forward/total seq	reverse/total seq	forward/seqpart	reverse/seqpart
Bos taurus	65	ATGGGGTAGCGCTGGGGAC	19	57.6	0.68	TCCACGACTCGCATGGAGGA	20	55.88	0.6	0.00	-2.7	-12.30	-13.4	-9.3	-42.7	-41.8	-42.0	-41.0
Callithrix jacchus	79	GATGGTGGCCATTTGGGAG	20	53.83	0.55	GGGGCAGGGGTCTAGTCA	18	54.9	0.67	-1.7	-2.5	-16.10	-12.7	-9.4	-40.0	-40.6	-38.2	-39.7
Oryctolagus cuniculus	71	TGGGTGGCATTTGGGGACA	20	53.83	0.55	AGGGCTGATGCACTGGACAC	20	55.88	0.6	-1.6	-1.9	-9.60	-8.7	-8.4	-40.0	-42.1	-39.4	-41.4
Pongo pygmaeus	72	TGGTGGCCTTTTTGGGGACA	20	53.83	0.55	GTGTCGAGTCACCGACCACA	20	55.88	0.6	-2.5	-5.3	-13.50	-16.0	-12.8	-39.7	-41.1	-39.2	-40.2
Rattus norvegicus	67	ATGGGGACCTCTTGGGGACA	20	55.88	0.6	AAACATGACAGGCCACACGG	20	53.83	0.55	-3.8	0.0	-14.30	-9.7	-7.7	-43.0	-38.6	-42.2	-37.8
Vicugna pacos	76	TTATGGGGGTAGGGCTGGGG	20	57.93	0.65	CGGGTCGGGTCCACGATTTG	20	57.93	0.65	0.0	-5.6	-5.10	-19.4	-10.5	-44.8	-41.3	-43.8	-40.5