

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	167	TCGACCAGATGACTCCC	19	55.4	0.63	AAGGCCCATTAAGGGCAACG	20	53.83	0.55	-1.6	-7.6	-9.1	-19.0	-5.8	-40.4	-29.2	-39.6	-37.7
Callithrix jacchus	134	TCGACCAGTCAGCCCTGGGT	20	57.93	0.65	AAGATATACCCCGCCCAGGG	20	55.88	0.6	-4.9	-3.5	-13.3	-9.4	-16.1	-44.4	-41.3	-43.6	-40.9
Oryctolagus cuniculus	167	TCGACCAGGTGGCCAGAG	18	54.9	0.67	CACCTACCTGCAGGTCCAAA	20	53.83	0.55	-1.1	-2.4	-11.7	-18.3	-12.6	-39.4	-39.0	-38.6	-38.9
Pongo pygmaeus	145	GTTCCGTGTCTGGCTGCCAT	20	55.88	0.6	TGGTGGCCCAAATGCCAAC	20	53.83	0.55	-0.6	-4.2	-13.5	-12.5	-14.1	-42.5	-39.3	-40.5	-38.4
Rattus norvegicus	105	CGACCAGTTGTTCTTTGCG	20	53.83	0.55	AAACATGACAGGCCACACGG	20	53.83	0.55	0.0	0.0	-5.8	-9.7	-4.3	-37.6	-38.6	-36.2	-37.8
Vicugna pacos	164	TCGATCAGATGGCCCCCGA	19	55.4	0.63	TCTGCTGGCAGAAAATGCC	20	53.83	0.55	0.0	-5.1	-7.5	-20.7	-5.1	-40.7	-39.4	-40.1	-38.8