

<b>Strong stop</b> (137 bases) <sup>a</sup>		
Forward Primer	oHC64	5' TAACTAGGGAACCCACTGC
Reverse Primer	oHC65	5' GCTAGAGATTTTCCCACTG
Probe	oHC66	5' FAM-ACACAACAGACGGGCACACACTA-TAMRA
<b>tRNA<sup>lys3</sup></b> (1 base) <sup>a</sup>		
Forward Primer	tRNA-15for	5' GTCGGTAGAGCATCAGACTTTTAATCT
Reverse Primer	Long PBS-G-(C)7rev	5' CCCCCCGTGGCGCCCGAACAGGGACTTGAAC
Probe	Probe-tRNA43for	5' FAM-AGGGTCCAGGGTTC-MGB
<b>Set 2</b> (16 bases) <sup>a</sup>		
Forward Primer	tRNA-15for	5' GTCGGTAGAGCATCAGACTTTTAATCT
Reverse Primer	PBSU5-rev	5' GGAAAATCTCTAGCAGTGGCG
Probe	Probe-tRNA43for	5' FAM-AGGGTCCAGGGTTC-MGB
<b>Set 3</b> (36 bases) <sup>a</sup>		
Forward Primer	tRNA-41for	5' TGAGGGTCCAGGGTTCAAGT
Reverse Primer	U5-rev-3	5' CAGACCCCTTTTAGTCAGTGTGGAA
Probe	Probe-PBSU5f-3	5' FAM-CCTGTTCGGGCGCCACTGCT-BHQ1

**Table S1.** Names and sequences of qPCR primers and probes used.

<sup>a</sup> Measured number of bases of cDNA from the 3' end of the tRNA<sup>lys3</sup> primer.