

Table S4. Sequencing primers.

Gene	Primer (5' -> 3') ¹	T _m (°C)	Length (bp)
<i>foraging</i>	F1: TCCCAAATACGATAAGGATG	55.11	821
	R1: CATCACGGAATTCTTCATTG	56.55	
	F2: ACTTGCTTGGTCATAGATCG	55.41	946
	R2: TTTGTAAACCCACCAATAG	55.12	
<i>Glutamate dehydrogenase</i>	F1: AAAGGCTTTGCCTACTGCTG	59.65	975
	R1: TGGTACCGTGTTTCGTTCTTG	59.61	
	F2: AACGAAGCCAACTACATGAGC	59.40	808
	R2: CATCACGGTAAGTGGTGAAG	56.59	
<i>Jonah65Aiv</i>	F: GCCTGGTGTGGTGGTTC	57.00	621
	R: CGATACCAGTATTAGATTTAATCC	57.00	
<i>Malvolio</i>	F1: CGTAAATTATGGGCCTTCAC	56.59	836
	R1: GAATGTTTGGCATCTTCRTAC	57.14	
	F2: GTTTTTGCTCATGGCATG	56.03	661
	R2: CGGCCATATGTATCACAAG	54.93	
<i>RPS6-p70-protein kinase</i>	F1: ATGGCTGACGTGASYG	51.18	790
	R1: AAGCACCTAATGACCACC	53.23	
	F2: CGTGGTCATGGTAAAGC	53.01	644
	R2: TGTACATCCATCATTTTCGTC	54.21	

¹ Primer sequences (F= forward; R= reverse). For sequences amplified in two discontinuous segments: F1= first forward fragment; R1= first reverse fragment; F2= second forward fragment; R2= second reverse fragment;)