Bactrocera tryoni

Taxonomic lineage: Brachycera > Muscomorpha > Tephritoidea > Tephritidae > Bactrocera > Bactrocera subgenus Bactrocera

Assembly: GCA_016617805.2_CSIRO_BtryS06_freeze2_genomic

Bac_try1 | CM028324.1:8522465-8523381 (-) | 917 nt | IncRNA:noe consensus e-value: 1.5e-19

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 8nt PSE: 0.84

Bac_try2 | CM028321.1:10235625-10236064 (-) | 440 nt | IncRNA:noe consensus e-value: 7.3e-03

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 4nt Trailing-T: 6nt PSE: 0.90

Bac_try3 | CM028321.1:9994957-9995391 (+) | 435 nt | IncRNA:noe consensus e-value: 5.7e-03

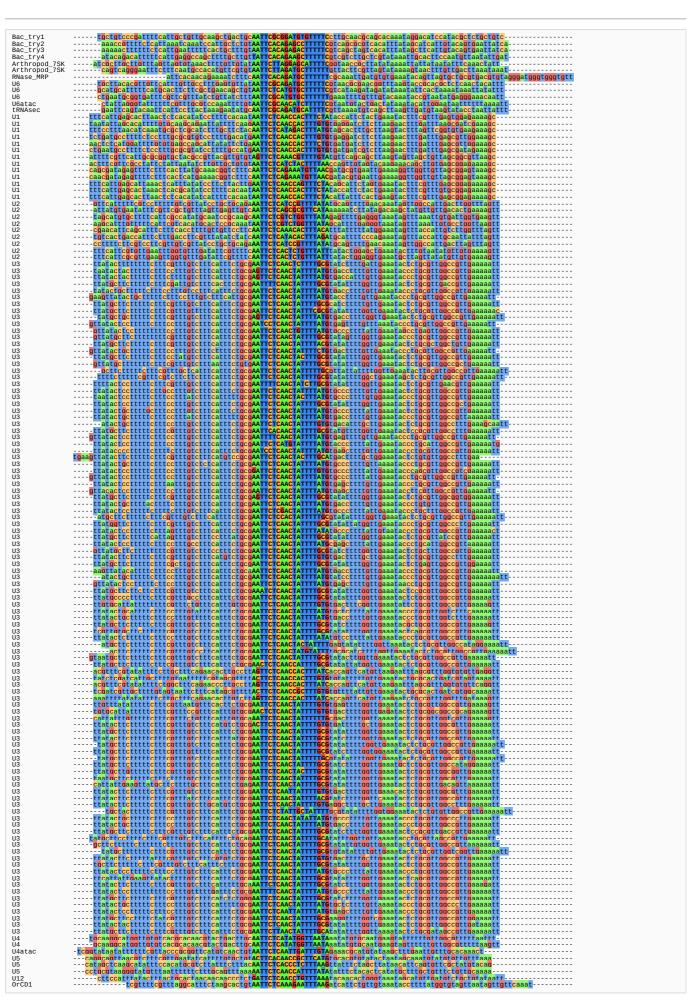
5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 4nt Trailing-T: 6nt PSE: 0.90

Bac_try4 | CM028323.1:56972100-56972394 (-) | 295 nt | IncRNA:noe consensus e-value: 1.6e-08

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 4nt Trailing-T: 6nt PSE: 0.93

	Bac_try1	Bac_try2	Bac_try3	Bac_try4
Bac_try1	-	29	30	21
Bac_try2		-	89	41
Bac_try3			-	42
Bac_try4				-

Pairwise sequence identity matrix (%) showing similarity between IncRNA sequences.



the alignment.