Zeugodacus scutellatus

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

Assembly: GCA_051201265.1_Zscut_ZJ_v1_genomic

Zeu_scu1 | CM118942.1:73478615-73479616 (-) | 1002 nt | IncRNA:noe consensus e-value: 6.9e-16

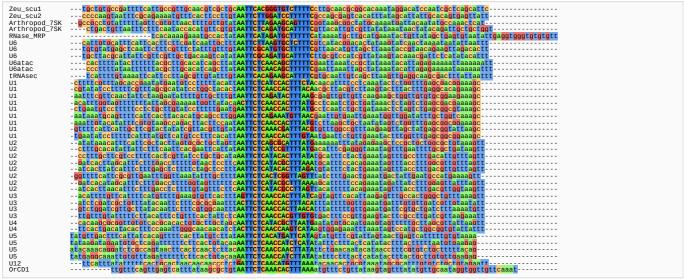
5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 7nt PSE: 0.83

Zeu_scu2 | CM118941.1:8247215-8247667 (-) | 453 nt | IncRNA:noe consensus e-value: 4.7e-04

5' motif: GCGAT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 6nt PSE: 0.88

	Zeu_scu1	Zeu_scu2
Zeu_scu1	-	29
Zeu_scu2		-

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



Upstream promoter alignment showing conserved regions. Sequences include Pol III promoters (noeCR34335, Arthropod_7SK, 7SL, RNase_MRP, RNaseP_nuc, U6, U6atac, tRNAsec, snoRNAMe18SA1806) and Pol II promoters (U1, U2, U3, U4, U4atac, U5, U7, U8, U11, U12, OrCD1). Conserved positions are highlighted in the alignment.