Epicampocera succincta

Taxonomic lineage: Brachycera > Muscomorpha > Oestroidea > Tachinidae > Epicampocera

Assembly: GCA_932526305.1_idEpiSucc1.1_genomic

Epi_suc1 | OW052035.1:64039701-64040211 (-) | 511 nt | IncRNA:noe consensus e-value: 5.1e-02

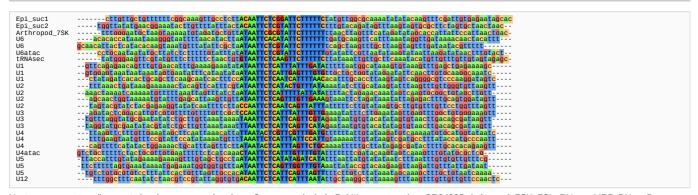
5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 4nt Trailing-T: 6nt PSE: 0.90

Epi_suc2 | OW052034.1:80765472-80765897 (+) | 426 nt | IncRNA:noe consensus e-value: 5.6e+00

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 6nt PSE: 0.94

	Epi_suc1	Epi_suc2
Epi_suc1	-	45
Epi_suc2		-

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.