Dexiosoma caninum

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

Assembly: GCA 963924685.1 idDexCann1.1 genomic

Dex_can1 | OZ004729.1:12634301-12634822 (+) | 522 nt | IncRNA:noe consensus e-value: 9.3e-02

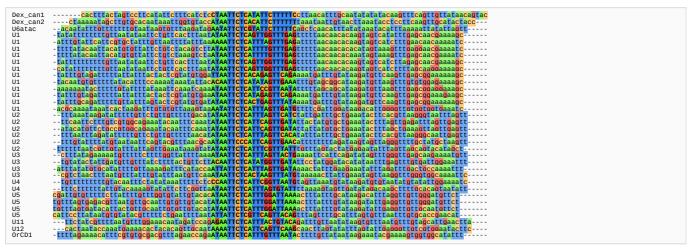
5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 4nt Trailing-T: 7nt

Dex_can2 | OZ004725.1:105539958-105540325 (-) | 368 nt | IncRNA:noe consensus e-value: 7.4e-01

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

	Dex_can1	Dex_can2
Dex_can1	-	43
Dex_can2		-

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.