Drosophila tanythrix

Taxonomic lineage: Brachycera > Muscomorpha > Ephydroidea > Drosophilidae > Drosophila > antopocerus group > diamphidiopoda subgroup

Assembly: GCA_035043275.1_ASM3504327v1_genomic

Dro_tany1 | JAWNNC010000517.1:10868671-10869682 (+) | 1012 nt | IncRNA:noe consensus e-value: 6e-158

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 6nt PSE: 0.84

Dro_tany2 | JAWNNC010000508.1:2733363-2733727 (+) | 365 nt | IncRNA:noe consensus e-value: 3.4e-03

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 4nt Trailing-T: 6nt PSE: 0.99

	Dro_tany1	Dro_tany2
Dro_tany1	-	24
Dro_tany2		-

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.