Drosophila demipolita

Taxonomic lineage: Brachycera > Muscomorpha > Ephydroidea > Drosophilidae > Drosophila

Assembly: GCA 035042405.1 ASM3504240v1 genomic

Dro_dem1 | JAWNLI010000179.1:932261-933249 (-) | 989 nt | IncRNA:noe consensus e-value: 1e-158

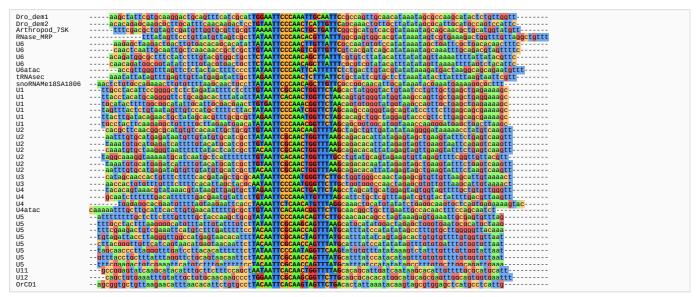
5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 5nt PSE: 0.84

Dro_dem2 | JAWNLI010000486.1:496758-497261 (+) | 504 nt | IncRNA:noe consensus e-value: 2.4e-09

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

	Dro_dem1	Dro_dem2
Dro_dem1	-	33
Dro_dem2		-

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.