Drosophila planitibia

Taxonomic lineage: Brachycera > Muscomorpha > Ephydroidea > Drosophilidae > Drosophila > planitibia group > planitibia subgroup

Assembly: GCA_035043785.1_ASM3504378v1_genomic

Dro_pla1 | JAWNMS010000077.1:1141079-1142106 (+) | 1028 nt | IncRNA:noe consensus e-value: 3e-167

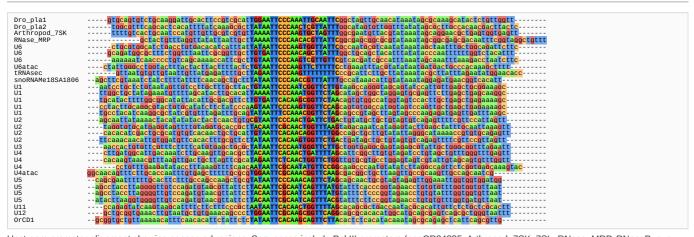
5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 6nt PSE: 0.88

Dro_pla2 | JAWNMS010000041.1:2253147-2253684 (-) | 538 nt | IncRNA:noe consensus e-value: 3.2e-10

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

	Dro_pla1	Dro_pla2
Dro_pla1	-	32
Dro_pla2		-

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.