Drosophila picticornis

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

Assembly: GCA_035043845.1_ASM3504384v1_genomic

Dro_pic1 | JAWNMR010000122.1:13167637-13168672 (-) | 1036 nt | IncRNA:noe consensus e-value: 3e-160

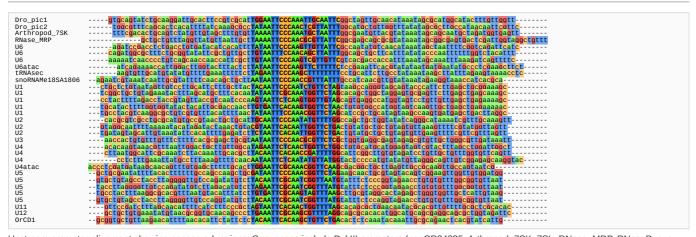
5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 6nt PSE: 0.92

Dro_pic2 | JAWNMR010000492.1:13858549-13859055 (-) | 507 nt | IncRNA:noe consensus e-value: 7.7e-15

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 4nt Trailing-T: 6nt PSE: 1.00

	Dro_pic1	Dro_pic2
Dro_pic1	-	29
Dro_pic2		-

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.