Drosophila anceps

Taxonomic lineage: Brachycera > Muscomorpha > Ephydroidea > Drosophilidae > Drosophila > Drosophila subgenus Drosophila > repleta group > mulleri subgroup

Assembly: GCA_035045945.1_ASM3504594v1_genomic

Dro_anc1 | JAWNON010000186.1:2947947-2948865 (-) | 919 nt | IncRNA:noe consensus e-value: 6e-169

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 7nt PSE: 0.88

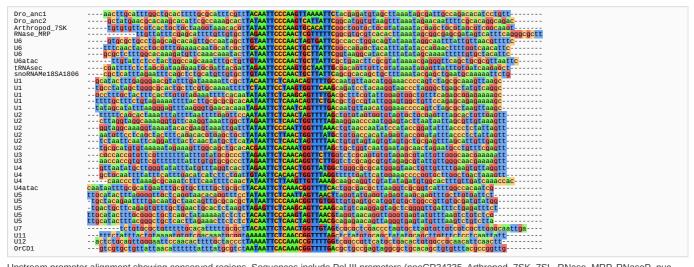
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Dro_anc2 | JAWNON010000004.1:1219053-1219759 (-) | 707 nt | IncRNA:noe consensus e-value: 1.1e+00

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 8nt PSE: 0.94

	Dro_anc1	Dro_anc2
Dro_anc1	-	47
Dro_anc2		-

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.