Drosophila mayaguana

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

Assembly: GCA_037075325.1_ASM3707532v1_genomic

Dro_may1 | JBAMCG010000042.1:5937900-5938820 (-) | 921 nt | IncRNA:noe consensus e-value: 6e-170

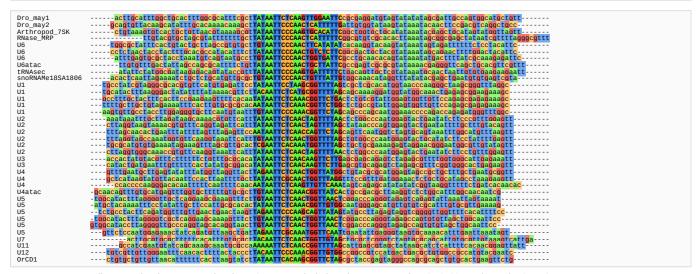
5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 7nt PSE: 0.85

Dro_may2 | JBAMCG010000077.1:1811460-1812124 (+) | 665 nt | IncRNA:noe consensus e-value: NA

5' motif: GCGGT 3' motif: ACCGC Internal max. Poly-T: 3nt Trailing-T: 15nt PSE: 0.98

	Dro_may1	Dro_may2
Dro_may1	-	44
Dro_may2		-

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.