Drosophila melanogaster

Taxonomic lineage: Brachycera > Muscomorpha > Ephydroidea > Drosophilidae > Drosophila > Sophophora > melanogaster group > melanogaster subgroup

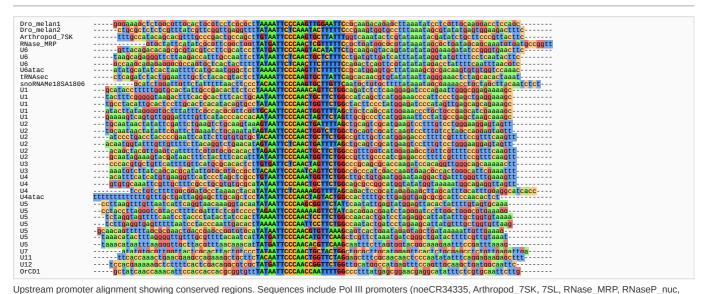
Assembly: GCA_000001215.4_Release_6_plus_ISO1_MT_genomic

Dro_melan2 | AE014298.5:3545750-3546110 (-) | 361 nt | IncRNA:noe consensus e-value: 1.8e-07
5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 4nt Trailing-T: 8nt PSE: 0.95

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	Dro_melan1	Dro_melan2
Dro_melan1	-	23
Dro_melan2		-

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.