Drosophila nigricruria

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

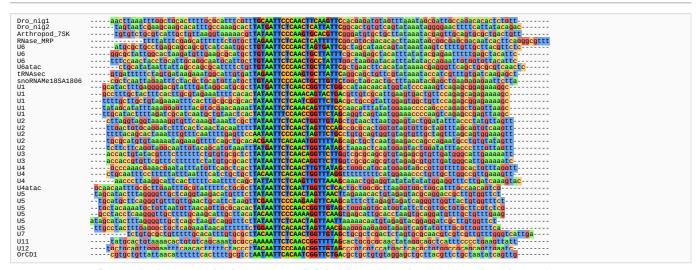
Assembly: GCA_037075265.1_ASM3707526v1_genomic

$Dro_nig2 \mid JBAMCl010000107.1:16458045-16458589 \text{ (-)} \mid 545 \text{ nt} \mid IncRNA: noe consensus e-value: NA}$

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 5nt PSE: 0.93

	Dro_nig1	Dro_nig2
Dro_nig1	-	40
Dro_nig2		-

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.