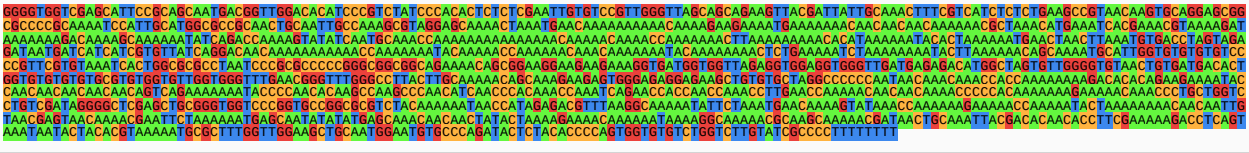
Drosophila prosaltans

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

Assembly: GCA_018151275.1_ASM1815127v1_genomic

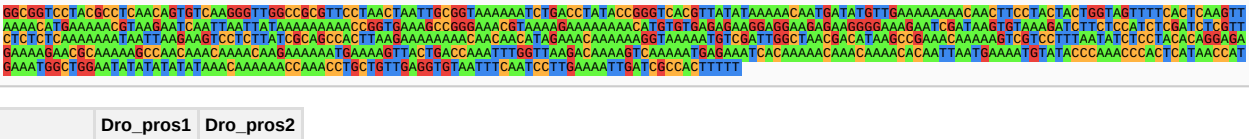
Dro_pros1 | JAECWY010002309.1:2487677-2489134 (+) | 1458 nt | lncRNA:noe consensus e-value: 1e-107

5' motif: GTGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 8nt PSE: 0.91



Dro_pros2 | JAECWY010001814.1:7343758-7344446 (+) | 689 nt | lncRNA:noe consensus e-value: 7.1e+00

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 4nt Trailing-T: 5nt PSE: 0.95



	Dro_pros1	Dro_pros2
Dro_pros1	-	32
Dro_pros2		-

Pairwise sequence identity matrix (%) showing similarity between lncRNA 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.