Drosophila pullipes

Taxonomic lineage: Brachycera > Muscomorpha > Ephydroidea > Drosophilidae > Drosophila > grimshawi group > grimshawi subgroup

Assembly: GCA_035043795.1_ASM3504379v1_genomic

Dro_pul1 | JAWNMT010000649.1:955633-956667 (+) | 1035 nt | IncRNA:noe consensus e-value: 2e-163

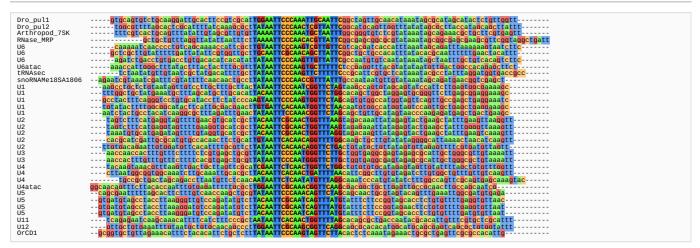
5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 6nt PSE: 0.88

Dro_pul2 | JAWNMT010000185.1:96937-97461 (-) | 525 nt | IncRNA:noe consensus e-value: 5.7e-10

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 6nt PSE: 1.00

	Dro_pul1	Dro_pul2
Dro_pul1	-	31
Dro_pul2		-

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.