Phortica variegata

Taxonomic lineage: Brachycera > Muscomorpha > Ephydroidea > Drosophilidae > Phortica > Phortica subgenus Phortica

Assembly: GCA_001014415.1_ASM101441v1_genomic

Pho_var1 | JXPM01001196.1:1056-2075 (+) | 1020 nt | IncRNA:noe consensus e-value: 9.4e-20

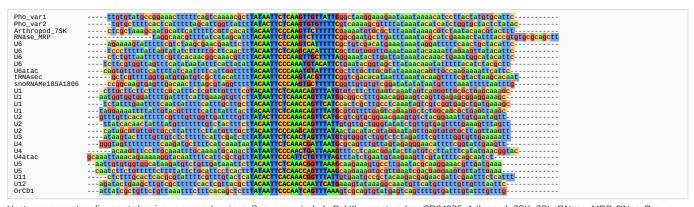
5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 5nt PSE: 0.87

Pho_var2 | JXPM01005486.1:5799-6195 (-) | 397 nt | IncRNA:noe consensus e-value: 3.0e-03

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

	Pho_var1	Pho_var2
Pho_var1	-	29
Pho_var2		-

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.