## Polietes domitor

Taxonomic lineage: Brachycera > Muscomorpha > Muscoidea > Muscidae > Polietes

Assembly: GCA 947397865.1 idPolDomi1.1 genomic

## Pol\_dom1 | OX377621.1:62711463-62711985 (-) | 523 nt | IncRNA:noe consensus e-value: 4.6e-02

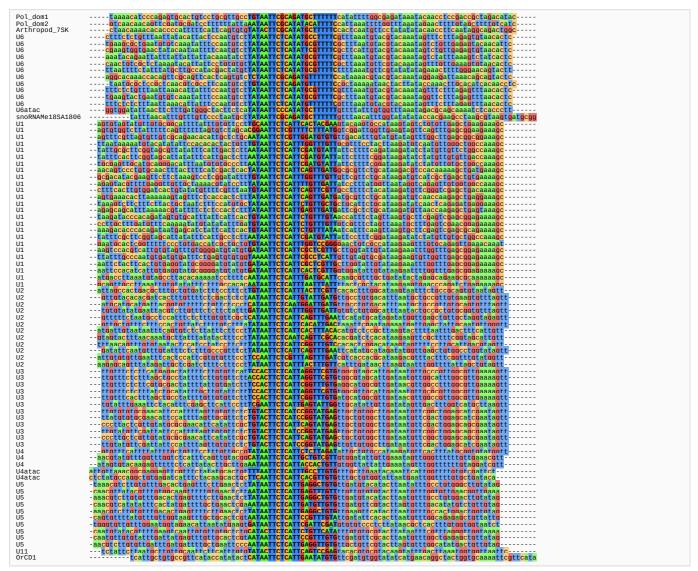
5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 6nt PSE: 0.96

## Pol\_dom2 | OX377618.1:161595879-161596379 (+) | 501 nt | IncRNA:noe consensus e-value: NA

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 4nt Trailing-T: 6nt PSE: 0.87

	Pol_dom1	Pol_dom2
Pol_dom1	-	45
Pol_dom2		-

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.