## Portevinia maculata

Taxonomic lineage: Brachycera > Muscomorpha > Syrphoidea > Syrphidae > Portevinia

Assembly: GCA 949715645.1 idPorMacu1.1 genomic

## Por\_mac1 | OX454526.1:53590523-53590769 (-) | 247 nt | IncRNA:noe consensus e-value: 3.1e-11

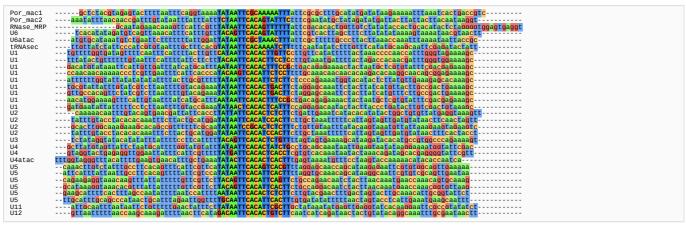
5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 5nt PSE: 0.94

## Por\_mac2 | OX454527.1:97797448-97797629 (-) | 182 nt | IncRNA:noe consensus e-value: 6.1e-03

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

	Por_mac1	Por_mac2
Por_mac1	-	54
Por_mac2		-

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.