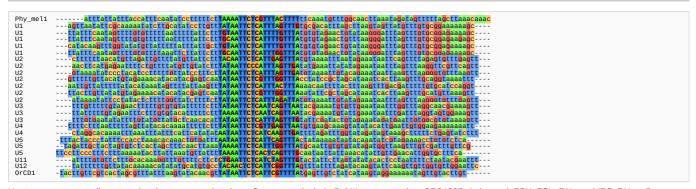
## Phyto melanocephala

Taxonomic lineage: Brachycera > Muscomorpha > Oestroidea > Rhinophoridae > Phyto

Assembly: GCA\_941918925.1\_idPhyMeIn1.1\_genomic

## Phy\_mel1 | OW799233.1:158530180-158530585 (+) | 406 nt | IncRNA:noe consensus e-value: 6.1e-01

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 10nt



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.