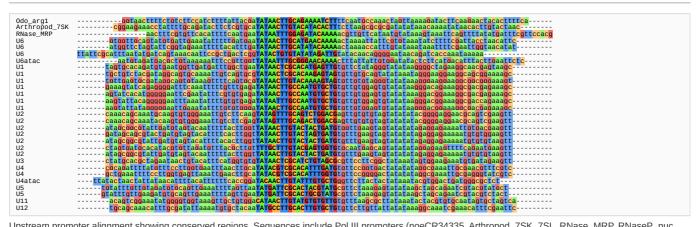
## Odontomyia argentata

Taxonomic lineage: Brachycera > Stratiomyomorpha > Stratiomyidae > Odontomyia

Assembly: GCA\_965615855.1\_idOdoArge1.hap1.1\_genomic

## Odo\_arg1 | OZ278058.1:39078606-39078840 (+) | 235 nt | IncRNA:noe consensus e-value: 6.8e-01

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



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.