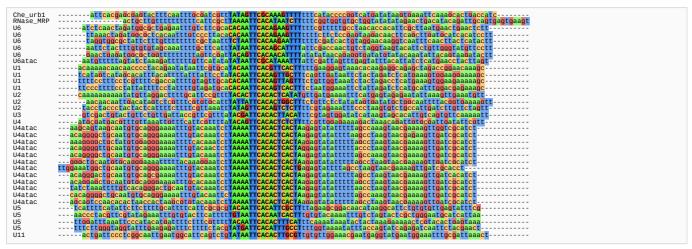
## Cheilosia urbana

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

Assembly: GCA 946477585.1 idCheUrba1.1 genomic

## Che\_urb1 | OX297854.1:24965313-24965580 (-) | 268 nt | IncRNA:noe consensus e-value: 6.5e-11

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 5nt PSE: 0.91



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.