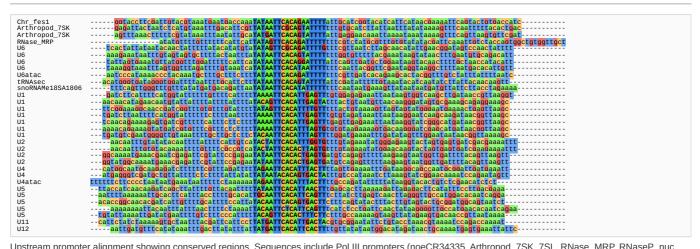
## Chrysotoxum festivum

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

Assembly: GCA 963931855.1 idChrFest1.1 genomic

## Chr\_fes1 | OZ007556.1:49382271-49382559 (+) | 289 nt | IncRNA:noe consensus e-value: 3.9e-11

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 5nt PSE: 1.00



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.