## Eristalinus aeneus

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

Assembly: GCA 955652365.1 idEriAene1.1 genomic

## Eri\_aen1 | OY019132.1:21552471-21552747 (+) | 277 nt | IncRNA:noe consensus e-value: NA

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 4nt Trailing-T: 5nt PSE: 0.89



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.