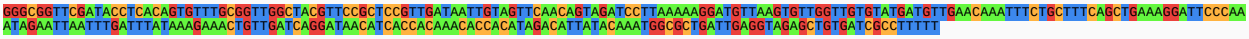
Eumerus sabulonum

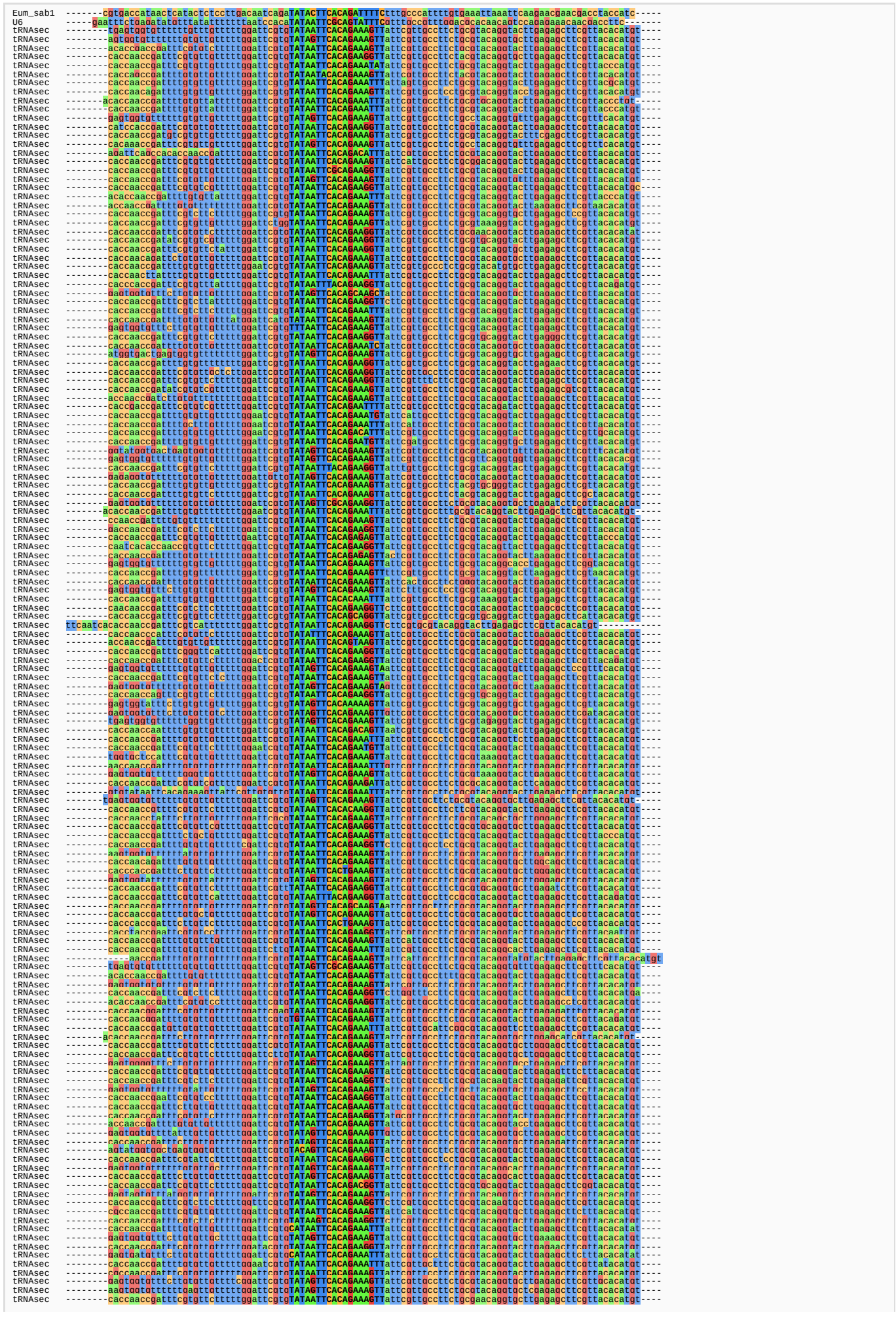
Taxonomic lineage: Brachycera > Muscomorpha > Syrphoidea > Syrphidae > Eumerus

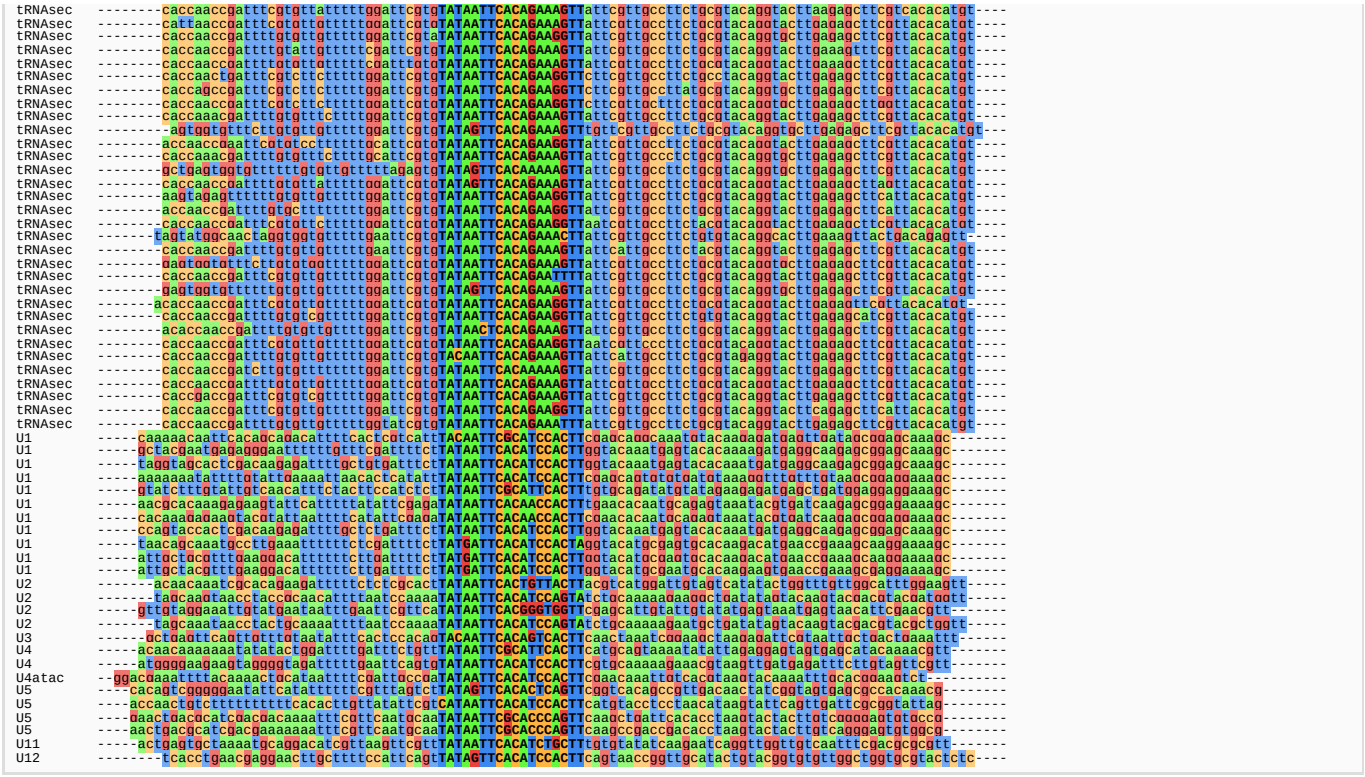
Assembly: GCA_951905685.1_idEumSabu1.1_genomic

Eum_sab1 | OX645064.1:76312061-76312323 (-) | 263 nt | lncRNA:noe consensus e-value: 1.3e-06

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 5nt PSE: 0.80







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.