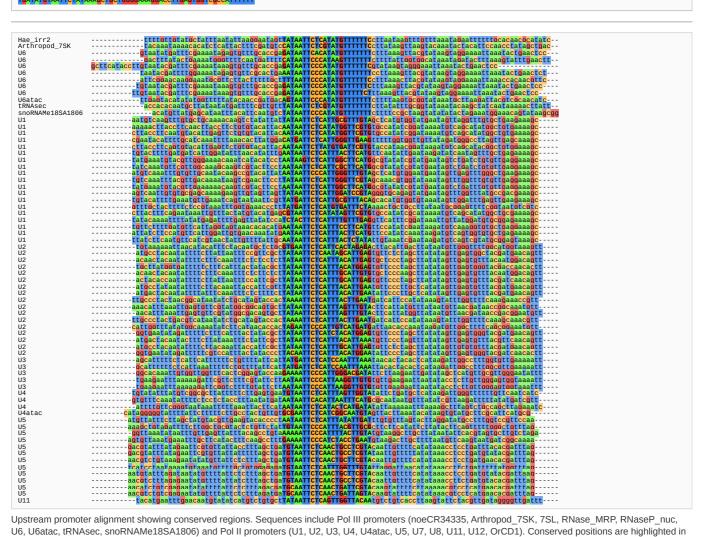
## Haematobia irritans

Taxonomic lineage: Brachycera > Muscomorpha > Muscoidea > Muscidae > Haematobia

Assembly: GCA 050003625.1 ASM5000362v1 genomic

## Hae\_irr2 | CM113865.1:154824607-154824813 (-) | 207 nt | IncRNA:noe consensus e-value: NA

5' motif: GCGGC 3' motif: GTCGC Internal max. Poly-T: 3nt Trailing-T: 6nt PSE: 0.99



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.