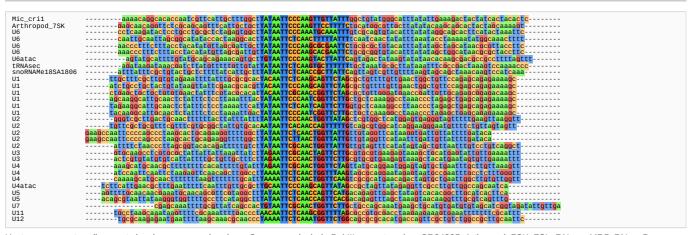
## Microdrosophila cristata

Taxonomic lineage: Brachycera > Muscomorpha > Ephydroidea > Drosophilidae > Microdrosophila

Assembly: GCA\_037075075.1\_ASM3707507v1\_genomic

## Mic\_cri1 | JBAMCQ010000563.1:3032-3619 (-) | 588 nt | IncRNA:noe consensus e-value: 4.0e-03

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 4nt Trailing-T: 5nt PSE: 0.95



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.