## Hydrotaea diabolus

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

Assembly: GCA 963513945.1 idHydDiab1.1 genomic

## Hyd\_dia1 | OY740716.1:180801983-180802469 (-) | 487 nt | IncRNA:noe consensus e-value: 2.6e-03

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 7nt PSE: 0.90



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.