Hydrotaea similis

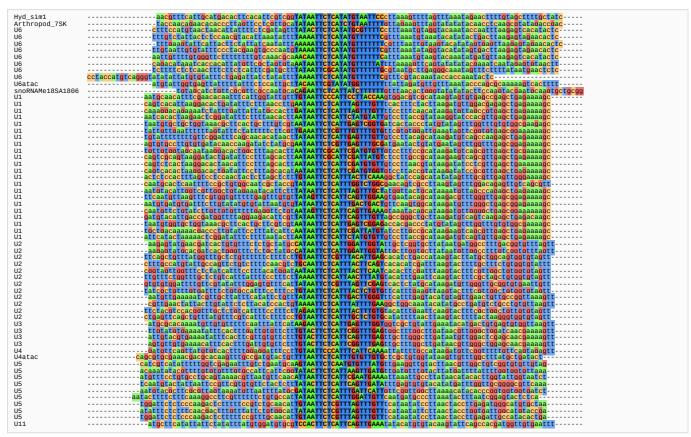
Taxonomic lineage: Brachycera > Muscomorpha > Muscoidea > Muscidae > Hydrotaea

Assembly: GCA 965643655.1 idHydSimi1.hap1.1 genomic

Hyd_sim1 | OZ286020.1:56866099-56866646 (-) | 548 nt | IncRNA:noe consensus e-value: NA

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 9nt PSE: 0.92

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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.