Hydrotaea militaris

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

Assembly: GCA 965614475.1 idHydMili2.hap1.1 genomic

Hyd_mil1 | OZ277508.1:145664652-145665173 (+) | 522 nt | IncRNA:noe consensus e-value: NA

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