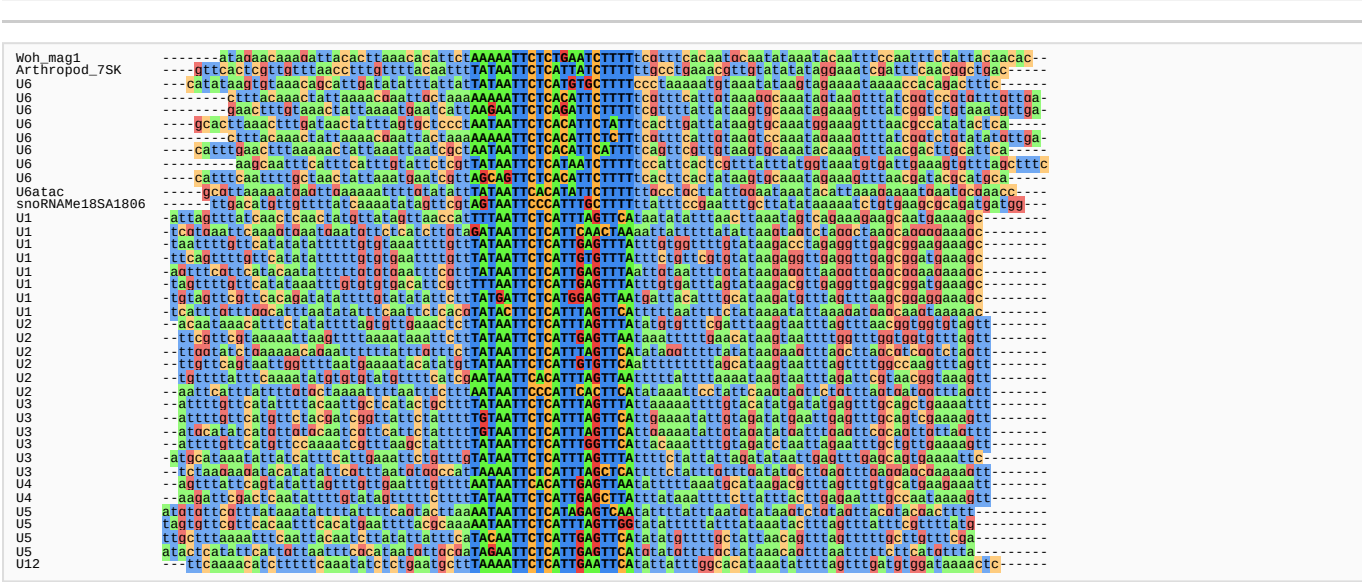
Wohlfahrtia magnifica

Taxonomic lineage: Brachycera > Muscomorpha > Oestroidea > Sarcophagidae > Wohlfahrtia

Assembly: GCA_023159175.1_FiWi_Woma_1.0_genomic

Woh_mag1 | JAKWBJ010000041.1:1402029-1402666 (-) | 638 nt | lncRNA:noe consensus e-value: 3.0e-02

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



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.