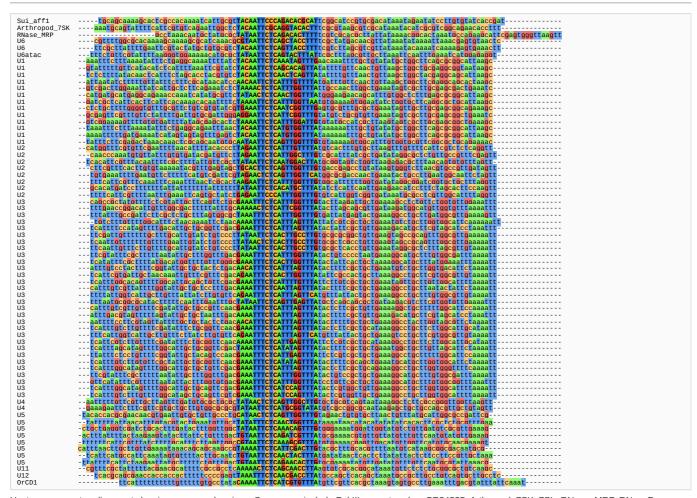
Suillia affinis

Taxonomic lineage: Brachycera > Muscomorpha > Sphaeroceroidea > Heleomyzidae > Suillia

Assembly: GCA 964659615.1 idSuiAffi2.hap1.1 genomic

Sui_aff1 | OZ211076.1:12427980-12428782 (+) | 803 nt | IncRNA:noe consensus e-value: 2.9e-34

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 5nt PSE: 0.88



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.