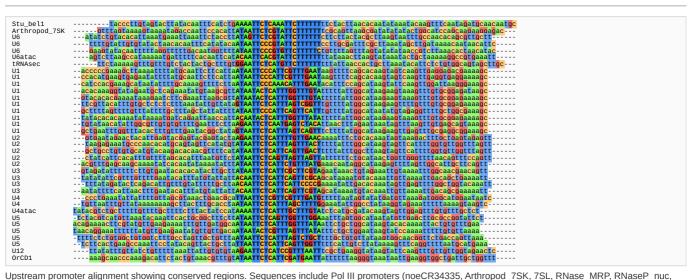
Sturmia bella

Taxonomic lineage: Brachycera > Muscomorpha > Oestroidea > Tachinidae > Sturmia

Assembly: GCA 963662145.1 idStuBell1.1 genomic

Stu_bel1 | OY759169.1:20086317-20086819 (-) | 503 nt | IncRNA:noe consensus e-value: NA

5' motif: GCGGT 3' motif: ACCGC Internal max. Poly-T: 3nt Trailing-T: 6nt 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.