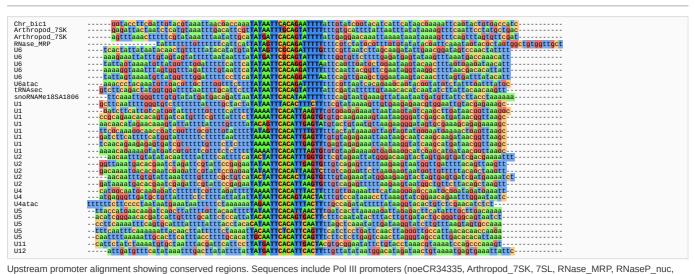
Chrysotoxum bicinctum

Taxonomic lineage: Brachycera > Muscomorpha > Syrphoidea > Syrphidae > Chrysotoxum

Assembly: GCA 911387755.1 idChrBici1.1 genomic

Chr_bic1 | OU426989.1:53307420-53307709 (+) | 290 nt | IncRNA:noe consensus e-value: 2.7e-11

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 5nt PSE: 1.00



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.