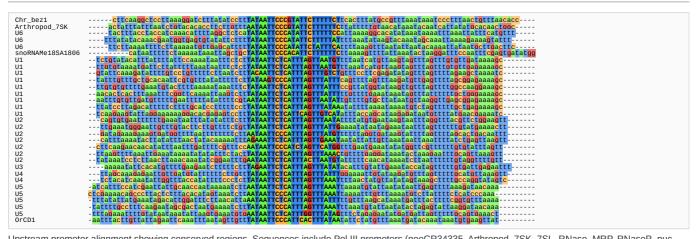
Chrysomya bezziana

Taxonomic lineage: Brachycera > Muscomorpha > Oestroidea > Calliphoridae > Chrysomya

Assembly: GCA_048593825.1_USP_Cbez_1.1_genomic

Chr_bez1 | JBLOGU010000838.1:71772-72247 (-) | 476 nt | IncRNA:noe consensus e-value: 2.0e-01

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 6nt PSE: 0.99



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.