

Taxonomic lineage: Brachycera > Muscomorpha > Syrphoidea > Syrphidae > Baccha
Assembly: GCA_951217065.2 idBacElon1.2 genomic

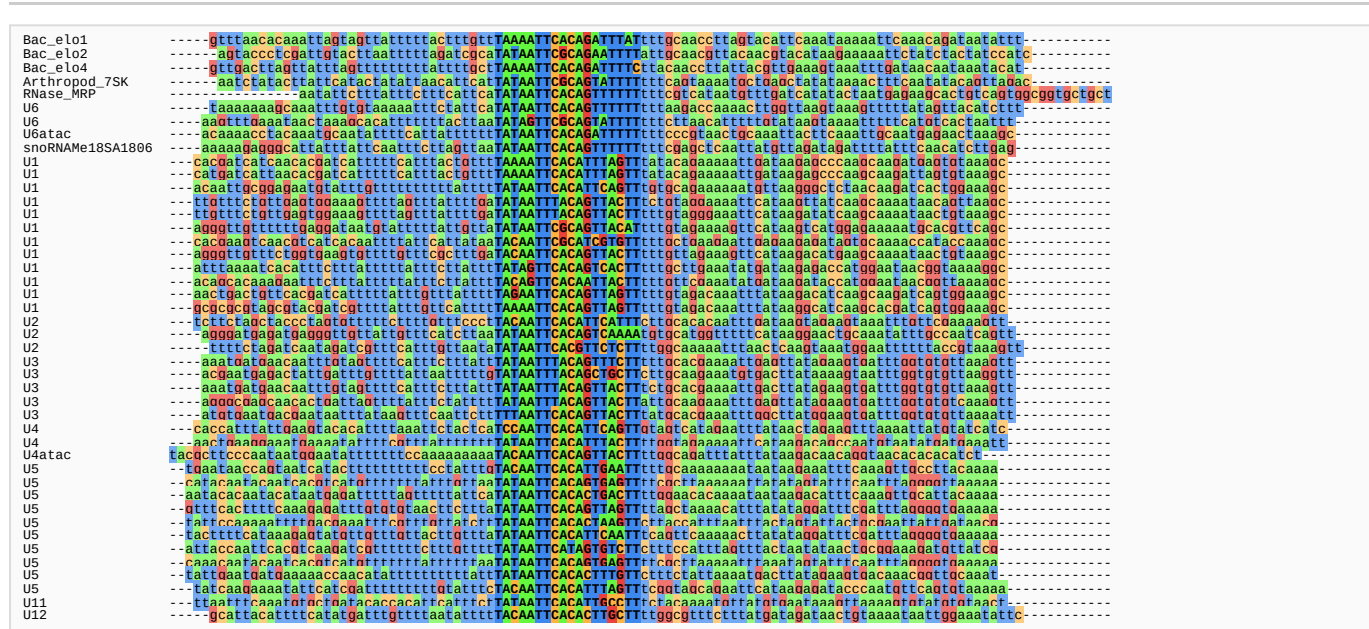
5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 4nt Trailing-T: 5nt PSE: 0.87

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	Bac_elo1	Bac_elo2	Bac_elo4
Bac_elo1	-	46	56
Bac_elo2		-	43
Bac_elo4			-

Pairwise sequence identity matrix (%) showing similarity between lncRNA sequences.



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.