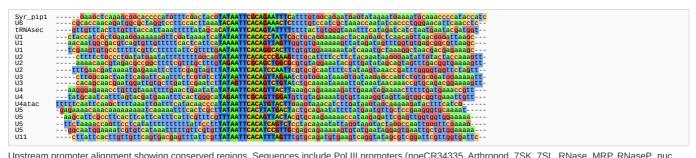
Syritta pipiens

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

Assembly: GCA_905187475.1_idSyrPipi1.1_genomic

Syr_pip1 | LR994571.1:19390290-19390567 (+) | 278 nt | IncRNA:noe consensus e-value: 1.7e-11

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 4nt Trailing-T: 5nt PSE: 0.94



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.