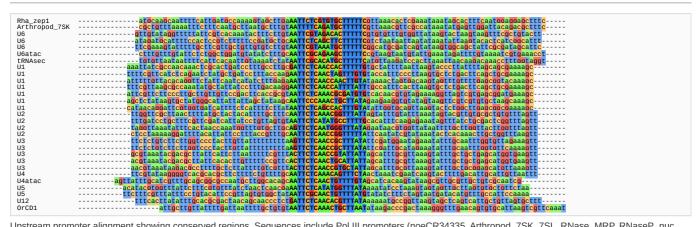
Rhagoletis zephyria

Taxonomic lineage: Brachycera > Muscomorpha > Tephritoidea > Tephritidae > Rhagoletis

Assembly: GCA_001687245.2_Rhagoletis_zephyria_1.1_genomic

Rha_zep1 | LYWK01054191.1:1228-1507 (-) | 280 nt | IncRNA:noe consensus e-value: 1.1e-09

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 4nt Trailing-T: 5nt PSE: 0.85



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.