Eristalis intricaria

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

Assembly: GCA 964034865.1 idEriIntr1.1 genomic

Eri_int1 | OZ035876.1:65450001-65450284 (-) | 284 nt | IncRNA:noe consensus e-value: 9.4e-10

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