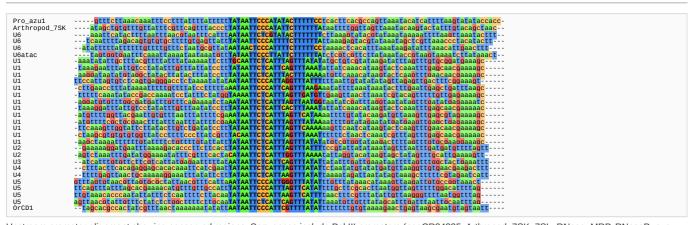
Protocalliphora azurea

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

Assembly: GCA 932274085.1 idProAzur1.1 genomic

Pro_azu1 | OW026519.1:127137211-127137637 (+) | 427 nt | IncRNA:noe consensus e-value: 5.7e-01

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 7nt PSE: 0.98



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.