Volucella bombylans

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

Assembly: GCA 949129095.1 idVolBomb1.1 genomic

Vol_bom1 | OX422143.1:113851371-113851652 (+) | 282 nt | IncRNA:noe consensus e-value: 2.8e-10

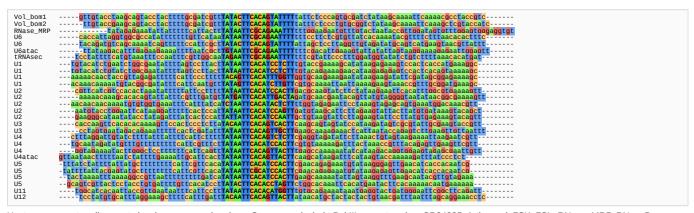
5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 4nt Trailing-T: 5nt PSE: 0.97

Vol_bom2 | OX422143.1:113850727-113850951 (+) | 225 nt | IncRNA:noe consensus e-value: 3.6e-10

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 5nt PSE: 0.97

	Vol_bom1	Vol_bom2
Vol_bom1	-	72
Vol_bom2		-

Pairwise sequence identity matrix (%) showing similarity between IncRNA 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.