

Volucella inflata

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

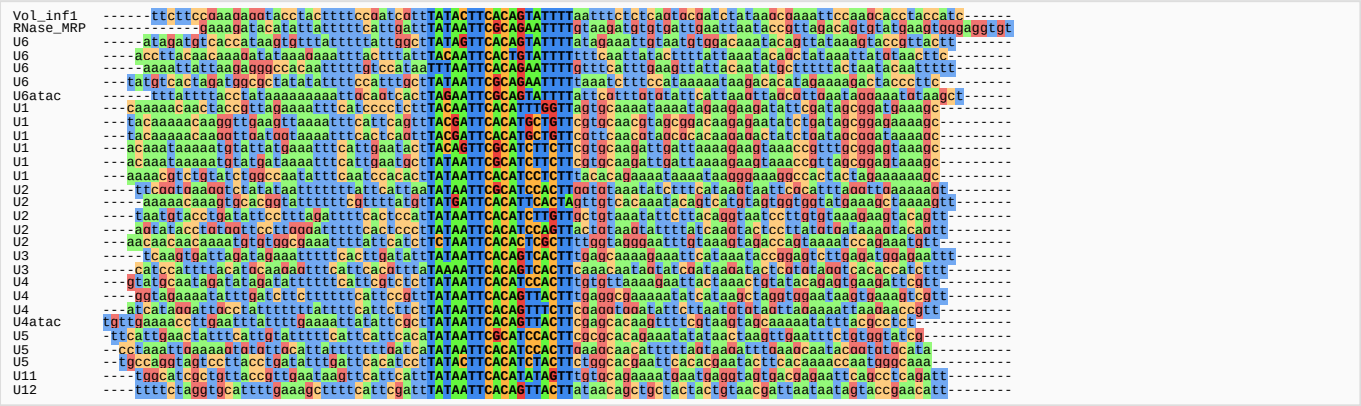
Assembly: GCA_928272305.1_idVollnf1.1_genomic

Vol_inf1 | OV743719.1:47339998-47340281 (-) | 284 nt | lncRNA: noe consensus e-value: 2.3e-10

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 4nt Trailing-T: 5nt PSE: 0.95

GGGCGGTTGATGCTCACAGTGTTCGGTGGTACATTCCGTTCCGCCAATTTGAGCTTCTAAATCGTATGCTTGACATCCATACCTATAATAATAAGATTATCTATATAAAAAAGTGTGTTTGA AAAAGTTTTCACCAAAAAACA

ACGCAAAACAAAAACAATCTTCTCTCTATCA TAAGATAAAATAAAATATAGAATATAAGCAATTCAGTCAATCGAGAAAAAAAAGTTTATCTAAAACGGTTCTGCTGGGGTAGATGCGATCGCCTTTTT



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.