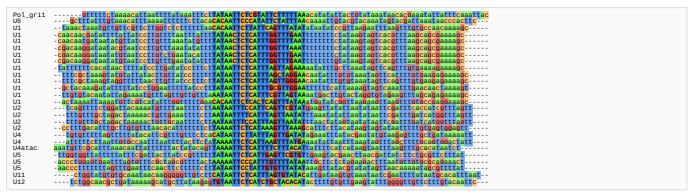
Pollenia griseotomentosa

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

Assembly: GCA_963931875.1_idPolGris2.1_genomic

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Pol_gri1 | OZ007562.1:86915746-86915954 (+) | 209 nt | IncRNA:noe consensus e-value: NA
5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 4nt Trailing-T: 7nt

GCGGTCTTGATCTCTCAAGTGAAACGGTGTTCACATCCCTATAATTGTGAGATTAAAACAACAACAACAACAACAACAACAACAACGAATGAAGTTGAAGTGGCGGAGTACCCCACAGTGAAGTAAAATACACCCATTGAAATTTCAAGCTTTTGGCAGGCGGTGTATAAATTACAAAGTTTTAAGGGGGGAGTACCCCATTGAAATTTCAAGCTTTTGGCAGGCGGTGTATAAATTACAAAGTTTTAAGGGGGGAGTACCACATTGAAAATTACAACGTTTTTTT
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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.