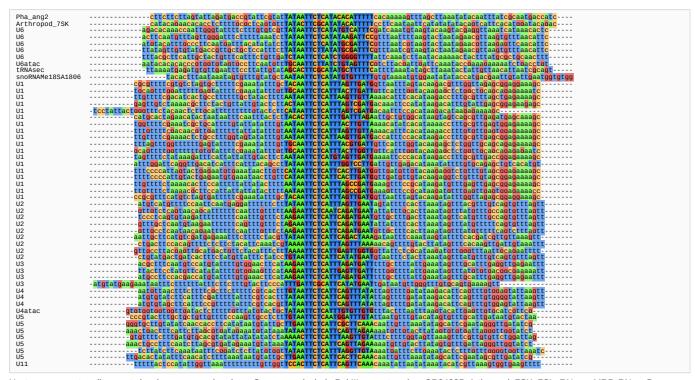
## Phaonia angelicae

Taxonomic lineage: Brachycera > Muscomorpha > Muscoidea > Muscidae > Phaonia

Assembly: GCA\_965111845.1\_idPhaAnge1.hap1.1\_genomic

## Pha\_ang2 | OZ217470.1:160807547-160808089 (+) | 543 nt | IncRNA:noe consensus e-value: NA

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 5nt PSE: 0.93



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.