Amiota albilabris

Taxonomic lineage: Brachycera > Muscomorpha > Ephydroidea > Drosophilidae > Amiota

Assembly: GCA 037040105.1 ASM3704010v1 genomic

Ami_alb1 | JBANCV010000447.1:9402-9772 (+) | 371 nt | IncRNA:noe consensus e-value: 7.7e-07

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



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.