Amiota mariae

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

Assembly: GCA 035041805.1 ASM3504180v1 genomic

Ami_mar1 | JAWNKV010000337.1:8231911-8232899 (+) | 989 nt | IncRNA:noe consensus e-value: 1.1e-11 5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 5nt PSE: 1.00



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.