## Amiota minor

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

Assembly: GCA 037043295.1 ASM3704329v1 genomic

## Ami\_min1 | JBAMAW010002187.1:6128270-6129279 (+) | 1010 nt | IncRNA:noe consensus e-value: 6.0e-12

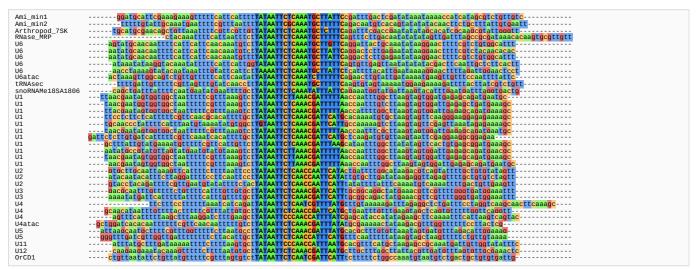
5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 5nt PSE: 1.00

## Ami\_min2 | JBAMAW010002240.1:198619-199009 (-) | 391 nt | IncRNA:noe consensus e-value: 5.3e-02

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

	Ami_min1	Ami_min2
Ami_min1	-	28
Ami_min2		-

Pairwise sequence identity matrix (%) showing similarity between IncRNA sequences.



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.