Amiota communis

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

Assembly: GCA 037043315.1 ASM3704331v1 genomic

Ami_com1 | JBAMAV010001125.1:4235188-4236123 (+) | 936 nt | IncRNA:noe consensus e-value: 4.2e-18

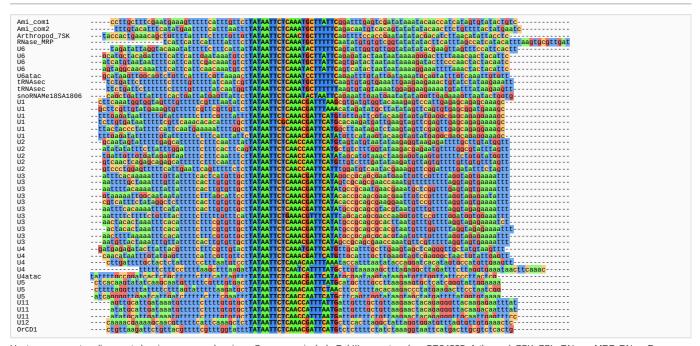
5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 5nt PSE: 1.00

Ami_com2 | JBAMAV010001050.1:1135461-1135833 (-) | 373 nt | IncRNA:noe consensus e-value: 1.1e-06

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 4nt Trailing-T: 5nt PSE: 1.00

	Ami_com1	Ami_com2
Ami_com1	-	26
Ami_com2		-

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.