Musca autumnalis

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

Assembly: GCA 047663935.1 ASM4766393v1 genomic

Mus_aut1 | JBKORA010000001.1:172836349-172836906 (+) | 558 nt | IncRNA:noe consensus e-value: NA

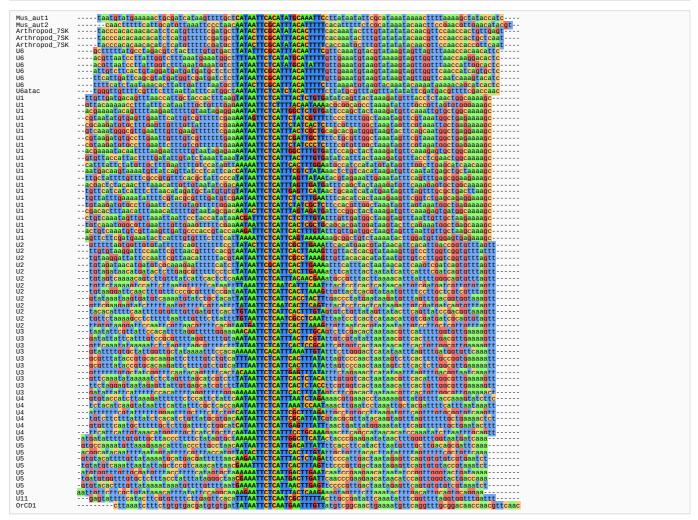
5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 8nt PSE: 0.87

Mus_aut2 | JBKORA010000004.1:73088789-73089266 (-) | 478 nt | IncRNA:noe consensus e-value: 1.6e-02

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 6nt PSE: 0.98

	Mus_aut1	Mus_aut2
Mus_aut1	-	49
Mus_aut2		-

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.