Xyphosia miliaria

Taxonomic lineage: Brachycera > Muscomorpha > Tephritoidea > Tephritidae > Xyphosia

Assembly: GCA_965200055.1_idXypMili3.hap1.1_genomic

Xyp_mil1 | OZ241499.1:7783468-7784339 (+) | 872 nt | IncRNA:noe consensus e-value: 2.9e-19

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 7nt PSE: 0.95

Xyp_mil2 | OZ241498.1:9699206-9699399 (-) | 194 nt | IncRNA:noe consensus e-value: 4.1e-03

5' motif: GCGAT 3' motif: ATCGC Internal max. Poly-T: 4nt Trailing-T: 5nt PSE: 0.88

	Xyp_mil1	Xyp_mil2
Xyp_mil1	-	15
Xyp_mil2		-

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.