Stomorhina lunata

Taxonomic lineage: Brachycera > Muscomorpha > Oestroidea > Rhiniidae > Stomorhina

Assembly: GCA 933228675.1 idStoLuna1.1 genomic

Sto_lun1 | OW121742.1:46158189-46158396 (-) | 208 nt | IncRNA:noe consensus e-value: NA

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 7nt PSE: 1.00

Sto_lun2 | OW121742.1:46162200-46162405 (-) | 206 nt | IncRNA:noe consensus e-value: NA

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 3nt Trailing-T: 6nt PSE: 1.00

	Sto_lun1	Sto_lun2
Sto_lun1	-	97
Sto_lun2		-

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.