Sarcophaga carnaria

Taxonomic lineage: Brachycera > Muscomorpha > Oestroidea > Sarcophagidae > Sarcophaga > Sarcophaga subgenus Sarcophaga

Assembly: GCA 958299815.1 idSarCarn1.1 genomic

Sar_car1 | OY283150.1:61821353-61821870 (-) | 518 nt | IncRNA:noe consensus e-value: 3.1e+00

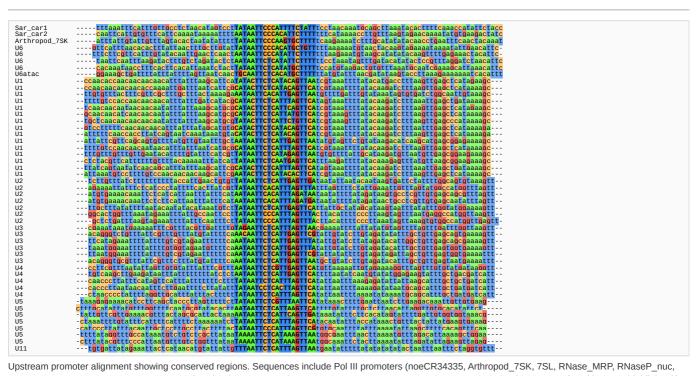
5' motif: GCGGT 3' motif: ACCGC Internal max. Poly-T: 3nt Trailing-T: 10nt PSE: 0.90

Sar_car2 | OY283151.1:8978686-8978910 (+) | 225 nt | IncRNA:noe consensus e-value: 1.4e-05

5' motif: GCGGT 3' motif: ACCGC Internal max. Poly-T: 4nt Trailing-T: 8nt PSE: 0.99

	Sar_car1	Sar_car2
Sar_car1	-	35
Sar_car2		-

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.