## Campiglossa producta

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

Assembly: GCA 964058745.1 idCamProd1.hap1.1 genomic

## Cam\_pro1 | OZ060071.1:6951345-6952188 (+) | 844 nt | IncRNA:noe consensus e-value: 3.0e-14

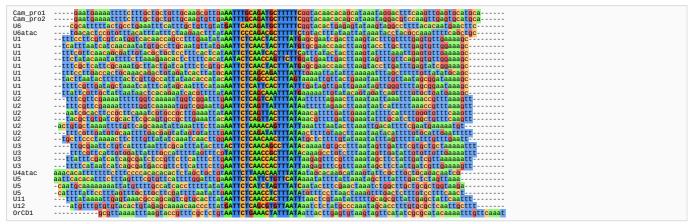
5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 4nt Trailing-T: 6nt PSE: 0.90

## Cam\_pro2 | CAXKVH010000746.1:62908-63746 (-) | 839 nt | IncRNA:noe consensus e-value: 1.1e-14

5' motif: GCGGT 3' motif: ATCGC Internal max. Poly-T: 4nt Trailing-T: 6nt PSE: 0.90

	Cam_pro1	Cam_pro2
Cam_pro1	-	99
Cam_pro2		-

Pairwise sequence identity matrix (%) showing similarity between lncRNA 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.