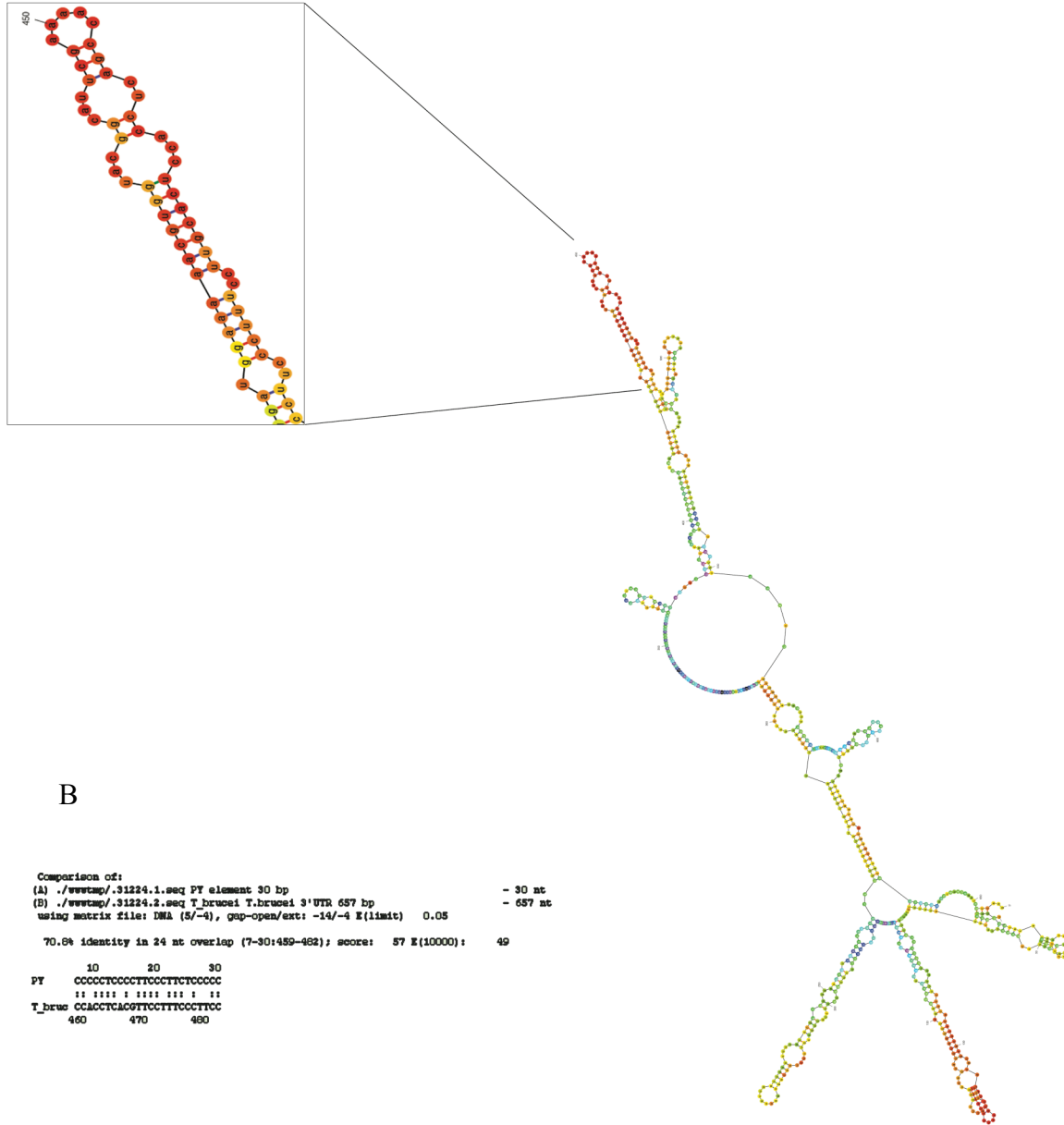


A



B

Supplementary Figure 3.

A. Structure prediction of the 3' UTR from Hsp83 of *Trypanosoma brucei*. The prediction was performed with the UNAFold V3.4-1 using Mfold utilities. The highly defined arm (red nucleotides) is enlarged.

B. A PPT with partial homology to the *Leishmania* regulatory sequence is shown.