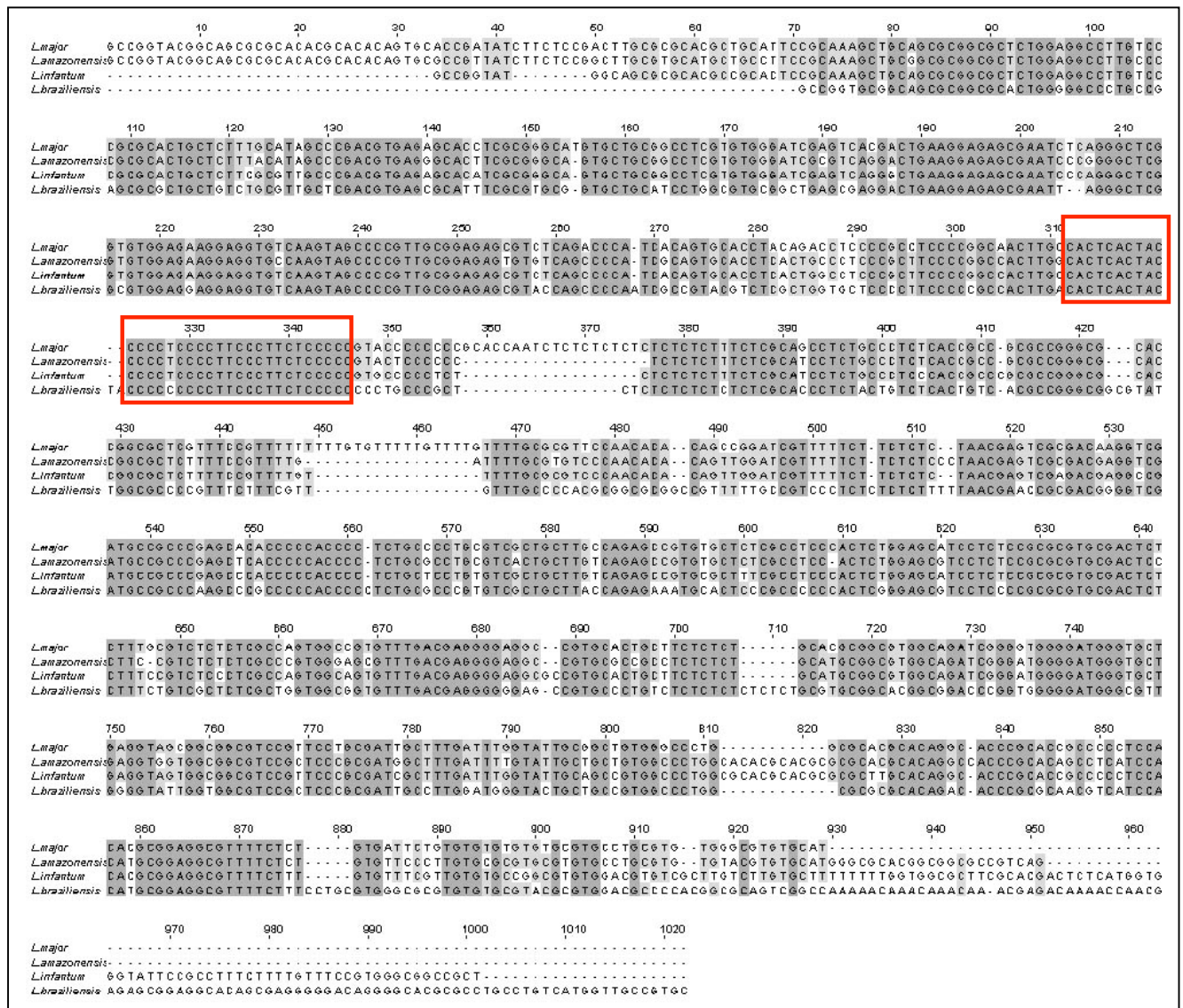




Supplementary Figure 2. Sequence alignment of the 3' UTRs of Hsp83 from four different *Leishmania* species. 3' UTRs from *L. amazonensis* (M92925.1); *L. Major* (S82021.1); *L. infantum* (Contig LinJ33_20070420 Location 113201-114101) and *L. braziliensis* (contig LbrM33_V2_October Location 115218-116118). The alignment was performed with ClustalW. Identical nucleotides are indicated in black. Similar nucleotides are indicated in gray boxes.