Figure S1. Secondary structure of IFN-ε 5'UTR generated using publicly-available prediction tools, and alignment of the region among several primates

A. KineFold (http://kinefold.curie.fr/)

B. RNAfold, MFE (http://rna.tbi.univie.ac.at)

C. RNAfold, Centroid (http://rna.tbi.univie.ac.at)