

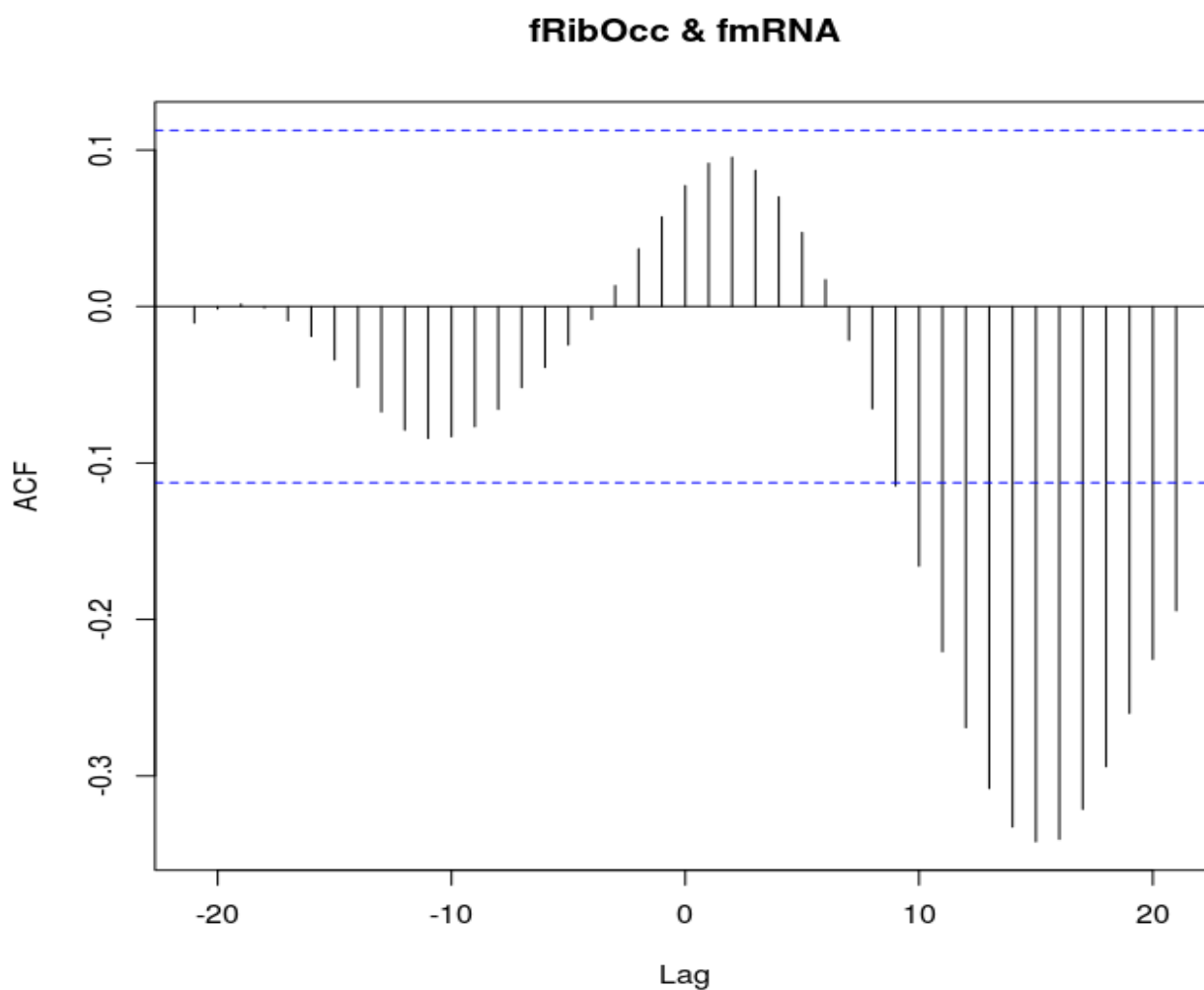
Amino-acid independent protein functional features coded in the mRNA

Daniel López & Florencio Pazos

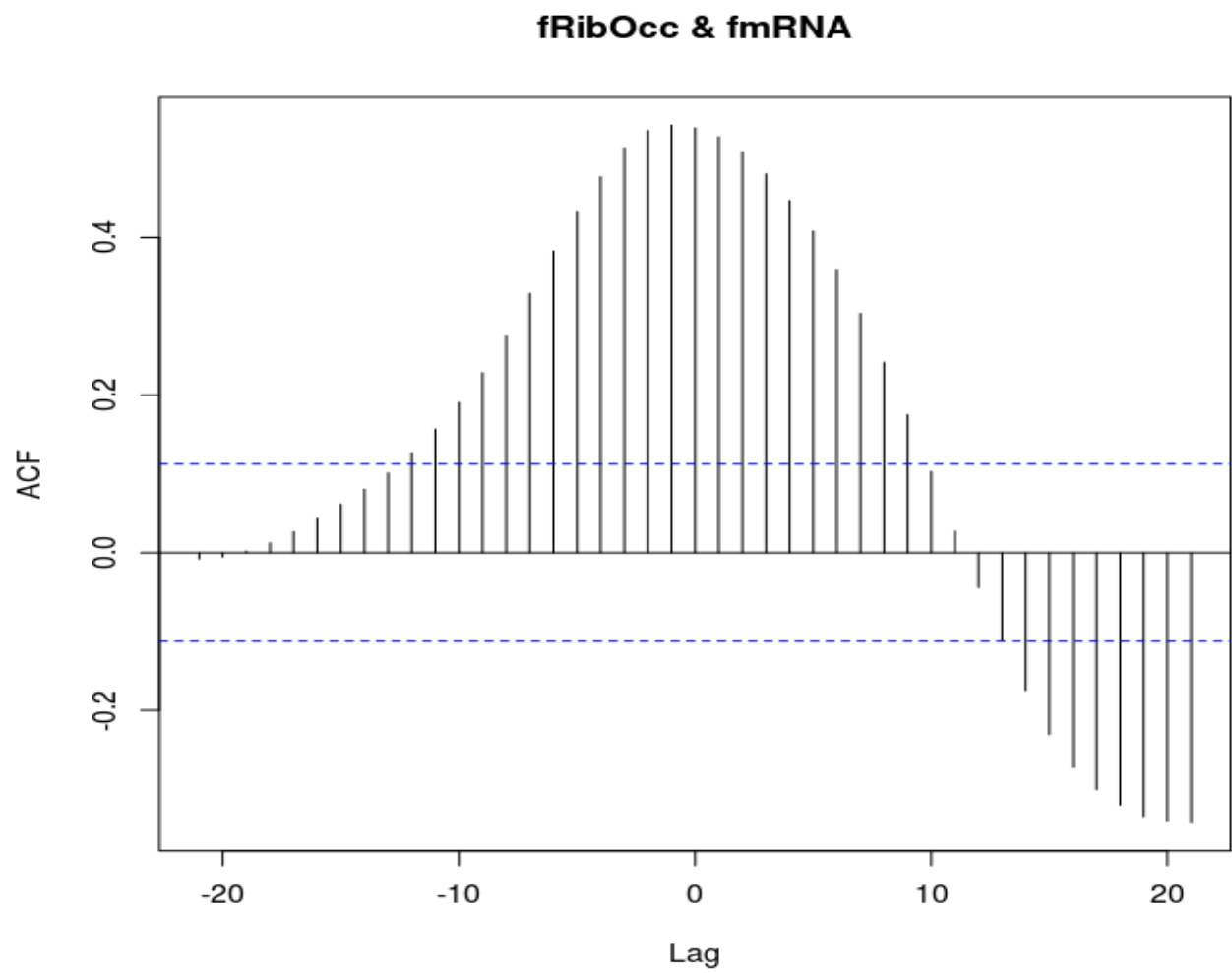
Supplementary File 3

Cross-correlation between the mRNA secondary structure and ribosome occupancy patterns for the protein functional features discussed in Figure 4. The plots show the correlation obtained (Y-axis) when one of the patterns is displaced a given number of positions (X-axis) respect to the other.

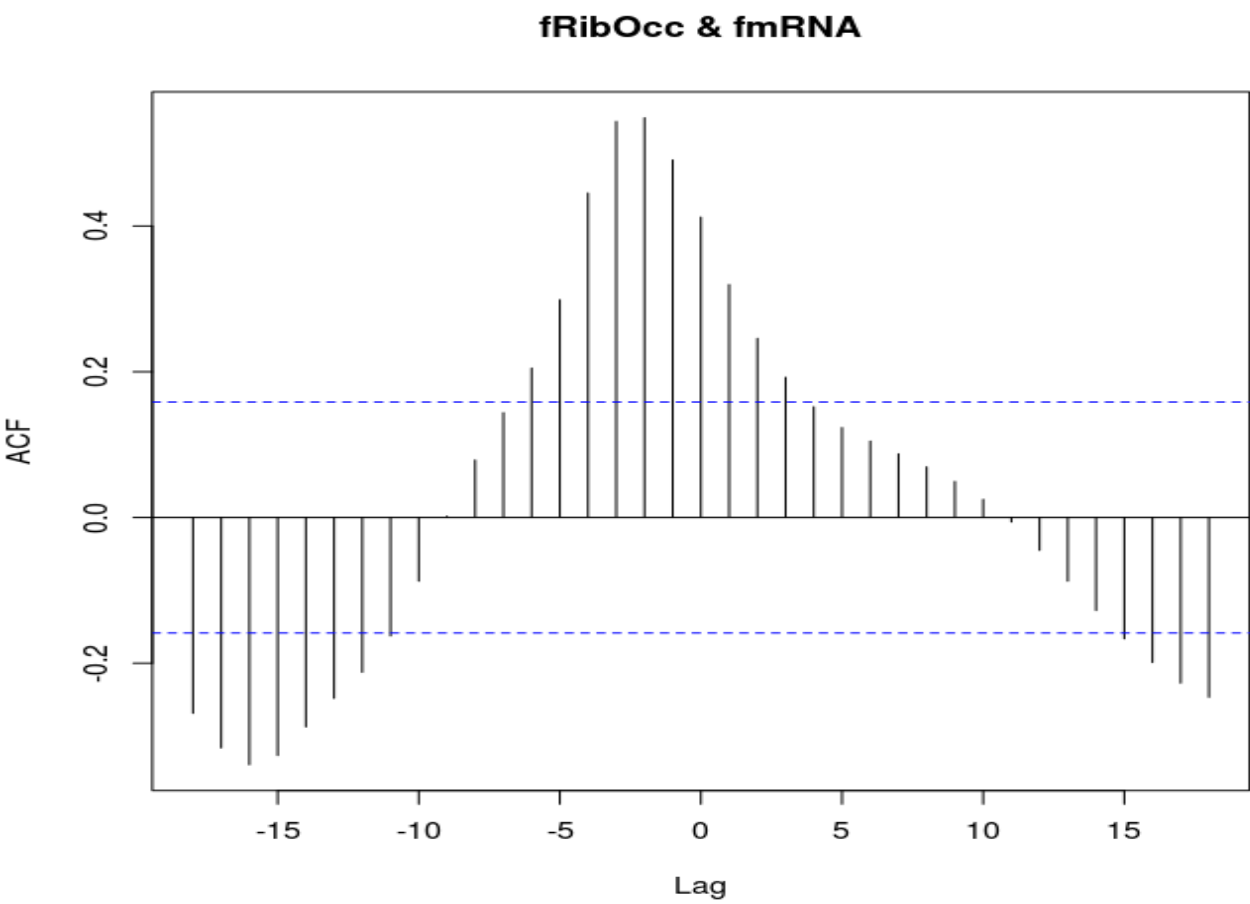
Domain



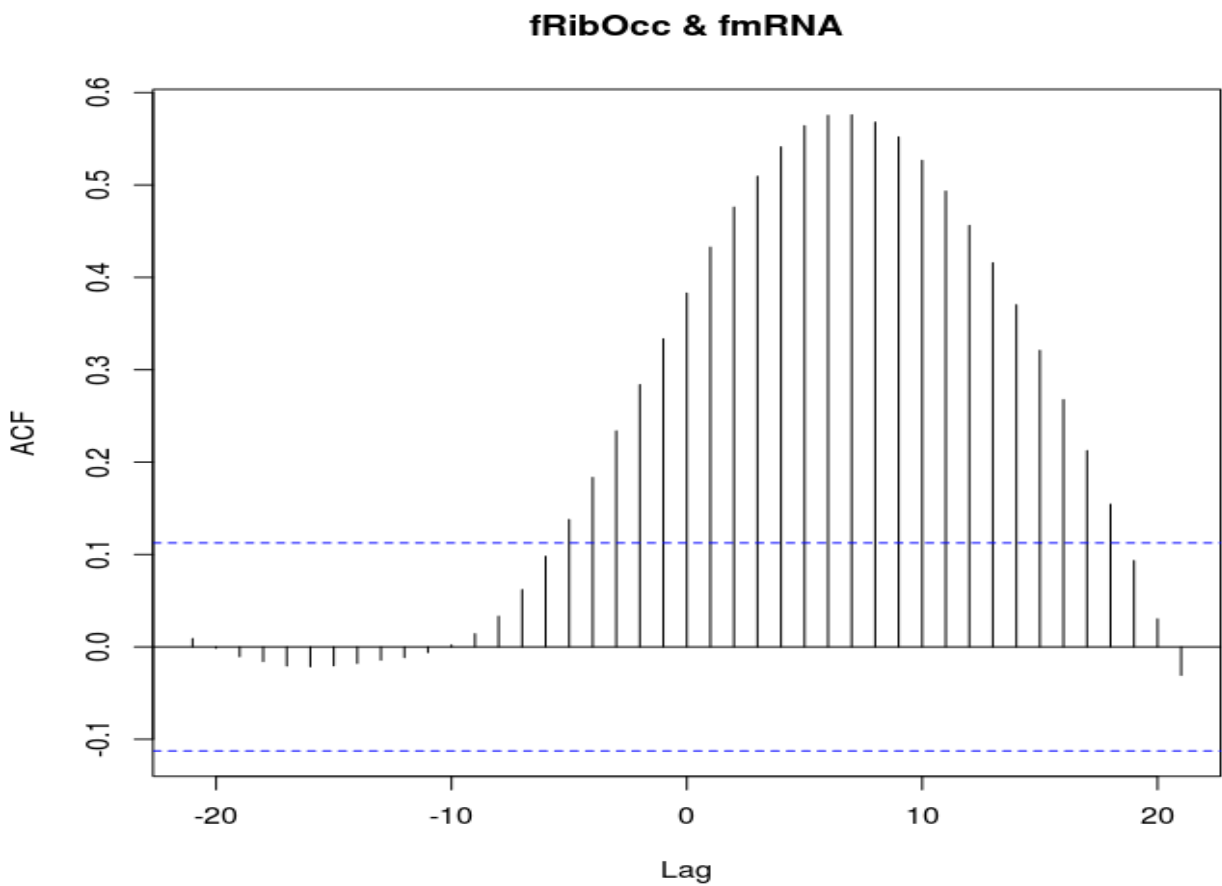
helix



initiator-metionine



nucleotide-phosphate-binding



Transmembrane region

