

11,000 unstable
eRNA 5' ends

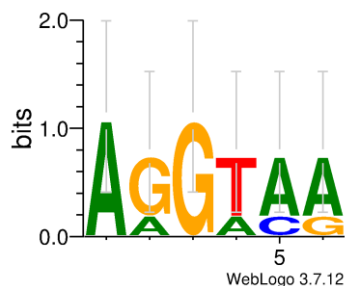
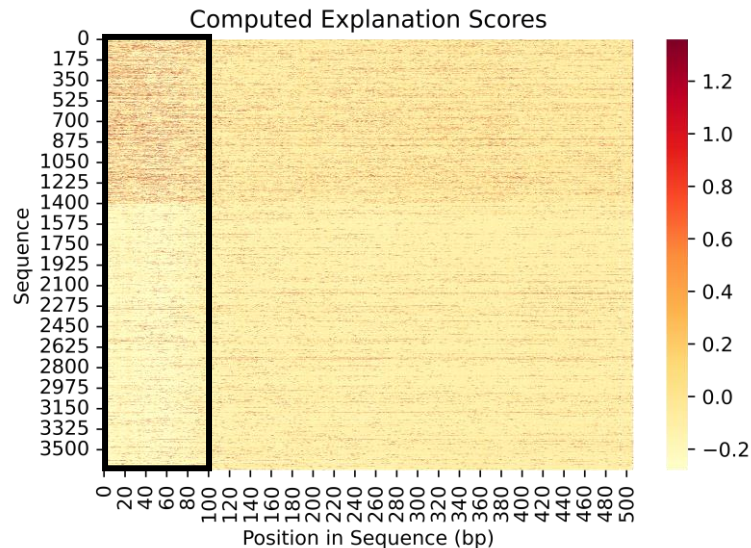
...

15,000 mRNA 5'
ends



DNABERT

Relevancy calculated
from attention scores



Cluster0:
AGGTAG
AGGTAA
AGGTCA
AAGAAA

...

Cluster17:



Clusters and aligns k-mers

Downstream analyses: motif
enrichment, positional biases, etc ...