

**mRNA/EST Translation**

- 1) Visit the following site:  
<https://www.ncbi.nlm.nih.gov/dbEST/>  
dbEST retired  
<https://ncbiinsights.ncbi.nlm.nih.gov/2019/07/25/est-and-gss-databases-now-retired/>
- 2) Read the paper – OrfPredictor at  
[http://proteomics.yzu.edu/courses/BIOL4800\\_6900/ExtraReadings/OrfPredictor.pdf](http://proteomics.yzu.edu/courses/BIOL4800_6900/ExtraReadings/OrfPredictor.pdf)

Go to the following site to practice using your own mRNA sequences or data downloaded from dbEST for any species - <http://proteomics.yzu.edu/tools/OrfPredictor.html>

- 3) Learn to use ORF-Finder at  
<https://www.ncbi.nlm.nih.gov/orffinder/>  
**This tool is designed for a researcher to do annotation before submitting mRNA sequences to GenBank.**

**RNA**

```
>NC_000014.9:c20609407-20609336 Homo sapiens chromosome 14, GRCh38.p13
GGCTCGTTGGTCTAGGGGTATGATTCTCGCTTAGGGTGCGAGAGGTCCCGGGTTCAAATCCCG
GACGAGCCC
>NC_000017.11:c8120395-8120314 Homo sapiens chromosome 17, GRCh38.p13
GGTAGCGTGGCCGAGCGGTCTAAGGCGCTGGATTTAGGCTCCAGTCTCTTCGGAGGCGTGGGT
TCGAATCCCACCGCTGCCA
```

**Using above two sequences to test if they are tRNAs by tRNAScan:**

<http://lowelab.ucsc.edu/tRNAScan-SE/>

Visit the following sites:

GEO: <http://www.ncbi.nlm.nih.gov/geo/>

RNA Central: <https://rnacentral.org/>

Extra-reading:

[http://proteomics.yzu.edu/courses/BIOL4800\\_6900/ExtraReadings/RNA\\_central2019.pdf](http://proteomics.yzu.edu/courses/BIOL4800_6900/ExtraReadings/RNA_central2019.pdf)