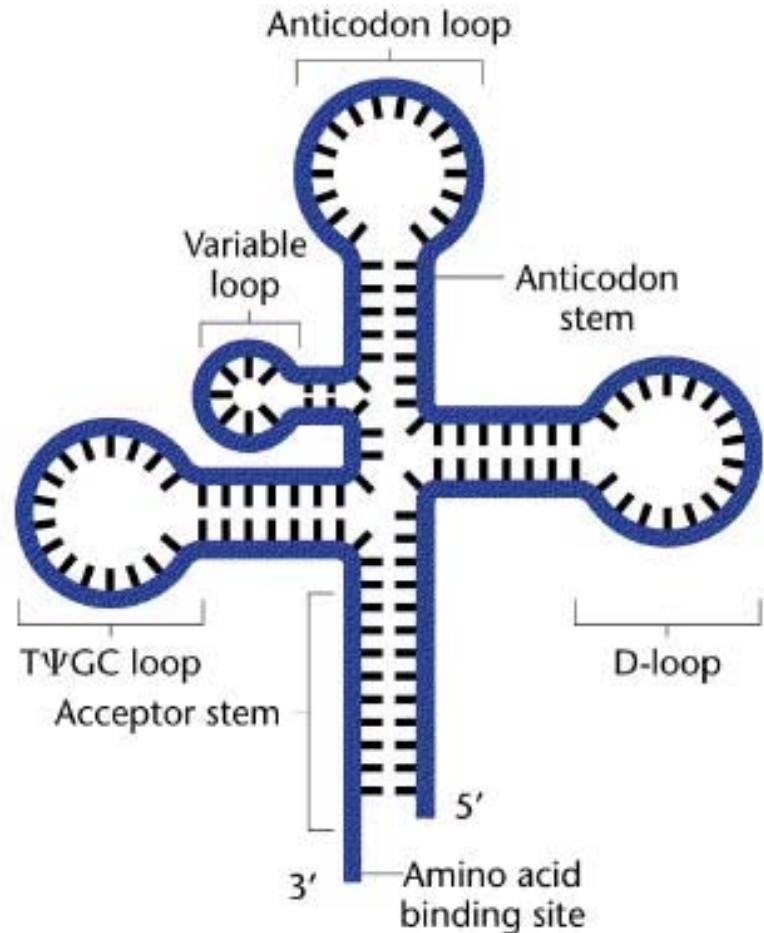


Translation

From RNA to Protein

Nucleic Acid to Protein



- How does the information in codons of mRNA get translated into amino acids in polypeptides?
- Through adapter molecules: [tRNA](#)
- tRNA has **anticodon** that base pairs with the **codon** in mRNA and carries an amino acid corresponding to that codon.

Transfer RNAs (tRNAs)

Any cell contains different types of tRNA molecules sufficient to incorporate all 20 amino acids into protein.

Some tRNAs can recognise more than one codon.

Short, about 80 nucleotides in length.

Complex secondary and tertiary structures.

Folding follows rules – complementary base pairs

- Stems and loops

- Contain non-standard base pairs

- Contain modified bases

Post-transcriptional processing of tRNA

Removal of introns

Separation of multiple tRNAs

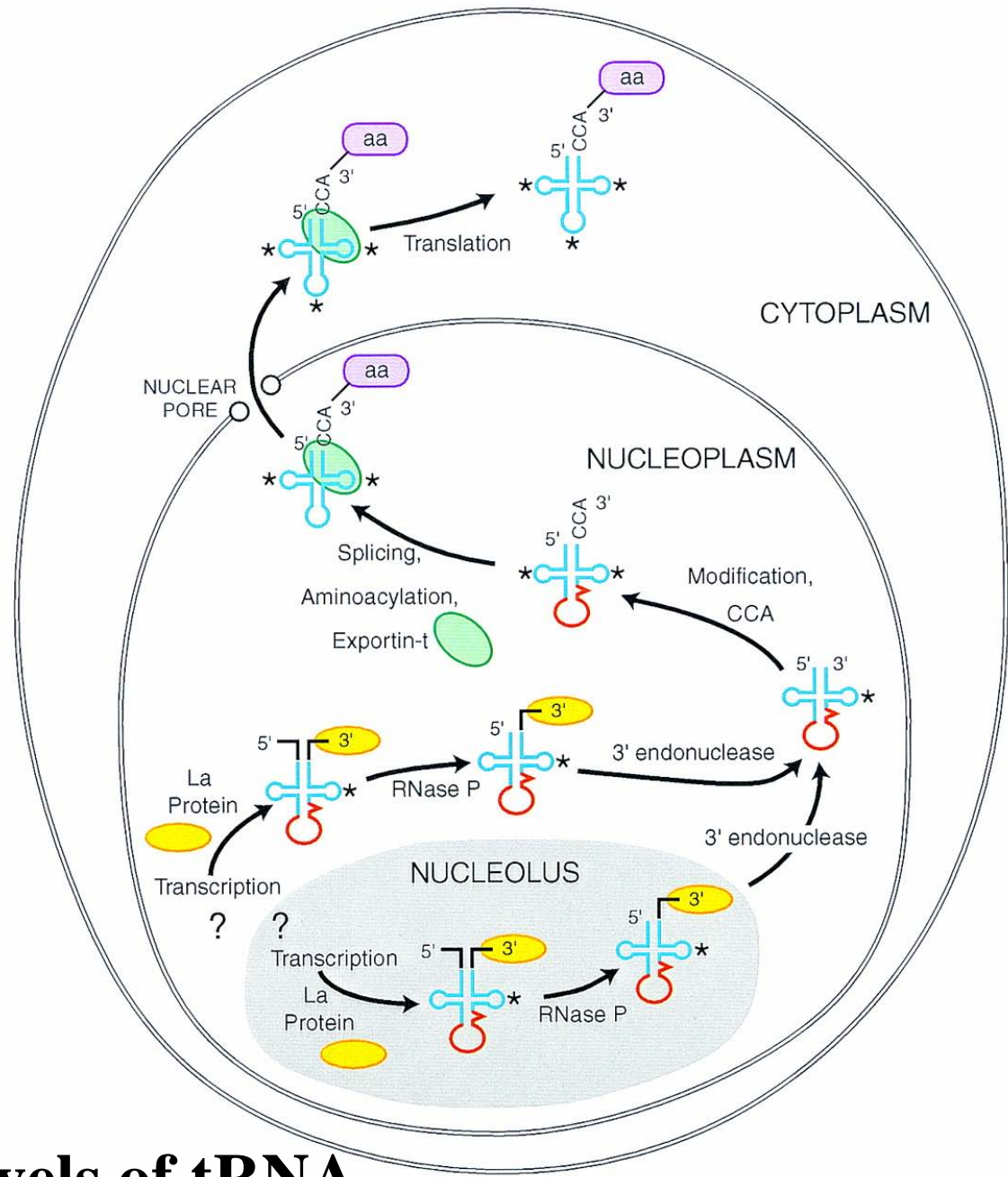
Processing of the 5' (RNaseP), addition of 3' CCA

Base modification

- Methylation

- Deamination

- Reduction



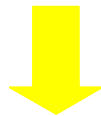
The trials and travels of tRNA

What is the correspondence between the mRNA nucleotides and the amino acids of the protein?

Proteins are formed from 20 amino acids in humans.

Codons of one nucleotide:

A
G
C
U



**Can only encode
4 amino acids**

Codons of two nucleotides:

AA GA CA UA
AG GG CG UG
AC GC CC UC
AU GU CU UU



**Can only encode
16 amino acids**

Note that 3rd Base Position is Variable

		Second position					
		U	C	A	G		
First position (5'-end)	U	UUU <i>phe</i> UUC	UCU UCC <i>ser</i> UCA UUG	UAU <i>tyr</i> UAC UAA <i>Stop</i> UAG <i>Stop</i>	UGU <i>cys</i> UGC UGA <i>Stop</i> UGG <i>trp</i>	Third position (3'-end)	U
	C	CUU CUC <i>leu</i> CUA CUG	CCU CCC <i>pro</i> CCA CCG	CAU <i>his</i> CAC CAA <i>gln</i> CAG	CGU CGC <i>arg</i> CGA CGG		U
	A	AUU AUC <i>ile</i> AUA AUG <i>met</i>	ACU ACC <i>thr</i> ACA ACG	AAU <i>asn</i> AAC AAA <i>lys</i> AAG	AGU <i>ser</i> AGC AGA <i>arg</i> AGG		C
	G	GUU GUC <i>val</i> GUA GUG	GCU GCC <i>ala</i> GCA GCG	GAU <i>asp</i> GAC GAA <i>glu</i> GAG	GGU GGC <i>gly</i> GGA GGG		A
						G	

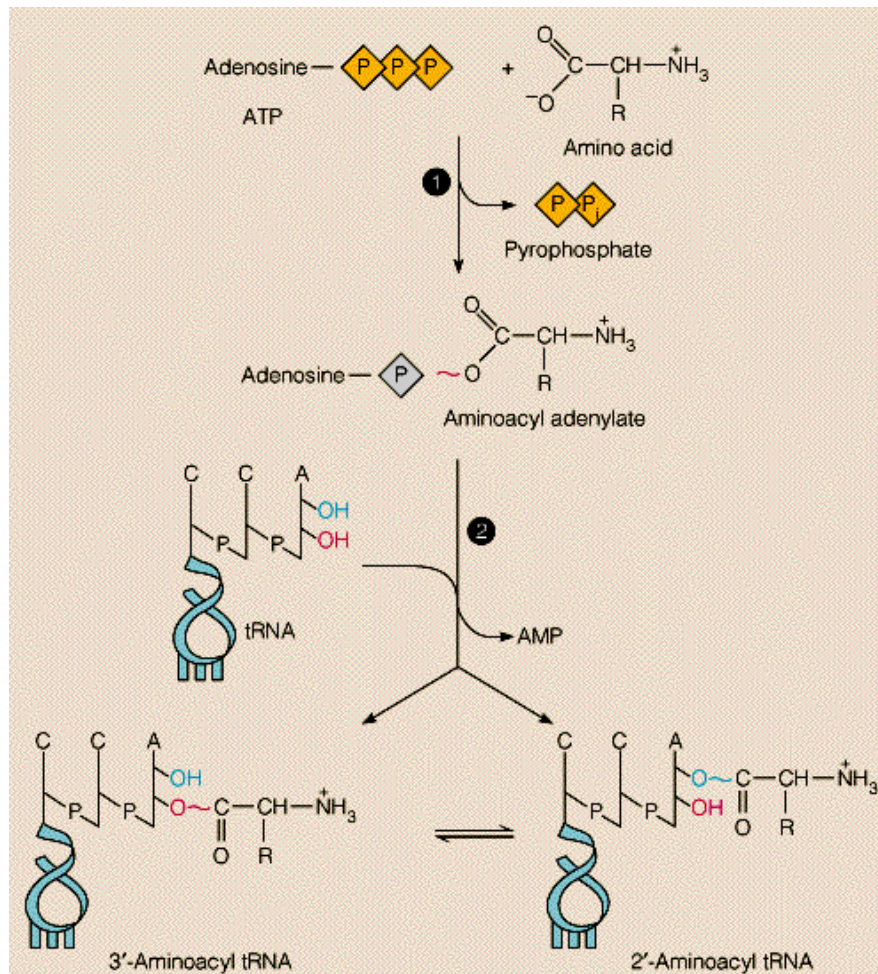
■ Initiation ■ Termination

The genetic code is nearly universal.

Exceptions:

- yeast
- mitochondria
- Tetrahymena*
- Mycoplasma*

Coupling of amino acids to tRNAs

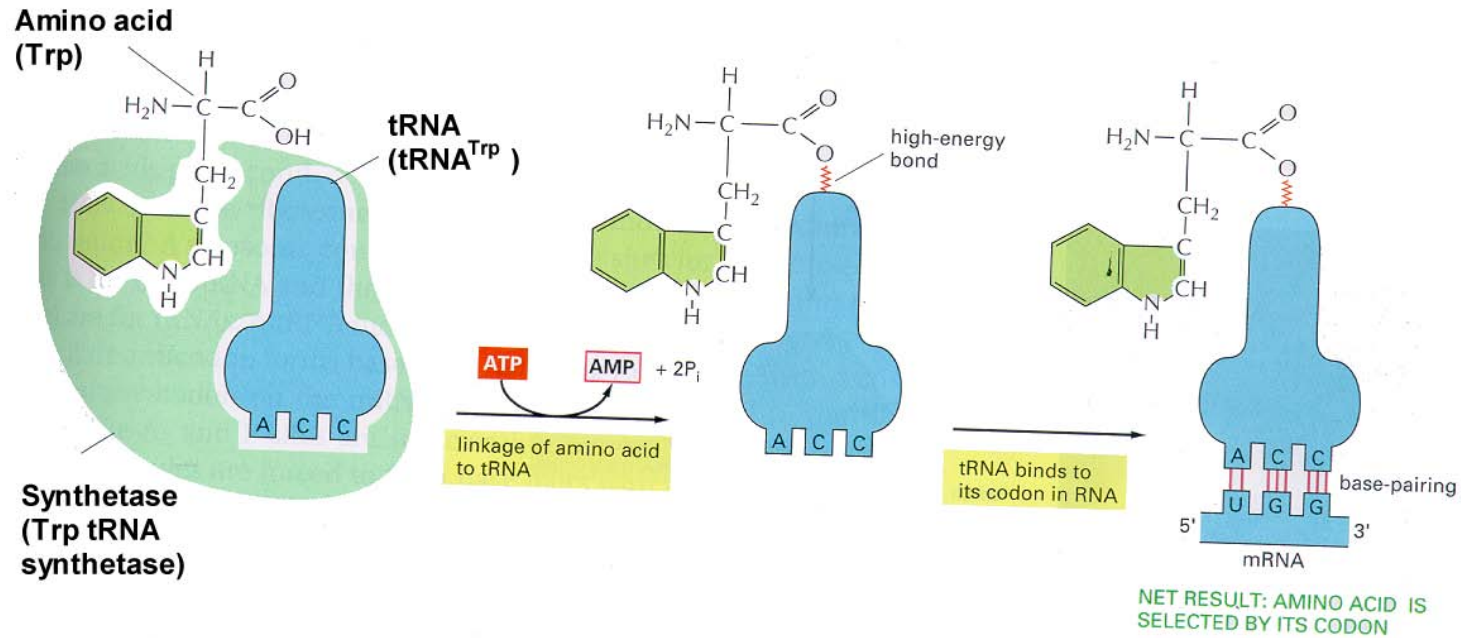


1. The amino acid is accepted by the aminoacyl-tRNA synthetase enzyme and is adenylated

2. The proper tRNA is accepted by the enzyme and the amino acid residue is transferred to the 2' or 3' OH of the 3'-terminal residue of the RNA

All reactions occur on the synthetase enzyme.

The Two Steps of Decoding



The genetic code is translated by means of two adaptors that act one after another.

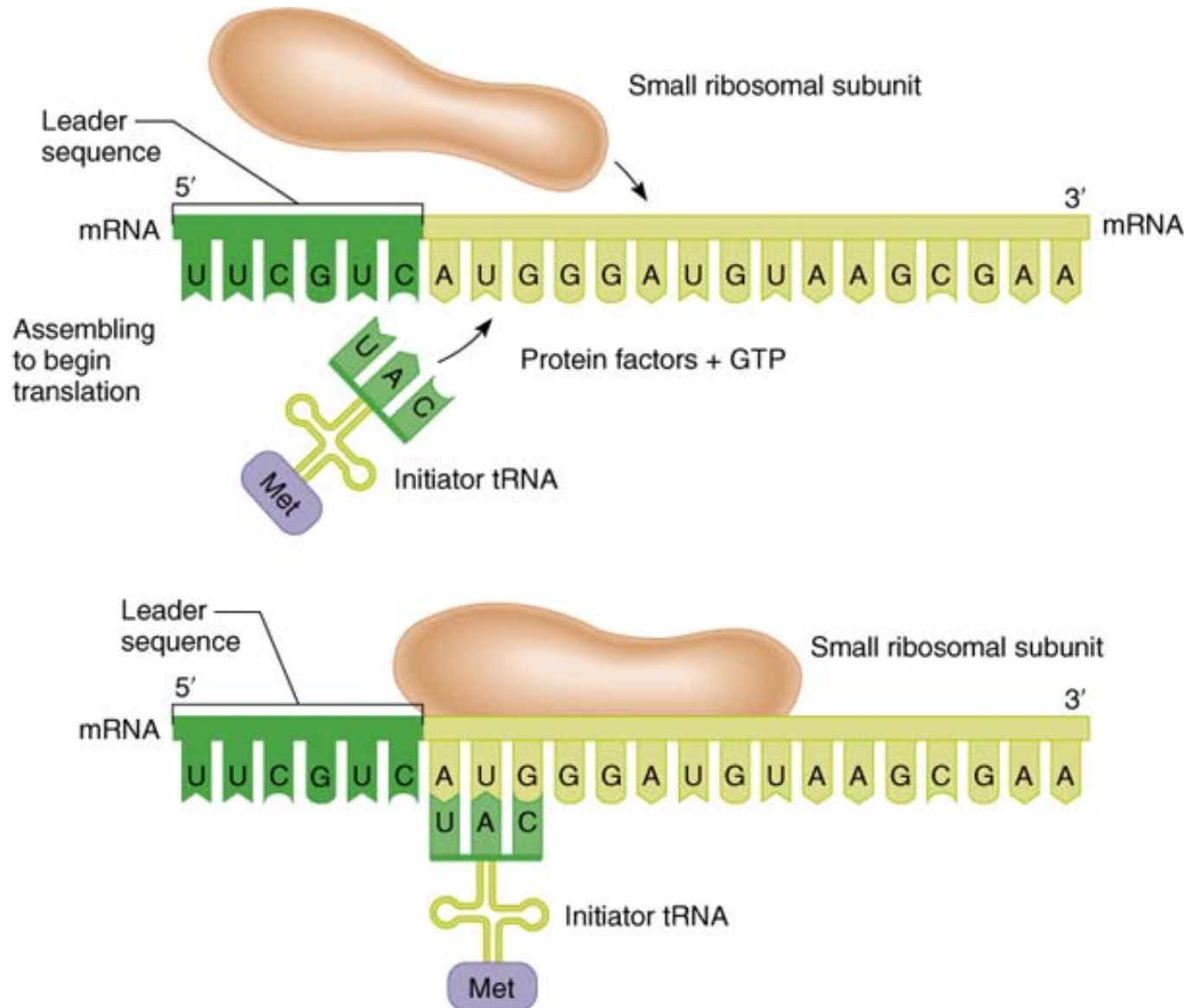
Translation – where all the guys from the last hour come together...

Initiation - translation begins at start codon
(AUG methionine)

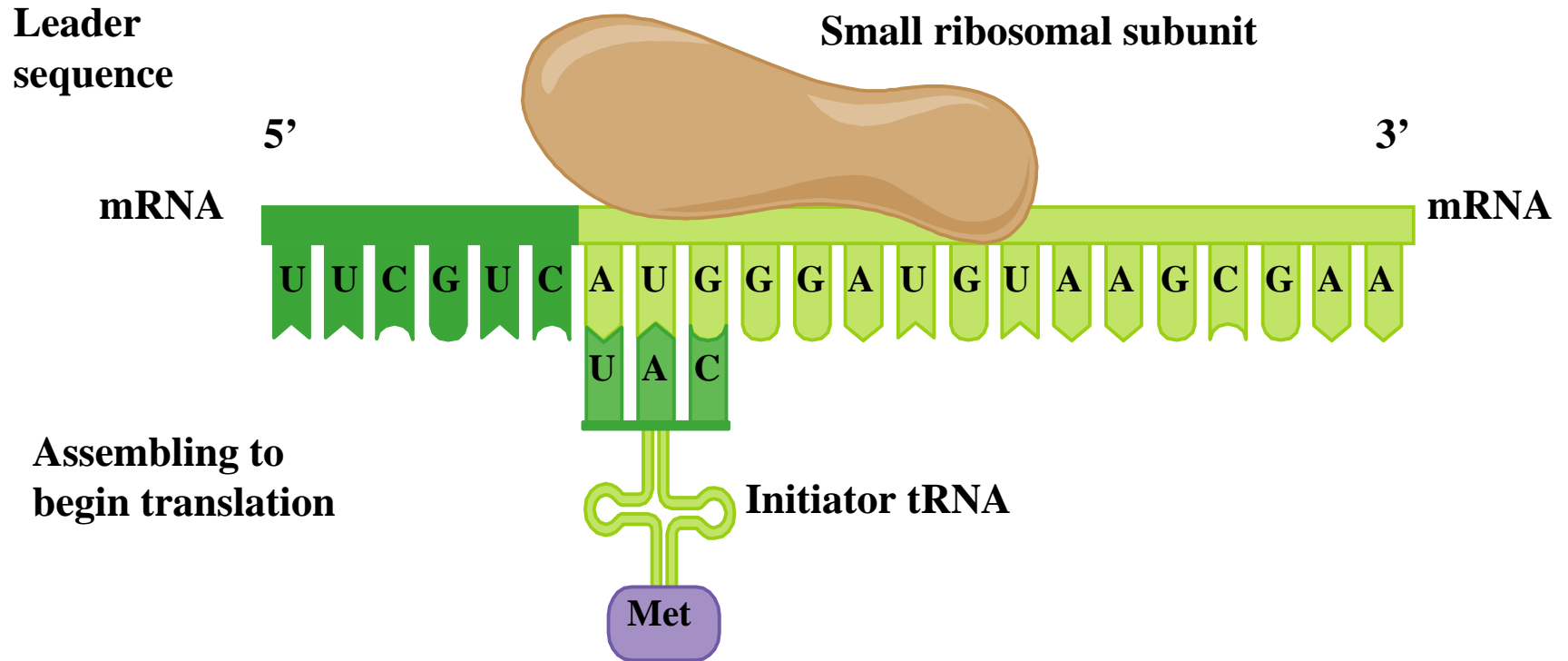
Elongation - the ribosome uses the tRNA anticodon to match codons to amino acids and adds those amino acids to the growing peptide chain

Termination - translation ends at the stop codon
UAA, UAG or UGA

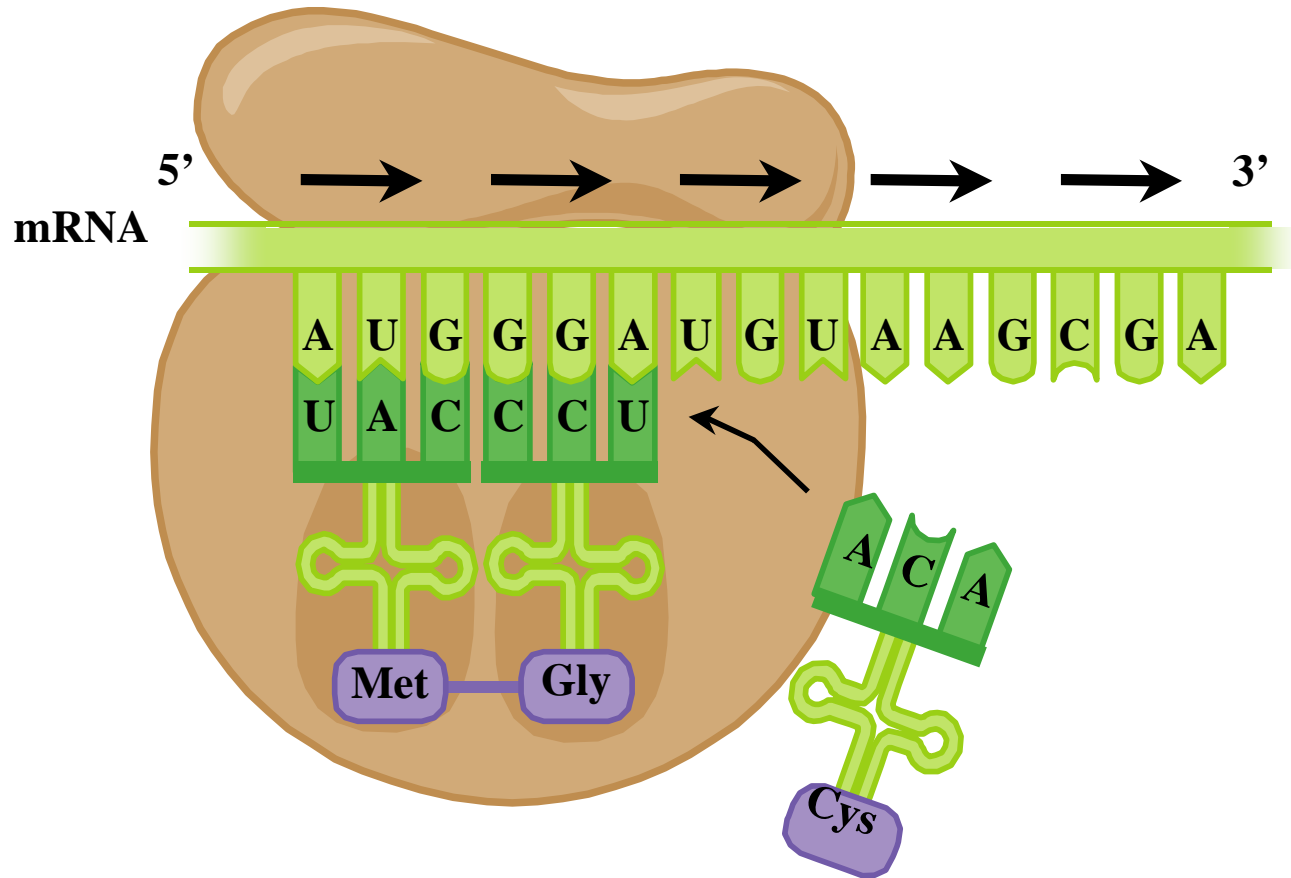
Translation initiation



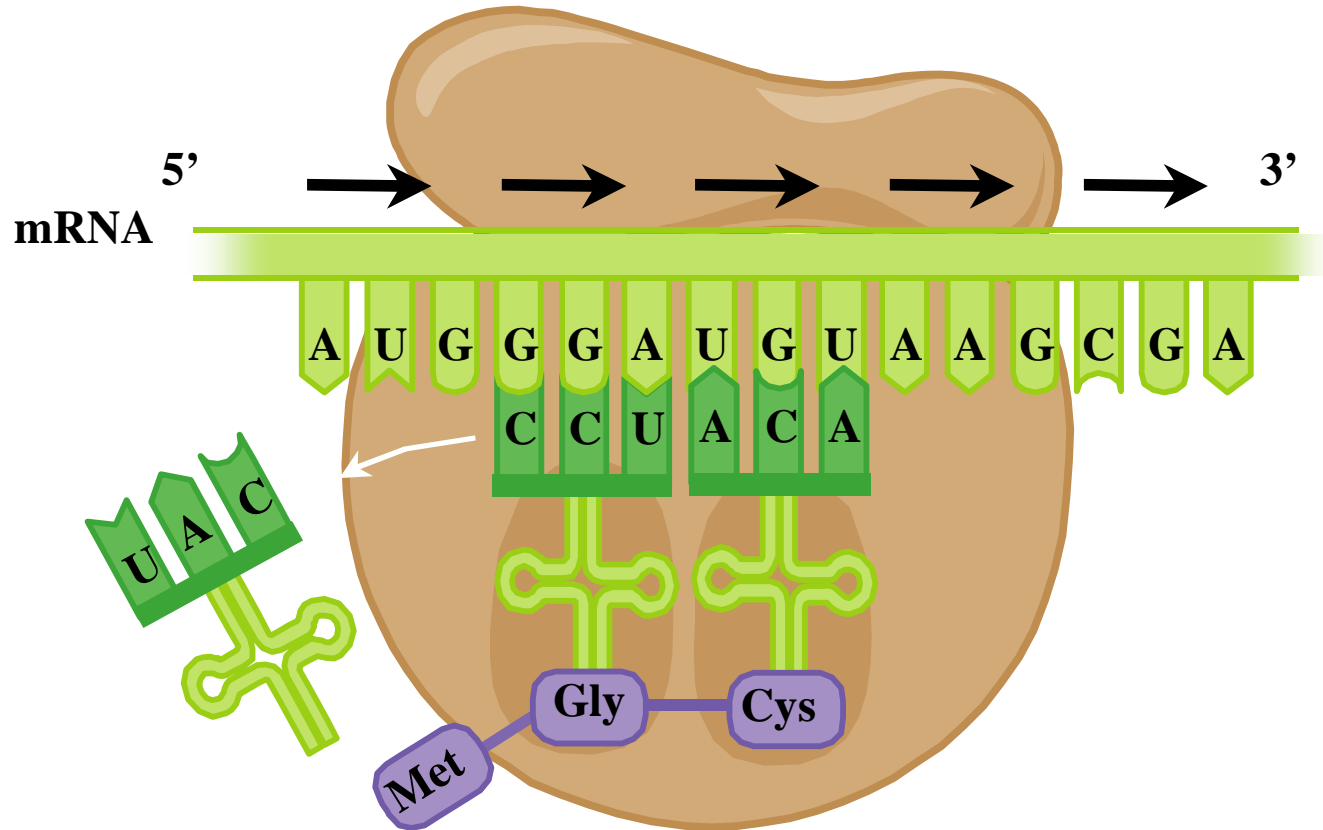
Translation initiation



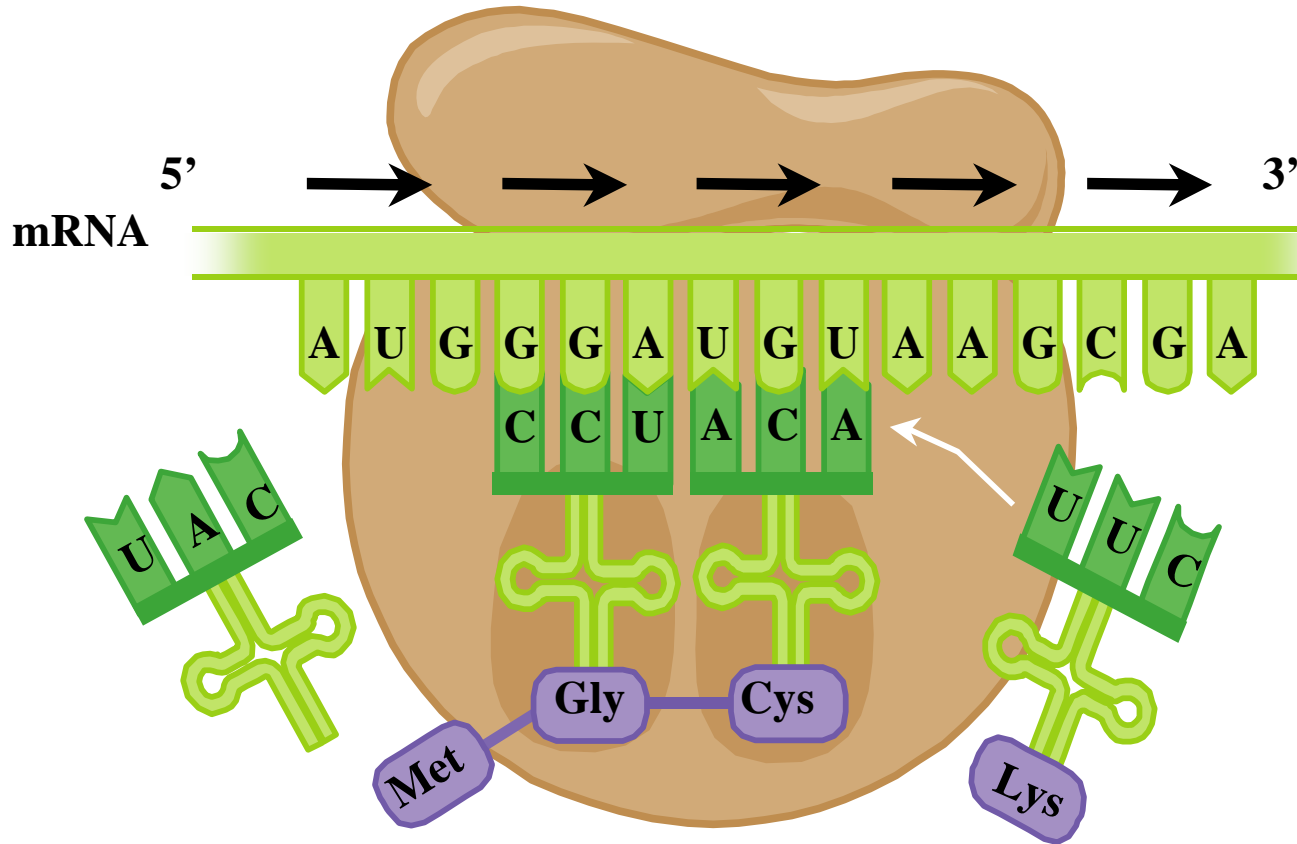
Translation Elongation



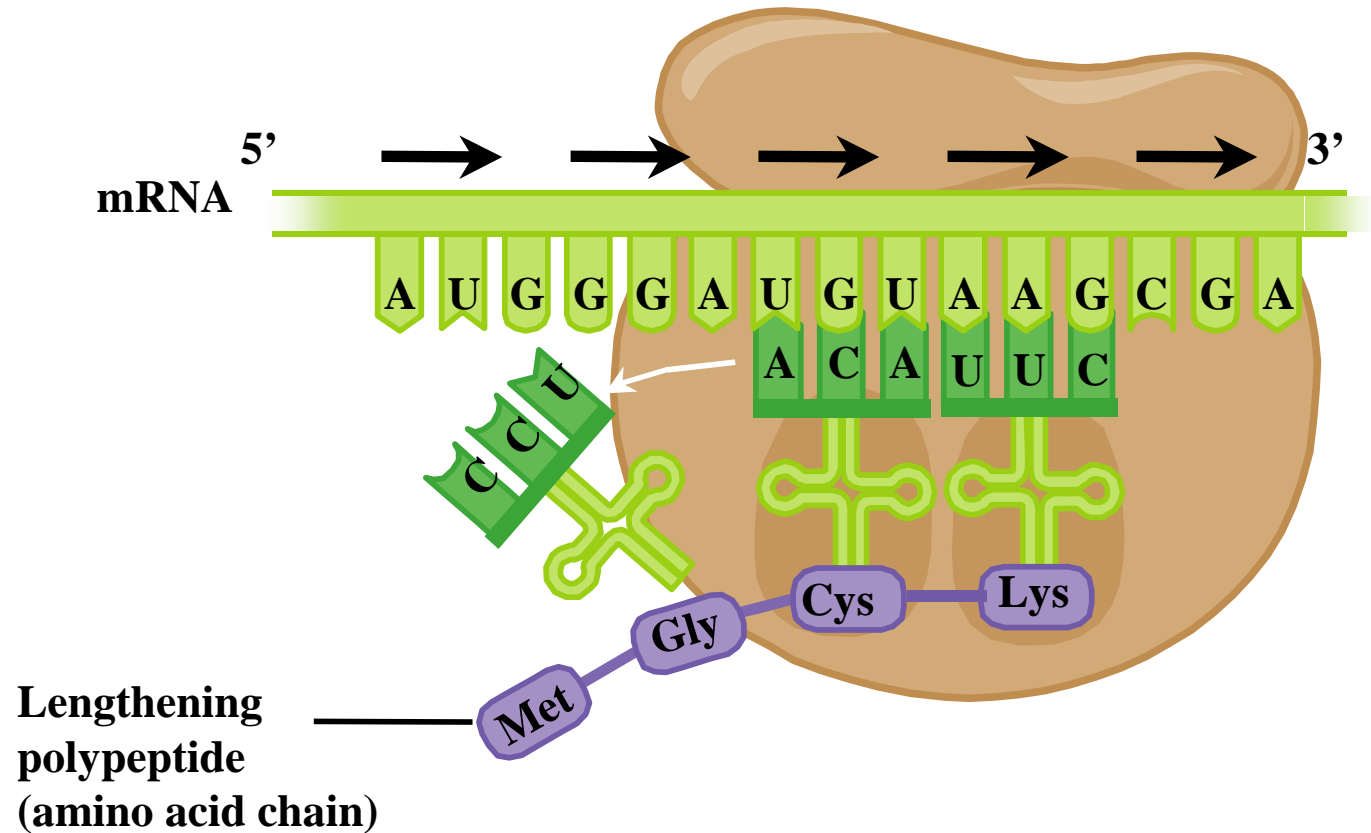
Translation Elongation



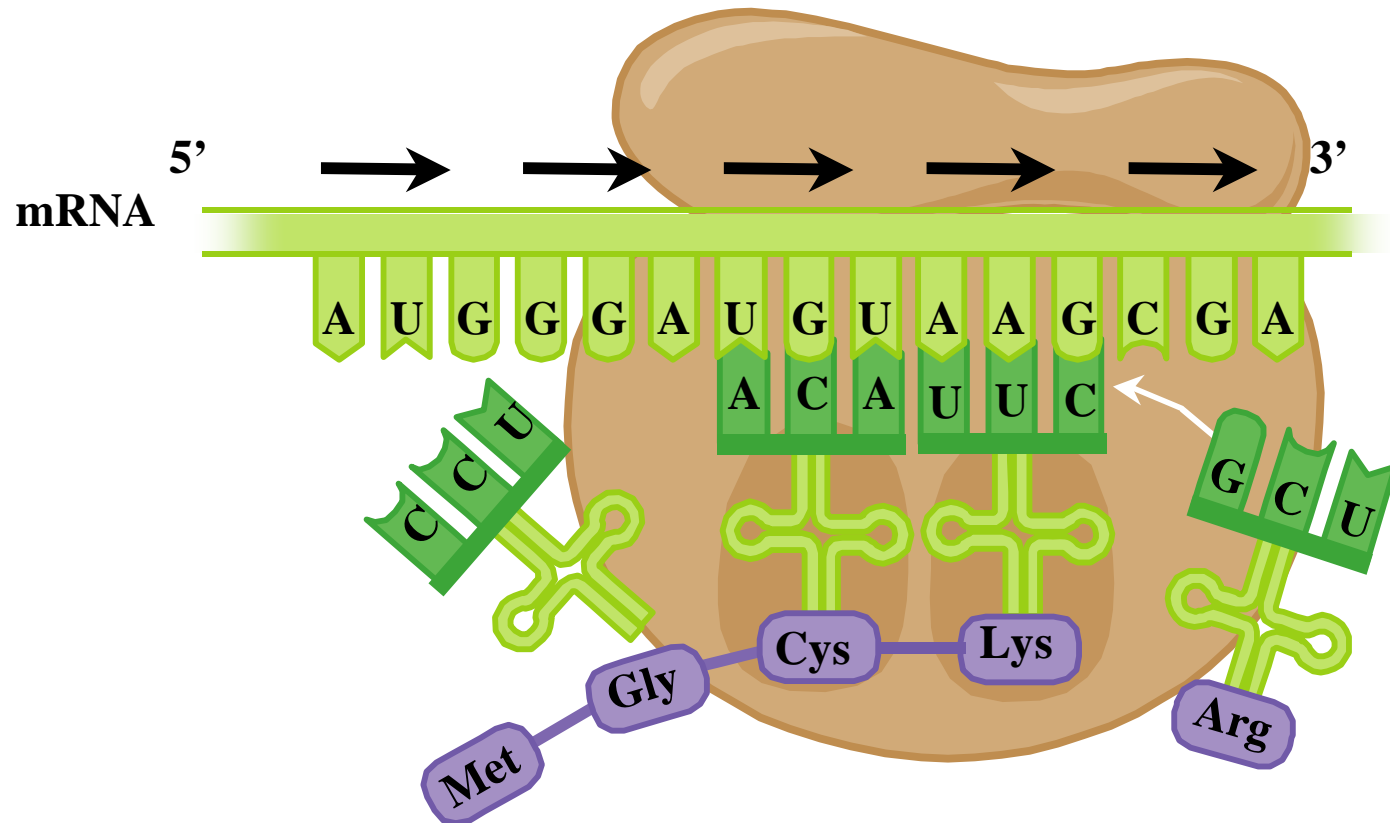
Translation Elongation



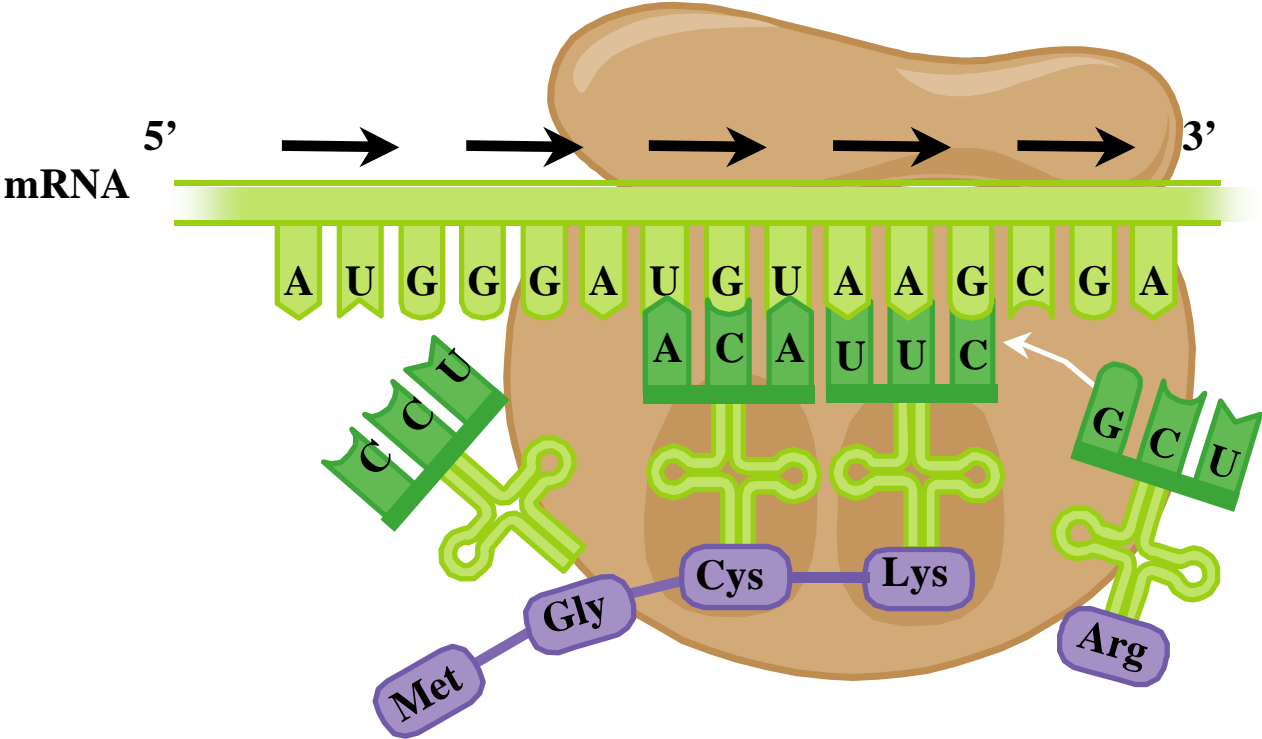
Translation Elongation



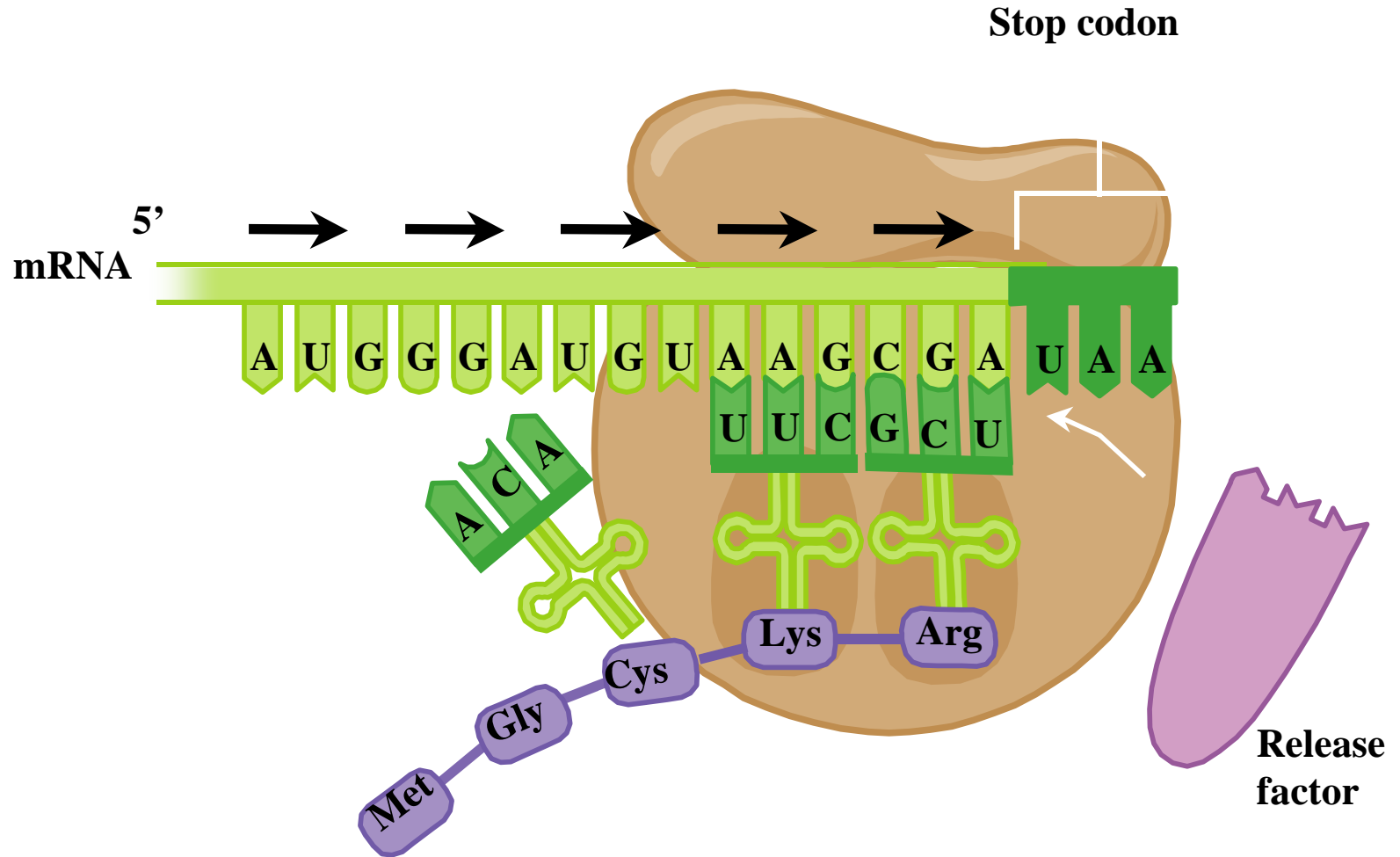
Translation Elongation



Translation Elongation

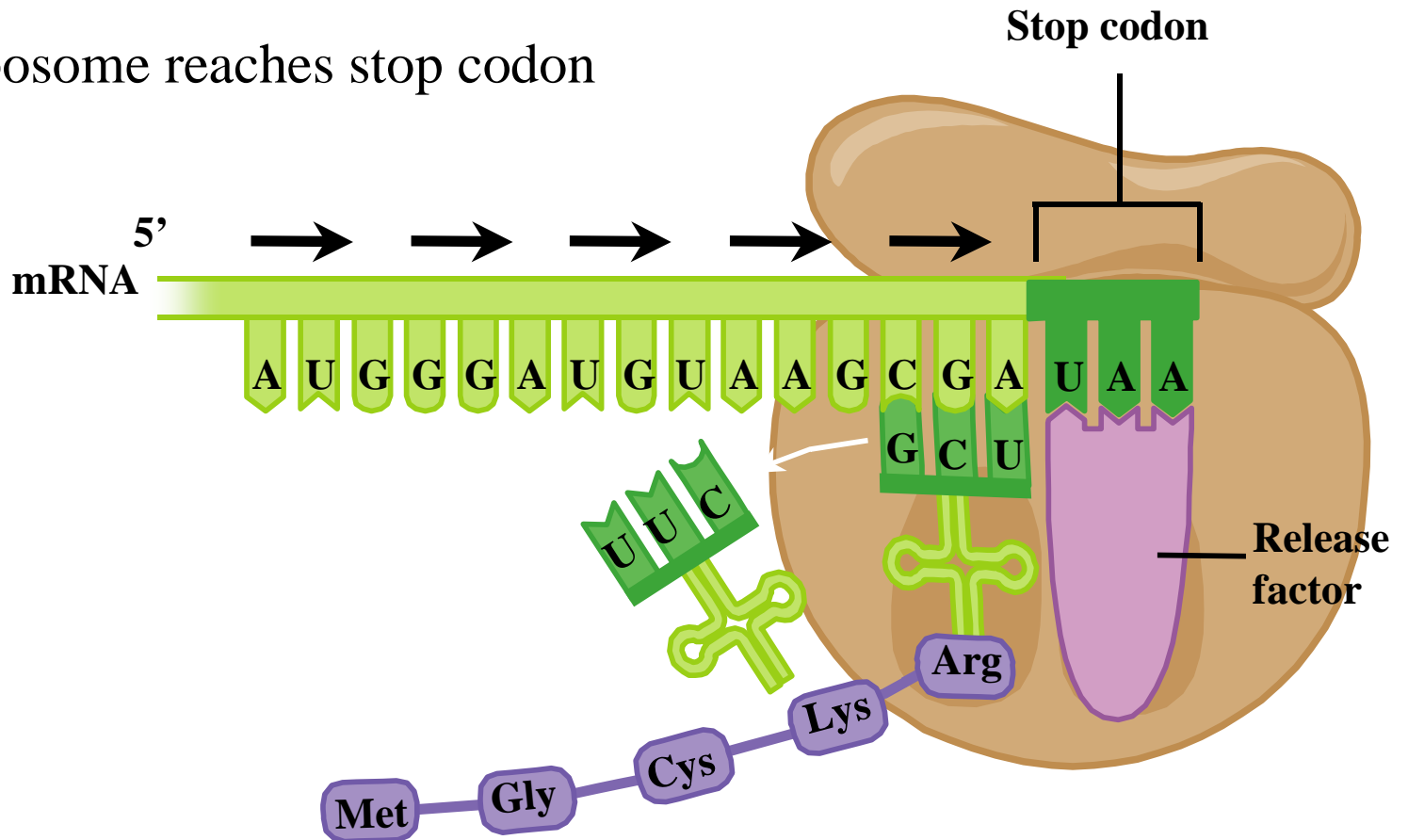


Translation Elongation



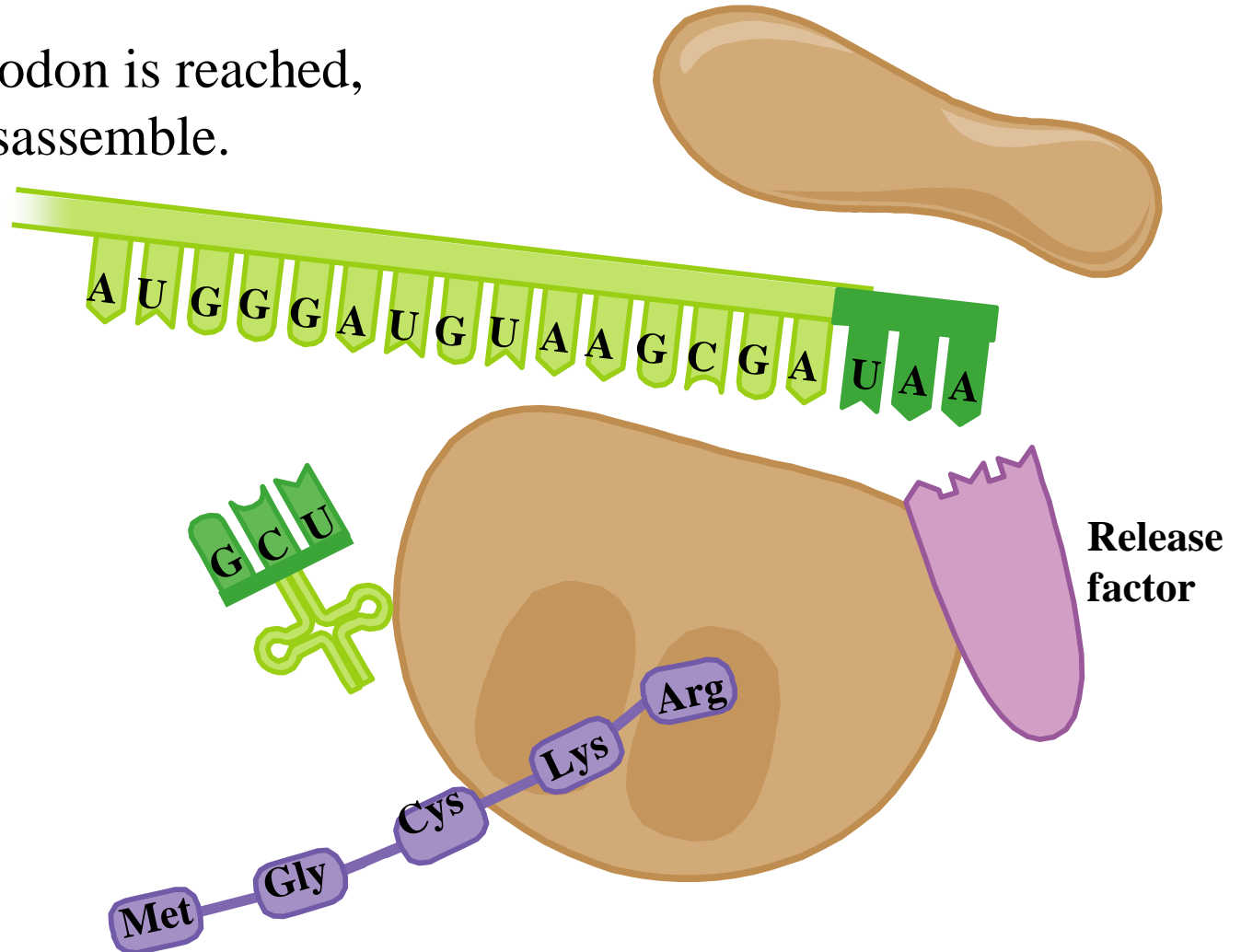
Translation Elongation

Ribosome reaches stop codon



Translation Termination

Once stop codon is reached,
elements disassemble.



Translation: multiple copies of a protein are made simultaneously

