# **Gene Expression Part II: Translation**

The process of transcribing an mRNA copy of the DNA code concludes with the exportation of the mature mRNA to the cytoplasm. This nucleotide sequence, however, cannot

have corporeal effects until it is translated into the proper amino acid sequence and arranged into a functional protein. The designated amino acids for each three nucleotides- known as the triplet codon- are outlined here:

As the enzymatic machinery moves along the mRNA molecule, the amino acids are bonded to form a polypeptide with a very specific order of amino acids via these steps:

#### U UUU }Phe UGU Cys UCU UAU Tyr C UCC Ser UUA T **UAA Stop** UCA UGA Stop UUG \Leu UCG UAG Stop UGG Trp G U CUU CCU CAUT CGU CUC CCC CAC CGC C C Pro Arg CAA CUA CCA CGA AG Gln CUG CCG CAG CGG AUU ACU AAU AGU AUC ACC AAC J AGC. C lle AUA AG ACA AAA T AGA AAG Lys AGG Arg ACG AUG U GUU GCU' GAU GGU GGC CA GUC GCC GAC G Gly Val Ala **GUA** GCA GAA GGA Glu G GUG GCG GAG GGG

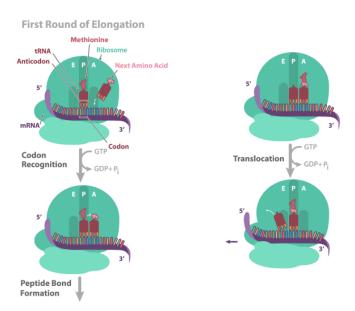
Second letter

#### Image credit: "The genetic code," by OpenStax College, Biology

### 1. Initiation

The large and small subunits of the ribosome assemble in conjunction with the mRNA molecule to

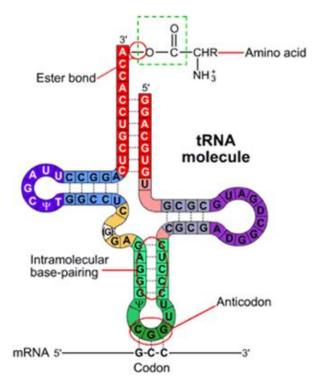
form the initiation complex. The large subunit has three sequential sites of binding- in the order E, P, and A along the 5' to 3' direction of the attached mRNA. When the mRNA start codon AUG enters the A site of the ribosome, it signals the beginning of the gene and translation will commence.



# 2. Elongation

The mRNA is read from 5' to 3' by complementary tRNA molecules whose anticodon recognizes the codon within the **A** site of binding based on the Watson-Crick base pairing rules that govern the processes of DNA synthesis and transcription, as well. That is to say, **A** binds to **U** for RNA (or **T** in DNA) and C binds G in both nucleic acid types.

Third letter



The corresponding amino acid is carried by the tRNA and transferred to the growing polypeptide chain. Elongation itself can be broken down into three repeating steps (each of which requires the input of GTP hydrolysis-derived energy) for each additional amino acid- codon recognition at the A site, peptide bond formation at the P site, and translocation as the tRNA exits from the E site.

# 3. Termination

When the mRNA's sequence reaches a stop codon (UAA, UAG, or UGA), no tRNA will recognize it, and it is instead bound to a release factor. The full polypeptide is then released, requiring the hydrolysis of another GTP to facilitate the energetic expenditure.