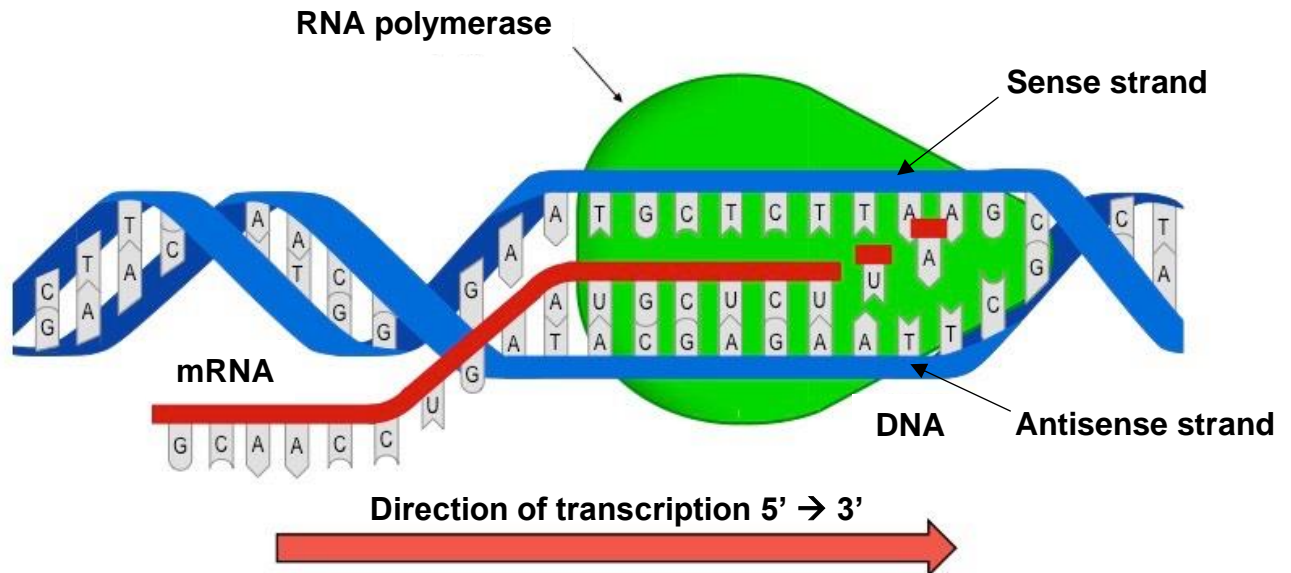
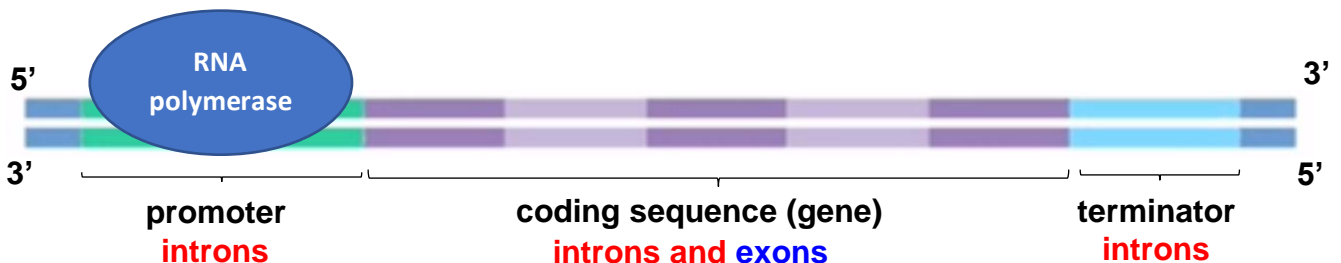


A. TRANSCRIPTION



- The strand that is **transcribed** is called the **antisense** strand and is **complementary** to the **mRNA** sequence (with T instead of U)
- The strand that is **not transcribed** is called the **sense** strand and is **identical** to the **mRNA** sequence (with T instead of U)

The process (DNA → mRNA)



What happens

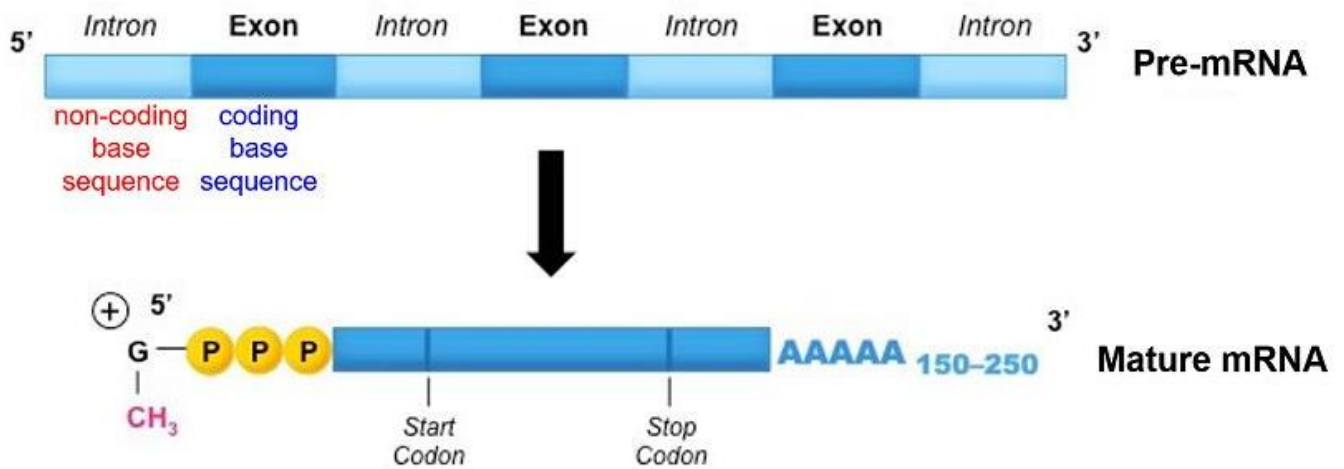
- **RNA polymerase** binds to **promoter** (at start of gene)
- (And) **separates strands** / unzips DNA / breaks hydrogen bonds
- **Antisense** strand acts as a **template**;
- **Free RNA nucleotides** added **5' → 3'** (direction);
- By **complementary base pairing** / adenine & uracil and cytosine & guanine;
- **RNA polymerase** joins **RNA nucleotides** together;
- **RNA polymerase** and **mRNA** detach at **terminator**;
- **DNA strands** attach **back** together / **H-bonds reform** between DNA strands;

How it is controlled

- **Transcription factors** bind to **promoter** and **help RNA polymerase** to bind
- **Repressor proteins** can bind to **promoter** and **prevent RNA polymerase** binding;
- **Methylation** (of cytosine bases) can **prevent transcription**;

B. POST-TRANSCRIPTION MODIFICATION OF mRNA

- The **introns** are **removed** and the **exons** are then **joined** together.
- **Only** happens in **eukaryotes** – **not prokaryotes**.



Capping	Splicing	Polyadenylation	Process
<p>Methyl (CH₃) group added to 5' end of mRNA to:</p> <ul style="list-style-type: none">- protect it from enzyme digestion- allow a ribosome to recognise it	<p>Introns are removed by splicing.</p> <p>Exons are joined together.</p>	<p>A long chain of adenine nucleotides (a poly-A tail) is added to the 3' end of the mRNA. This:</p> <ul style="list-style-type: none">- makes it more stable- helps it to leave the nucleus	

- Splicing of exons is the reason we can produce so **many different types of antibody**, each having a **specific shape**.

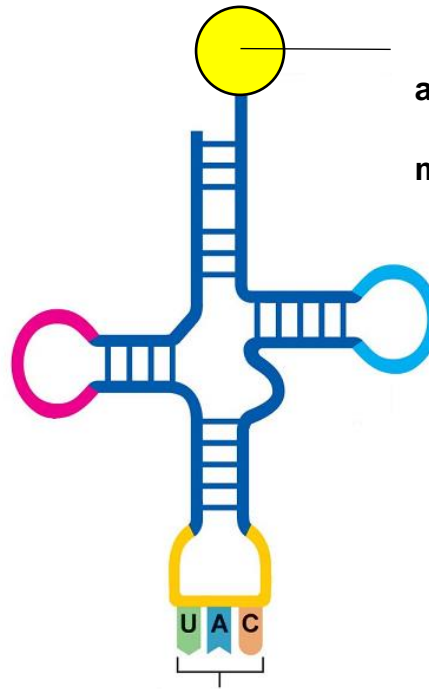
C. FUNCTIONS OF DNA BASE SEQUENCES THAT DO NOT CODE FOR PROTEINS

- **Non-coding** DNA base sequences can have **four** main roles:
 1. **Introns** – involved in **processing mRNA**
 2. **Coding for tRNA and rRNA** – these are involved in **translation**
 3. **Controlling gene expression/transcription** – **binding sites** for **proteins** that can **allow** or **prevent** transcription.
 4. **Telomeres** – **repetitive base sequences** at the **ends** of **chromosomes**, which **prevent** parts of genes here from being **lost** each time the **DNA** is **replicated**.

tRNA (transfer RNA)

The **specific amino acid** carried is **determined** by the **anticodon**

i.e. all tRNA molecules with the anticodon **UAC** will carry the amino acid **methionine** (Met)



Anticodon

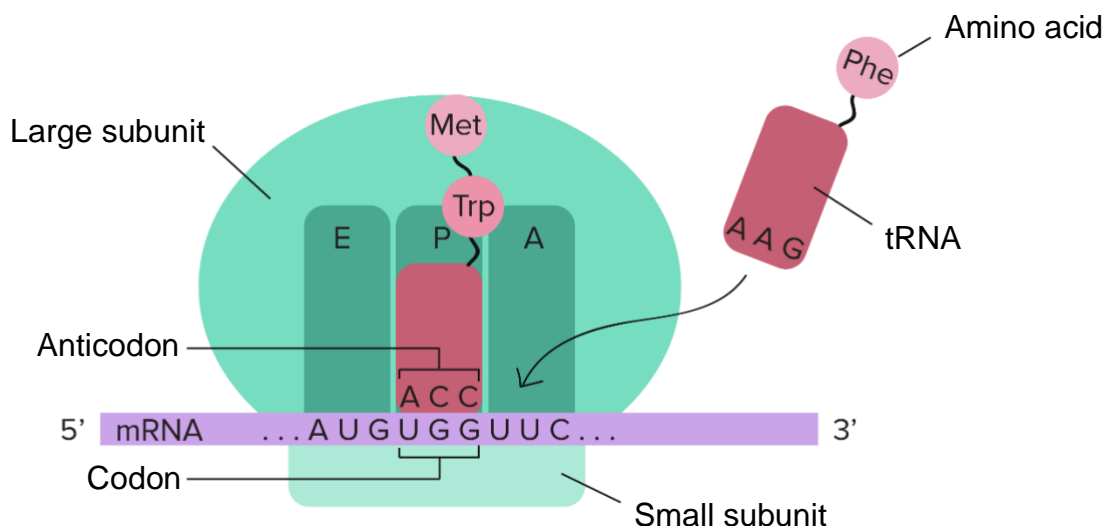
Three bases that bind to a **specific codon** on **mRNA**

Each tRNA molecule **carries** a **specific amino acid**

tRNA-activating enzyme **attaches** this **specific amino acid** to the **3' end** of a **tRNA molecule**, using **energy** from **ATP**

A ribosome

- Has **large** and **small subunits**.



- **A site** = tRNA carrying a new **Amino acid** binds here
- **P site** = tRNA carrying the growing **Protein** binds here
- **E site** = tRNA that has lost its amino acid **Exits** here

D. THE GENETIC CODE

- The **base sequence** of a **mRNA** molecule **codes for** the production of a **polypeptide**
- This **base sequence** is read by a ribosome in **triplets of bases** called **codons**
- Each **codon** codes for one **specific amino acid** in a protein chain
- The **order** of the **codons** in an **mRNA** determines the **order** of **amino acids** in a protein

The **genetic code** is **UNIVERSAL**

The **same mRNA codons** code for the **same amino acids** in **all organisms**

		Second letter					
		U	C	A	G		
First letter	U	UUU } Phe UUC } UUA } Leu UUG }	UCU } UCC } Ser UCA } UCG }	UAU } Tyr UAC } UAA Stop UAG Stop	UGU } Cys UGC } UGA Stop UGG Trp	U C A G	Third letter
	C	CUU } CUC } Leu CUA } CUG }	CCU } CCC } Pro CCA } CCG }	CAU } His CAC } CAA } Gln CAG }	CGU } CGC } Arg CGA } CGG }	U C A G	
	A	AUU } AUC } Ile AUA } AUG Met	ACU } ACC } Thr ACA } ACG }	AAU } Asn AAC } AAA } Lys AAG }	AGU } Ser AGC } AGA } Arg AGG }	U C A G	
	G	GUU } GUC } Val GUA } GUG }	GCU } GCC } Ala GCA } GCG }	GAU } Asp GAC } GAA } Glu GAG }	GGU } GGC } Gly GGA } GGG }	U C A G	

mRNA codons and the **amino acids** they code for

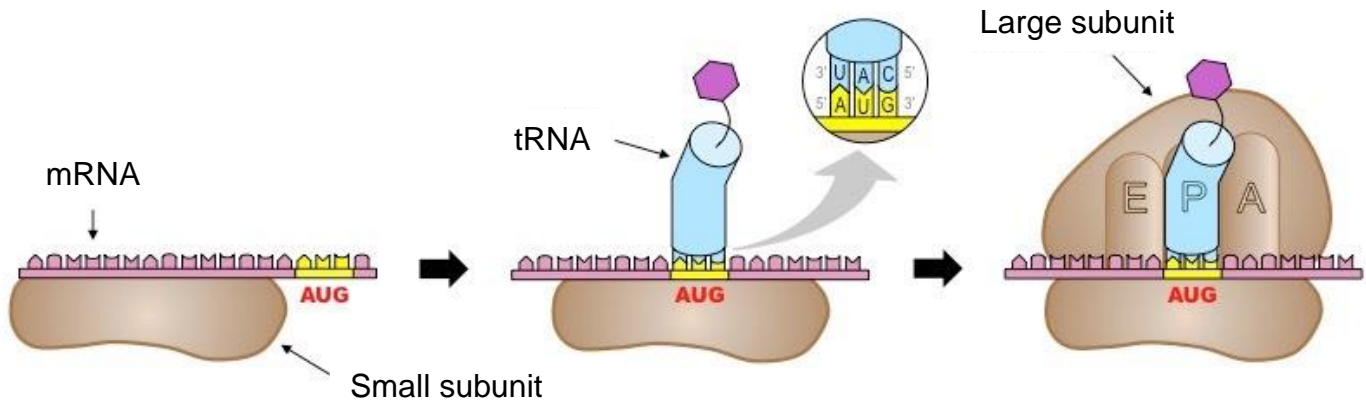
The **genetic code** is **DEGENERATE**

Different mRNA codons can code for the **same amino acid**

- The **mRNA codon** CCU codes for the **amino acid** Proline

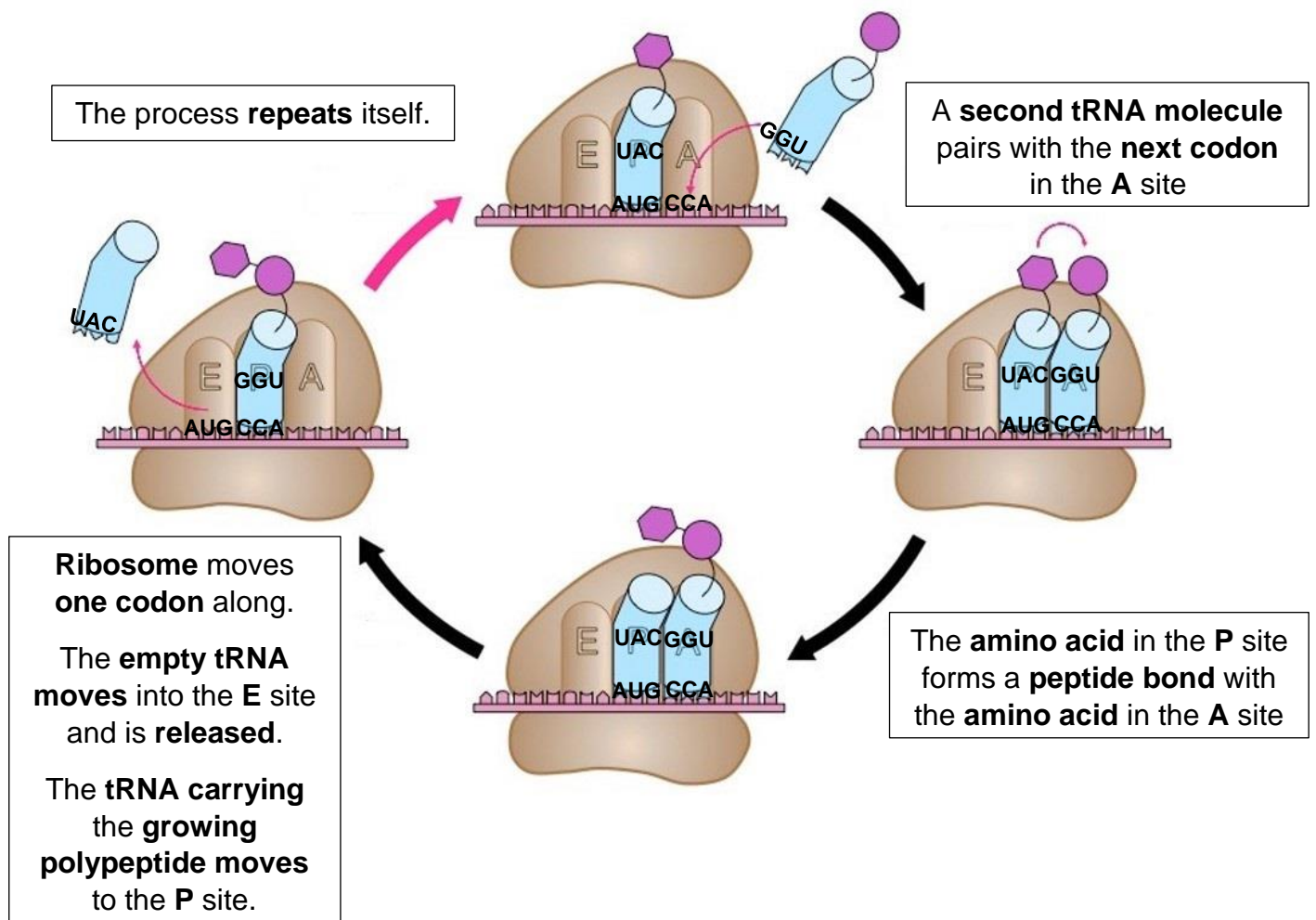
E. TRANSLATION (mRNA → POLYPEPTIDE/PROTEIN)

Stage 1: Initiation

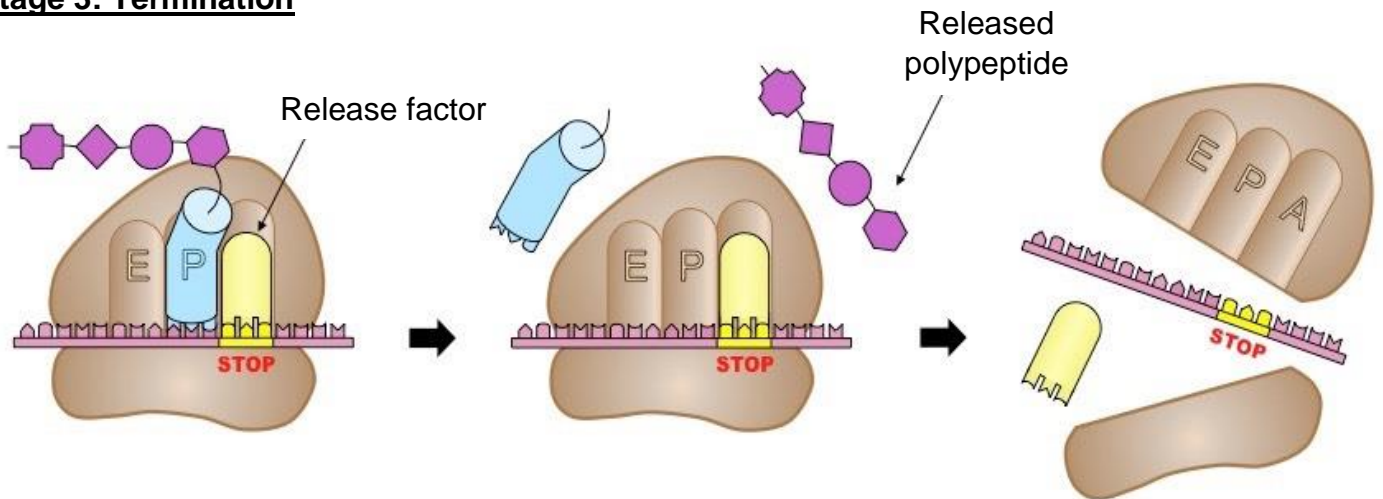


- **Small ribosome subunit** attaches to **5' end** of mRNA
- **First tRNA** with **anticodon UAC** attaches to **first codon (AUG)**;
- (tRNA) carries the amino acid **methionine/met**;
- **Large ribosome subunit** attaches (forming a complex);
- **First tRNA** is in the **P site**;

Stage 2: Elongation



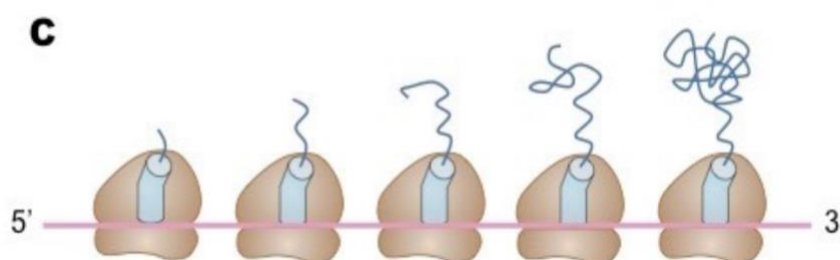
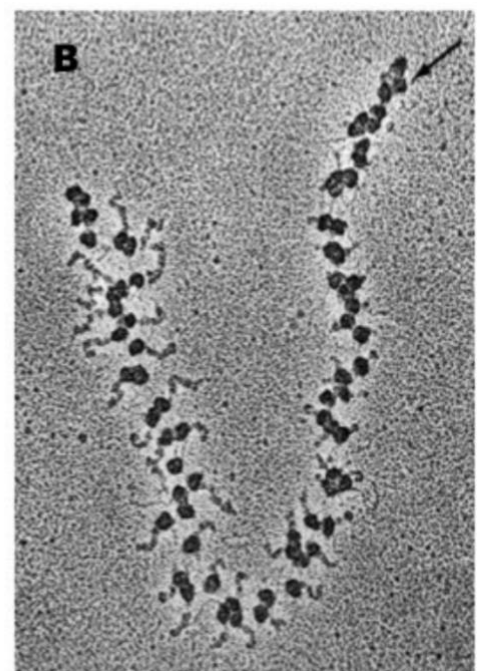
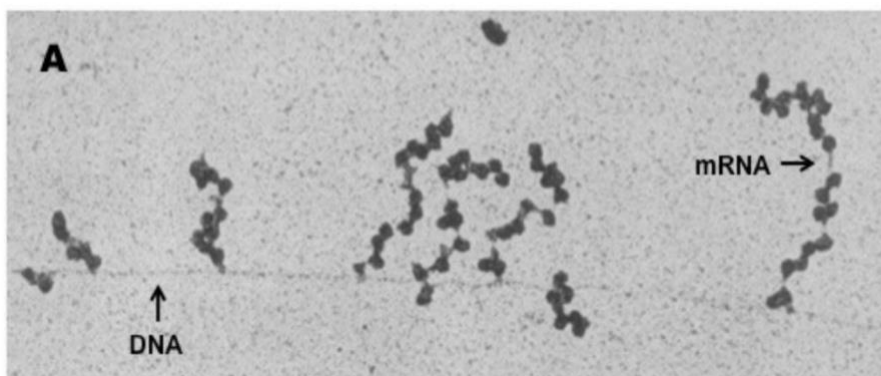
Stage 3: Termination



- **Ribosome** reaches a **stop codon**;
- This causes a **release factor** to **attach**;
- The **polypeptide** is **released**;
- The **ribosome separates** back into its **two subunits**;

Polysome

- A group of **two or more ribosomes** translating a mRNA base sequence **simultaneously**.
- They appear as **beads** on a **string** (bead = ribosome; string = mRNA).



F. COMPARE AND CONTRAST DNA REPLICATION & TRANSCRIPTION

DNA replication	Transcription
DNA → DNA	DNA → mRNA
Involves DNA polymerase	Involves RNA polymerase
Involves thymine	Involves uracil
Product is double stranded	Product is single stranded
Both strands act as templates	One strand acts as a template
Complete strand copied	Part of a strand copied
Happens in the nucleus	
DNA unzips/strands separate	
Complementary copy/strand produced	
DNA molecule zips back up (at the end)	

G. WHY IS THE DNA BASE SEQUENCE SO IMPORTANT?

DNA base sequence determines:

mRNA base sequence determines:

amino acid sequence determines:

tertiary structure of protein determines:

if the protein is functional or non-functional