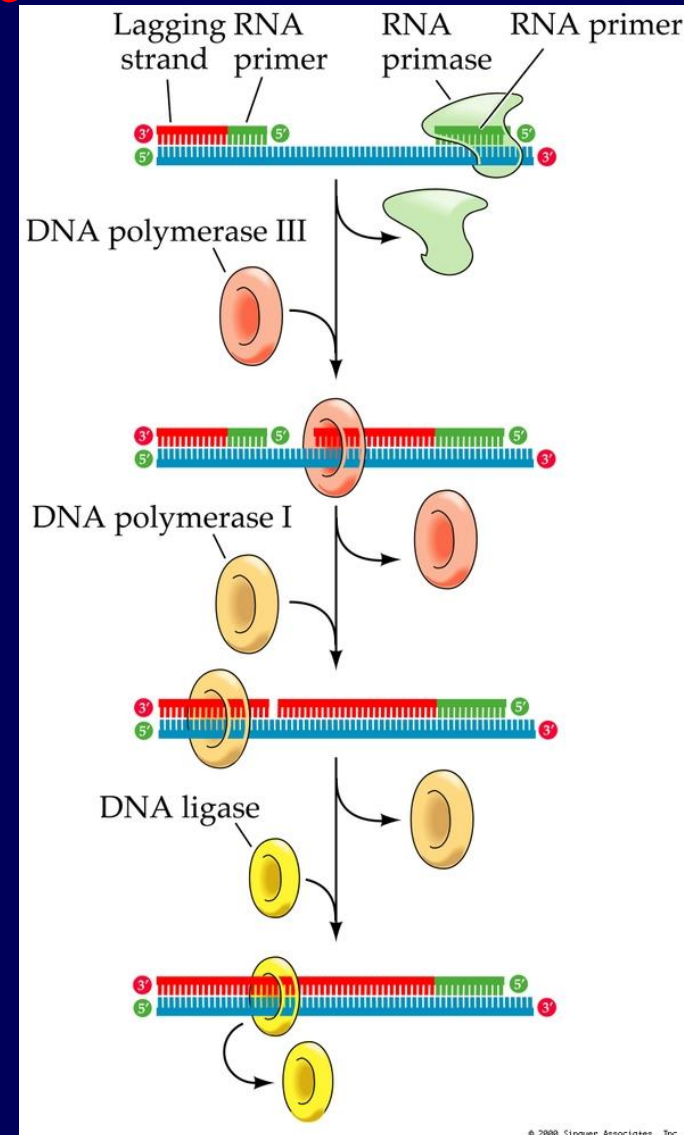


The Mechanism of DNA Replication

- Many proteins assist in DNA replication
- DNA helicases unwind the double helix, the template strands are stabilized by other proteins
- Single-stranded DNA binding proteins make the template available
- RNA primase catalyzes the synthesis of short RNA primers, to which nucleotides are added.
- DNA polymerase III extends the strand in the 5'-to-3' direction
- DNA polymerase I degrades the RNA primer and replaces it with DNA
- DNA ligase joins the DNA fragments into a continuous daughter strand



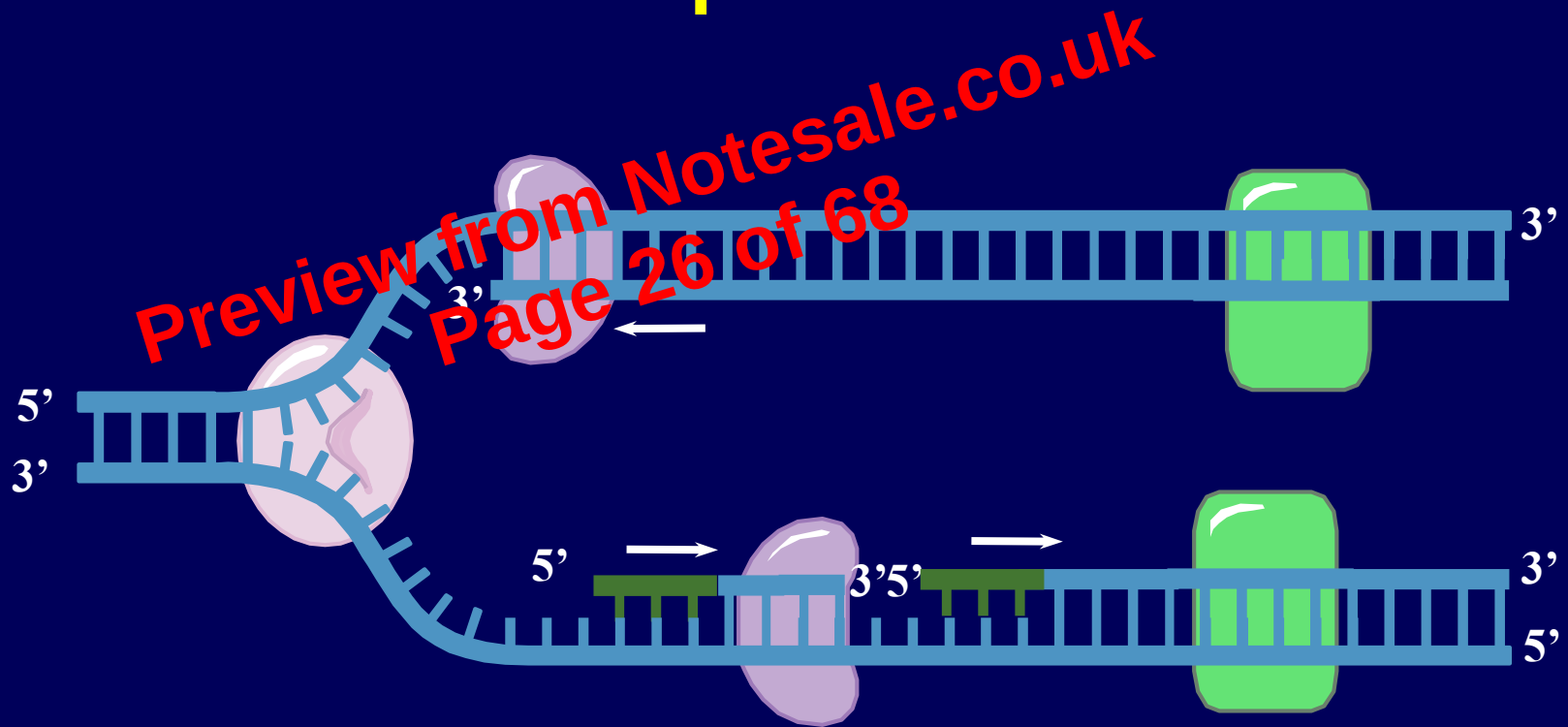
Replication



Leading strand synthesis continues in a 5' to 3' direction.

Discontinuous synthesis produces 5' to 3' DNA segments called Okazaki fragments.

Replication

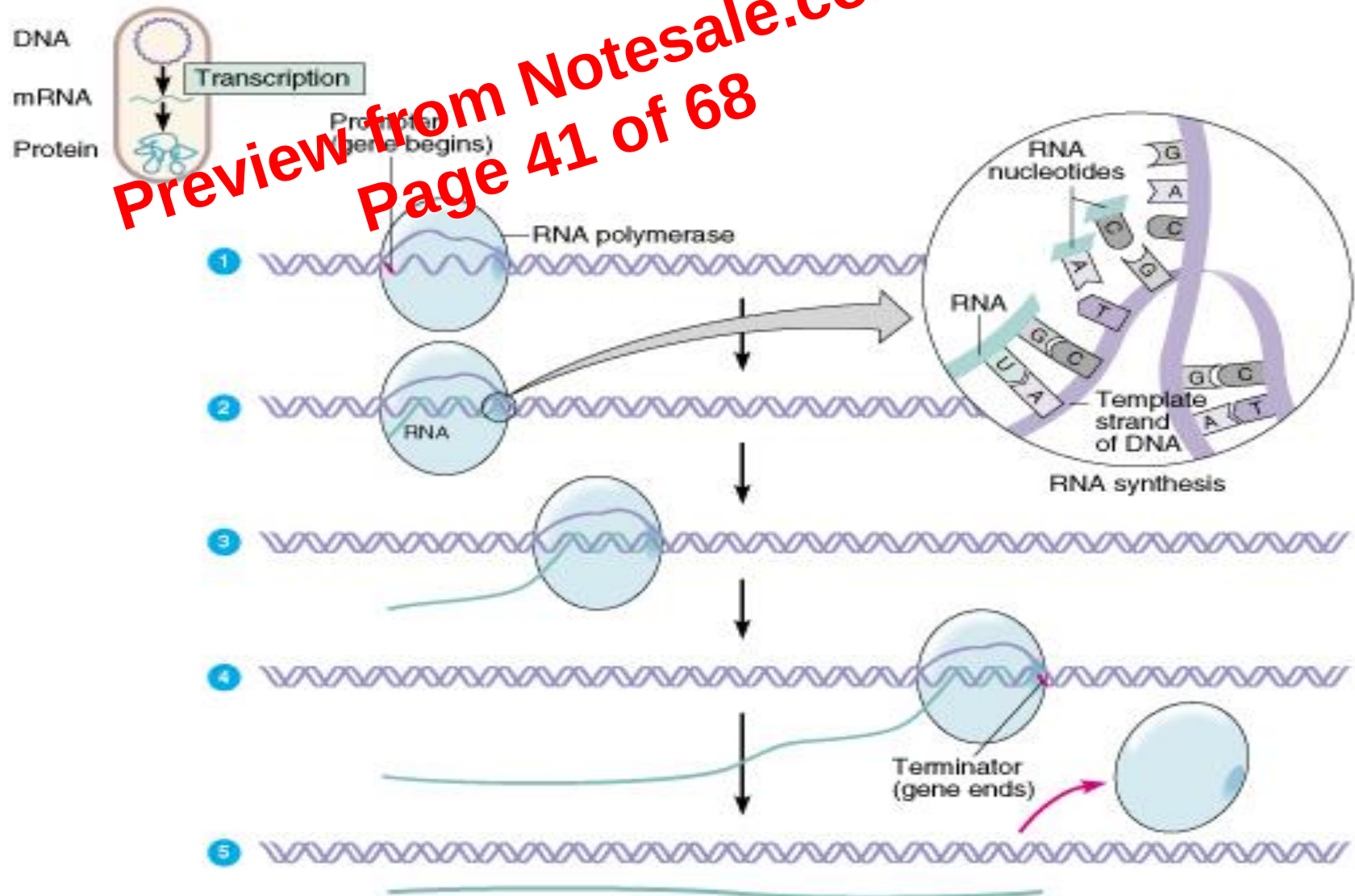


Polymerase activity of DNA polymerase I fills the gaps.

Ligase forms bonds between sugar-phosphate backbone.

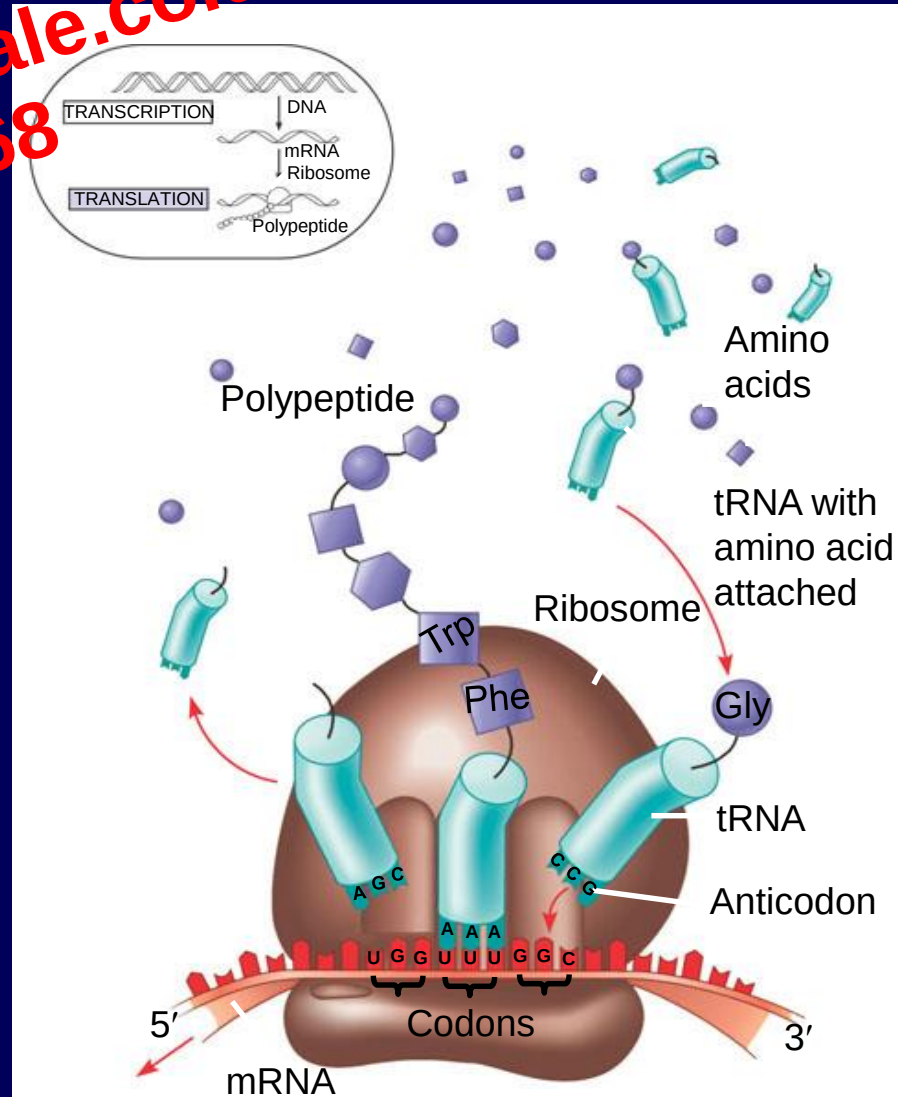
Transcription Overview

Preview from Notesale.co.uk
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Translation

- Translation is the RNA-directed synthesis of a polypeptide
- Translation involves
 - mRNA
 - Ribosomes - Ribosomal RNA
 - Transfer RNA
 - Genetic coding - codons



The Genetic Code

- Codons: 3 base code for the production of a specific amino acid, sequence of three of the four different nucleotides
- Since there are 4 bases and 3 positions in each codon, there are $4 \times 4 \times 4 = 64$ possible codons
- 64 codons but only 20 amino acids, therefore most have more than 1 codon
- 3 of the 64 codons are used as STOP signals; they are found at the end of every gene and mark the end of the protein
- One codon is used as a START signal: it is at the start of every protein
- Universal: in all living organisms

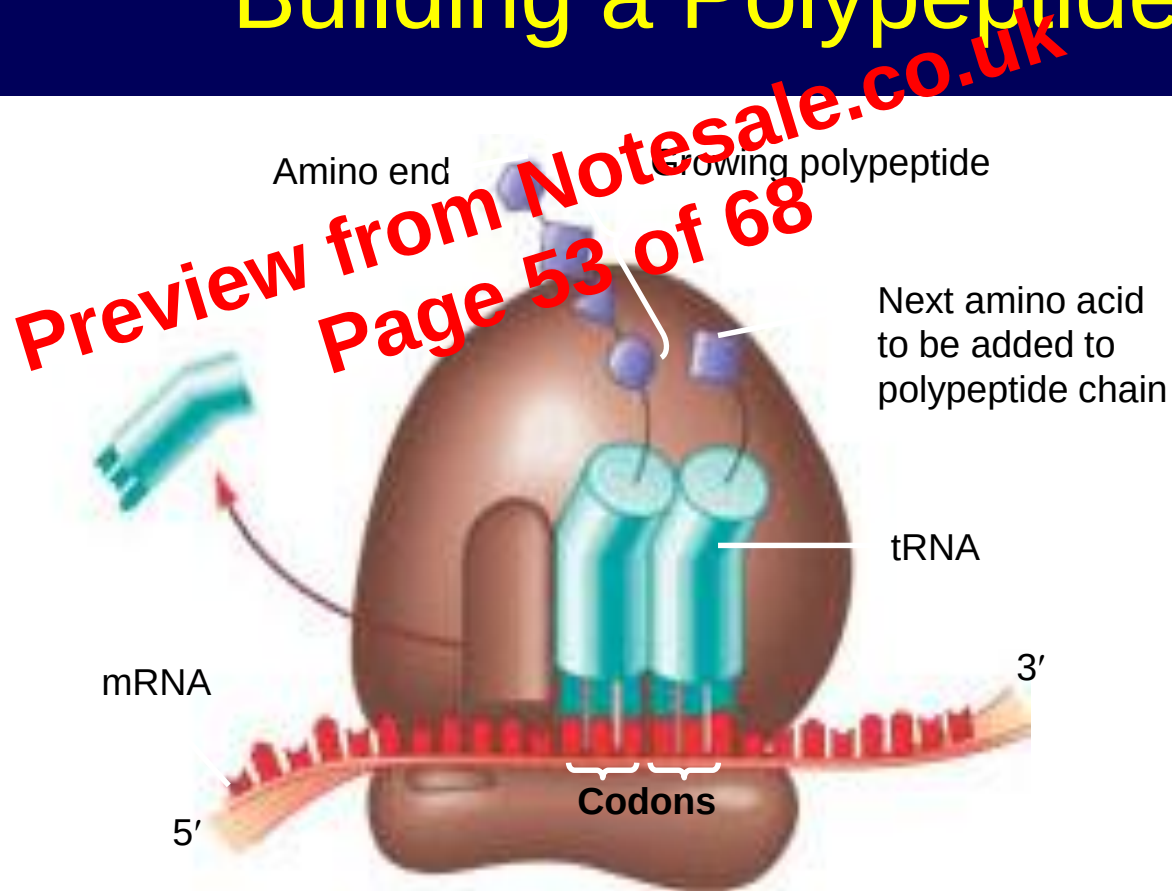
The Genetic Code

- A codon in messenger RNA is either translated into an amino acid or serves as a translational start/stop signal

A codon in messenger RNA is either translated into a amino acid or serves as a translational start/stop signal

		Second mRNA base										
		U		C		A		G				
First mRNA base (5' end)	U	UUU	Phe	UCU	Ser	UAU	Tyr	UGU	Cys	U		
		UUC		UCC		UAC		UGC			C	
		UUA		UCA		UAA Stop		UGA Stop				A
		UUG		UCG		UAG Stop		UGG Trp				
	C	CUU	Leu	CCU	Pro	CAU	His	CGU	Arg	U		
		CUC		CCC		CAC		CGC			C	
		CUA		CCA		CAA		CGA				A
		CUG		CCG		CAG		CGG				
	A	AUU	Ile	ACU	Thr	AAU	Asn	AGU	Ser	U		
		AUC		ACC		AAC		AGC			C	
		AUA		ACA		AAA		AGA				A
		AUG Met or start		ACG		AAG		AGG				
	G	GUU	Val	GCU	Ala	GAU	Asp	GGU	Gly	U		
		GUC		GCC		GAC		GGC			C	
		GUA		GCA		GAA		GGA				A
		GUG		GCG		GAG		GGG				
		Third mRNA base (3' end)										

Building a Polypeptide



(c) **Schematic model with mRNA and tRNA.** A tRNA fits into a binding site when its anticodon base-pairs with an mRNA codon. The P site holds the tRNA attached to the growing polypeptide. The A site holds the tRNA carrying the next amino acid to be added to the polypeptide chain. Discharged tRNA leaves via the E site.

Mutagens

- **Mutagens** are chemical or physical agents that interact with DNA to cause mutations.
- Physical agents include high-energy radiation like X-rays and ultraviolet light
- Chemical mutagens fall into several categories.
 - Chemicals that are base analogues that may be substituted into DNA, but they pair incorrectly during DNA replication.
 - Interference with DNA replication by inserting into DNA and distorting the double helix.
 - Chemical changes in bases that change their pairing properties.
- Tests are often used as a preliminary screen of chemicals to identify those that may cause cancer
- Most carcinogens are mutagenic and most mutagens are carcinogenic.

Viral Mutagens

- Scientists have recognized a number of *tumor viruses* that cause cancer in various animals, including humans
- About 15% of human cancers are caused by viral infections that disrupt normal control of cell division
- All tumor viruses transform cells into cancer cells through the integration of viral nucleic acid into host cell DNA.

Substitutions

- A base-pair substitution is the replacement of one nucleotide and its partner with another pair of nucleotides
 - Silent - changes a codon but codes for the same amino acid
 - Missense - substitutions that change a codon for one amino acid into a codon for a different amino acid
 - Nonsense - substitutions that change a codon for one amino acid into a stop codon

