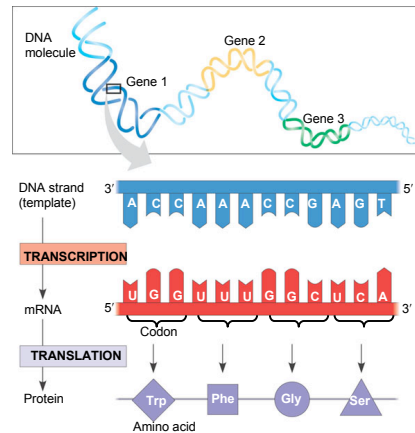


From Gene to Phenotype- part 2



Lecture Outline 11/7/05

- The central dogma:
 - DNA->RNA->protein
- Various types of RNA
- The genetic code
- Mechanisms of translation
 - Initiation, Elongation, Termination

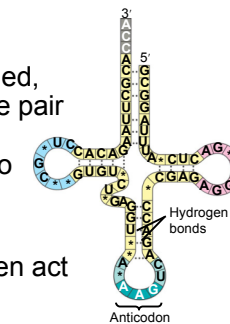
Exam 3 is next Monday. It will cover mitosis and meiosis, DNA synthesis, transcription, translation, genetics of viruses. (chapters 12, 13, 16, 17, part of 18)

Four types of RNA

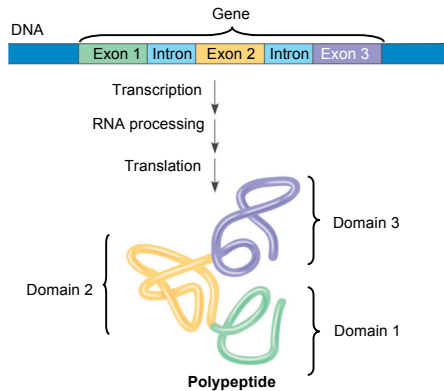
- mRNA
 - Messenger RNA, encodes the amino acid sequence of a polypeptide
- rRNA
 - Ribosomal RNA, forms complexes with protein called ribosomes, which translate mRNA to protein
- tRNA
 - Transfer RNA, transports amino acids to ribosomes during protein synthesis
- snRNA
 - Small nuclear RNA, forms complexes with proteins used in eukaryotic RNA processing

RNA will fold to specific shapes

- Because RNA is single-stranded, parts of the molecule can base pair with other parts of the same molecule, causing it to fold into defined shapes.
- Some RNA molecules can even act as enzymes (ribozymes)



Correspondence between exons and protein domains



The genetic code

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

The code is a triplet code

- the new boy saw the big cat eat the hot dog
- The first evidence for triplet code came from deletion experiments
 - the new boy saw the big cat eat the hot dog
 - the neoy boy saw the big cat eat the hot dog
 - the neoy awt heb icg ate att heh otd og
- Function could be restored by an insertion nearby
 - the new boy saw the big cat eat the hot dog
 - the neoy aawt heb icg ate att heh otd og

The code is a triplet code

- . . . or by two more deletions to get it back into the correct “reading frame”
 - the neo boy saw the big cat eat the hot dog
 - the neoy awt heb icg ate att heh otd og
 - the neosa wte big cat eat the hot dog

Deciphering the Code

- Your book tells how people used synthetic mRNAs and *in vitro* translation to determine decipher the codons.
- E.g. UUUUUUUU -> phenylalanine only
 - UCUCUCUCUCU -> mix of leucine and serine
 - Why is it a mixture?

The code is redundant

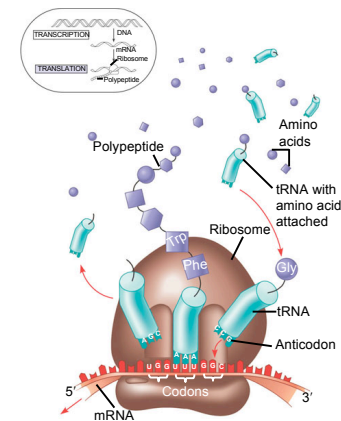
- Several different codons encode the same amino acid

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

The code is comma free

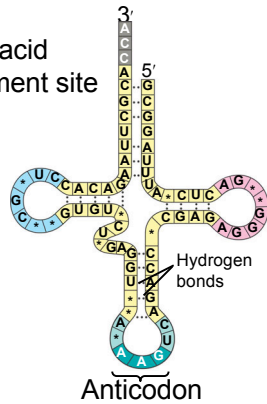
- No punctuation between words.
 - Therefore deletions cause frameshifts
- It does have start and stop signals, however.
 - Start: AUG
 - Stop: UAG, UAA, UGA

Translation: the basic concept

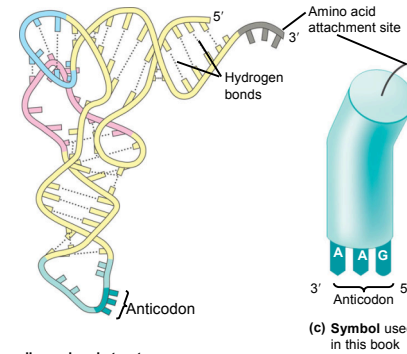


The structure of transfer RNA (tRNA)

Amino acid attachment site

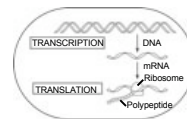


tRNA



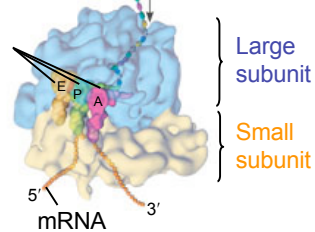
(b) Three-dimensional structure

Ribosomes



Growing polypeptide

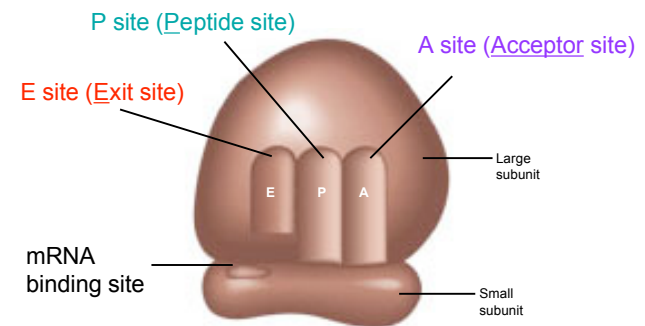
tRNA molecules



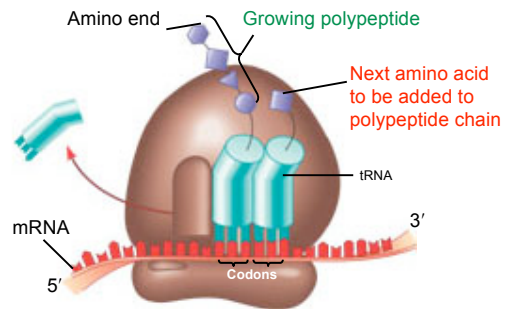
RNA + protein.

We now think that RNA catalyzes the reactions.

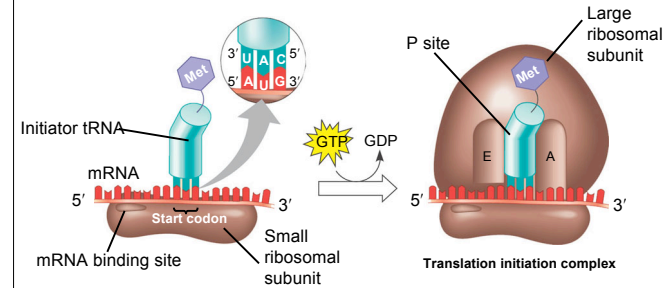
EPA model of a ribosome



New amino acids are added to the carboxyl end of the polypeptide

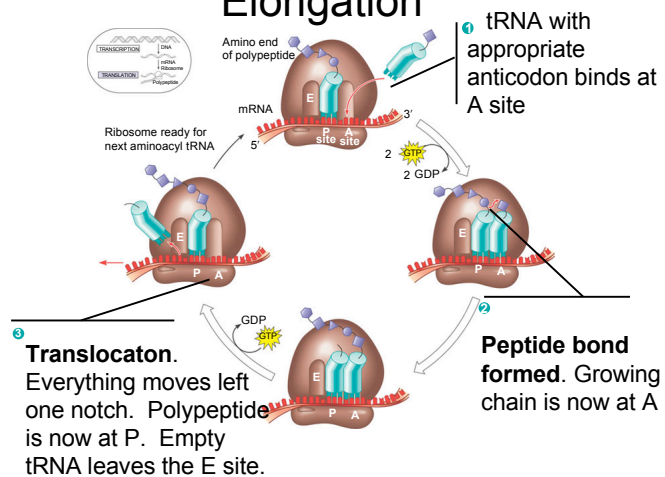


The initiation of translation

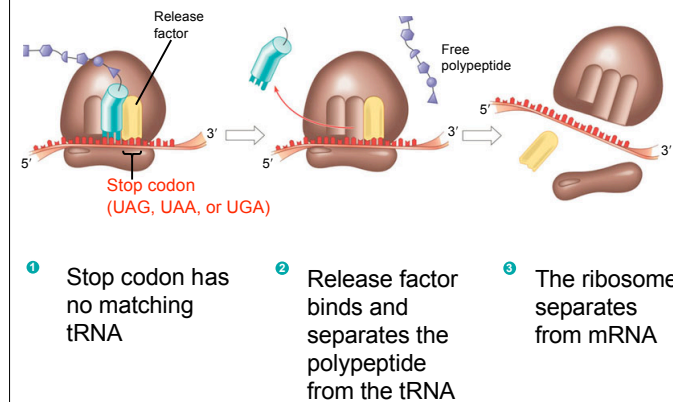


- 1 Small subunit binds just upstream of start
Special initiator methionine binds to the start codon (AUG)
- 2 The large subunit can now bind to make the active ribosome

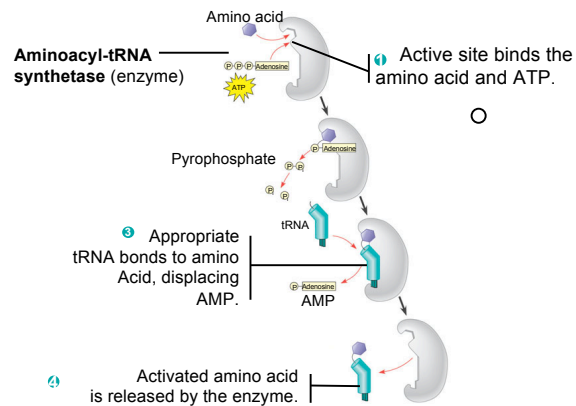
Elongation



Termination

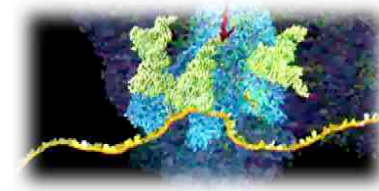


An aminoacyl-tRNA synthetase joins a specific amino acid to a tRNA



See the Animation

- www.dnai.org



Inhibition of protein synthesis

Toxin	Mode of action	Target
Puromycin	forms peptidyl-puromycin, prevents translocation	Prokaryotes
Tetracycline	blocks the A-site, prevents binding of aminoacyl tRNAs	Prokaryotes
Chloramphenicol	blocks peptidyl transfer	Prokaryotes
Cycloheximide	blocks peptidyl transferase	Eucaryotes
Streptomycin	inhibits initiation at high concentrations	Prokaryotes
Diphtheria toxin	catalyzes ADP-ribosylation of residue in eEF2	Eucaryotes
Erythromycin	binds to 50S subunit, inhibits translocation	Prokaryotes
Ricin	inactivates 60S subunit, depurinates an adenosine in 23S rRNA	Eucaryotes

NOTE: Prokaryotes (this generally includes protein synthesis in mitochondria and chloroplasts)

Summary of transcription and translation in a eukaryotic cell

