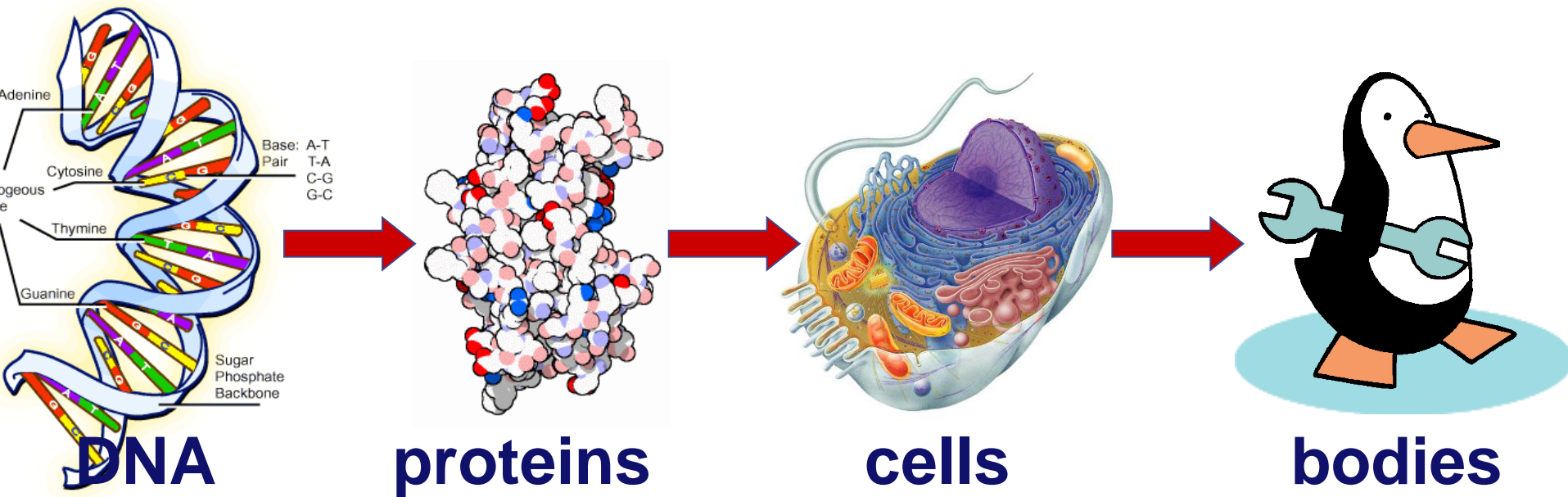
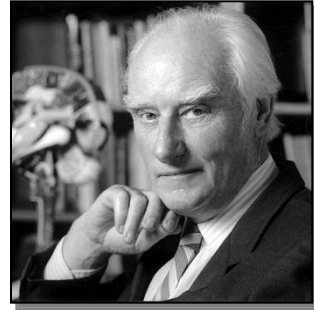


What do genes code for?

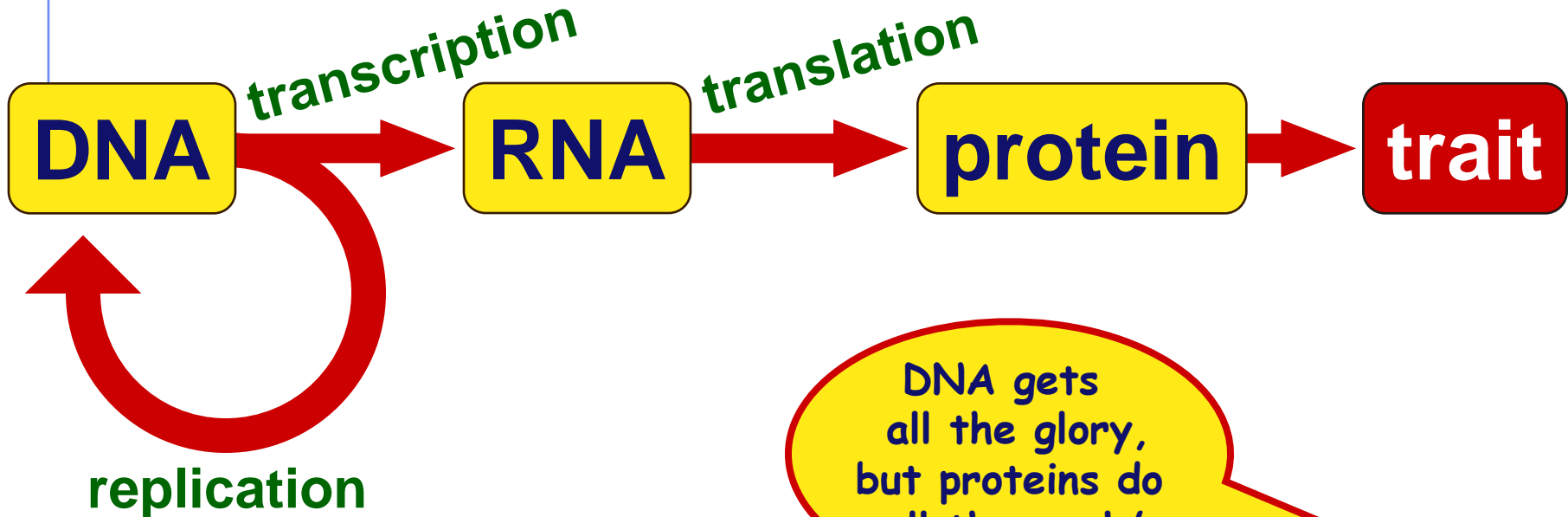
- How does DNA code for cells & bodies?
 - ◆ how are cells and bodies made from the instructions in DNA



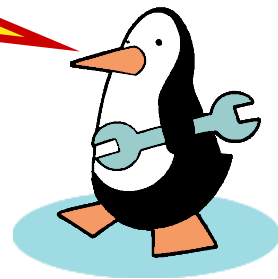
The “Central Dogma”



- Flow of genetic information in a cell
 - ◆ How do we move information from DNA to proteins?



DNA gets all the glory, but proteins do all the work!

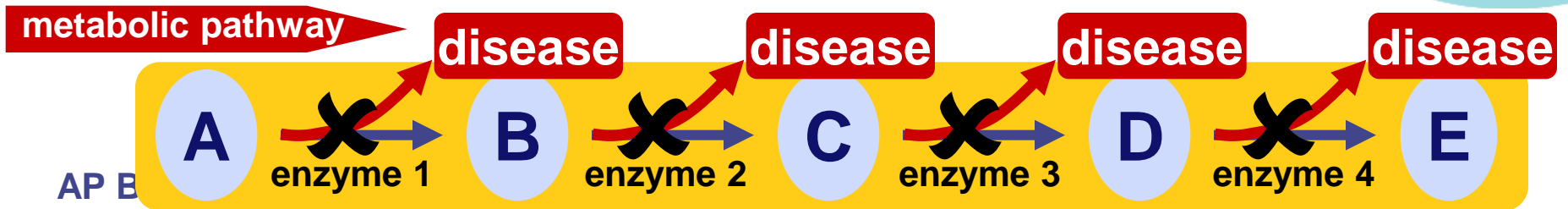
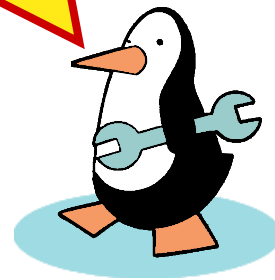


Metabolism taught us about genes

■ Inheritance of metabolic diseases

- ◆ suggested that genes coded for enzymes
- ◆ each disease (phenotype) is caused by non-functional gene product
 - lack of an enzyme
 - Tay Sachs
 - PKU (phenylketonuria)
 - albinism

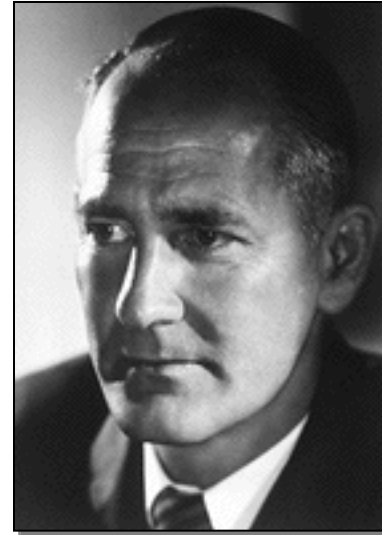
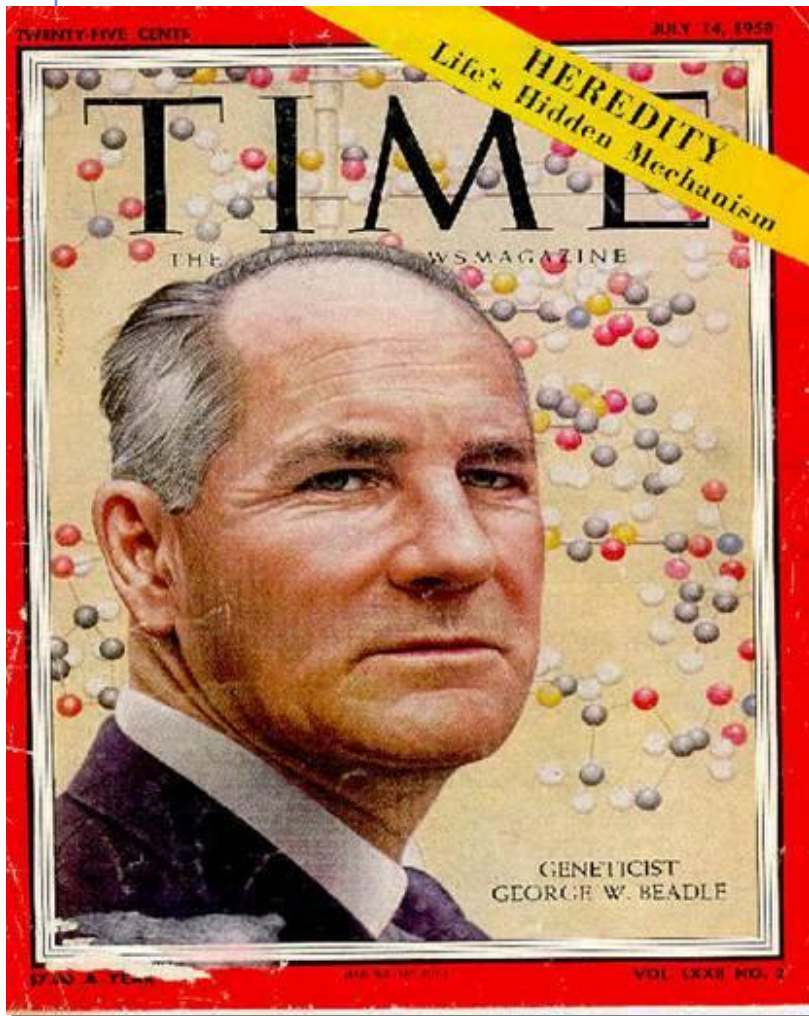
Am I just the sum of my proteins?



1941 | 1958

Beadle & Tatum

one gene : one enzyme hypothesis



George Beadle



Edward Tatum

"for their discovery that genes act by regulating definite chemical events"

Beadle & Tatum

Wild-type
Neurospora

Minimal
medium

X rays or ultraviolet light

create mutations

asexual
spores

Growth on
complete
medium

positive control

spores

Select one of
the spores

Test on minimal
medium to confirm
presence of mutation

negative control

Grow on
complete medium

Minimal media supplemented only with...

experimentals

Pyridoxine

amino acid
supplements

Benzoic acid

Choline

Inositol

Nucleic
acid

Folic
acid

Arginine

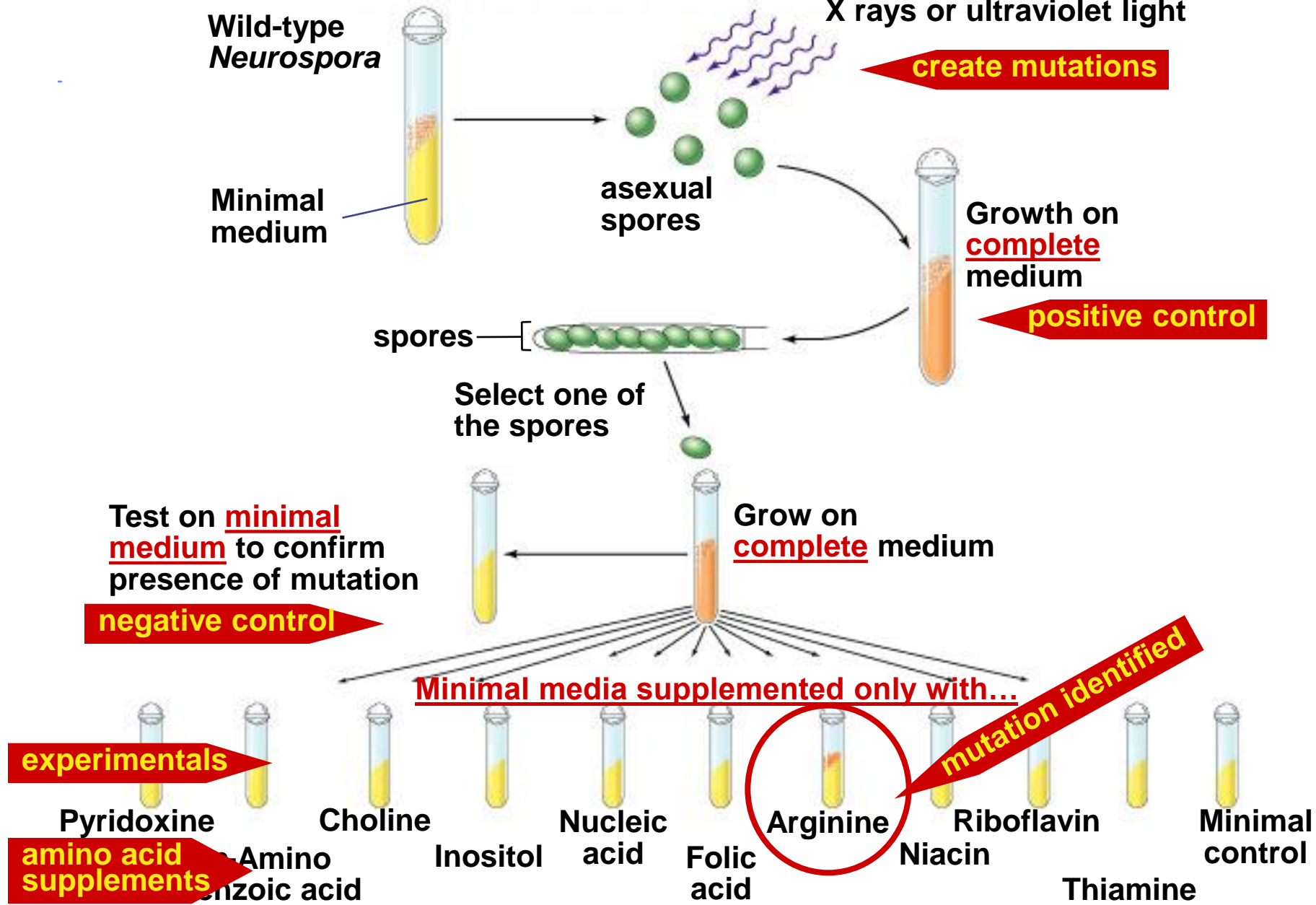
Niacin

mutation identified

Riboflavin

Thiamine

Minimal
control



From gene to protein

nucleus

cytoplasm

DNA

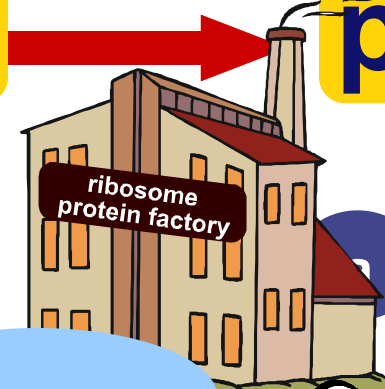
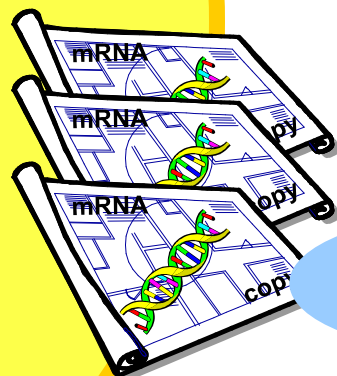
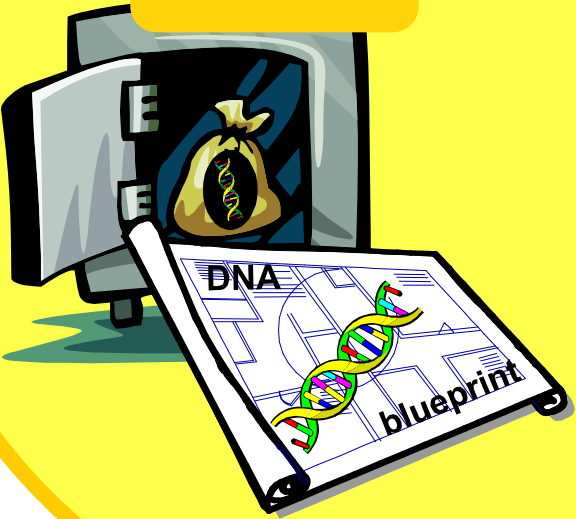
transcription

mRNA

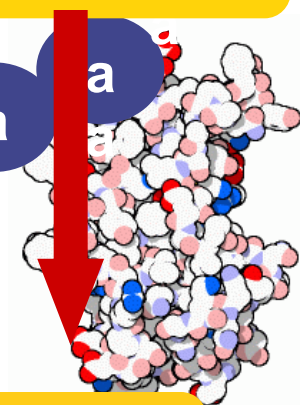
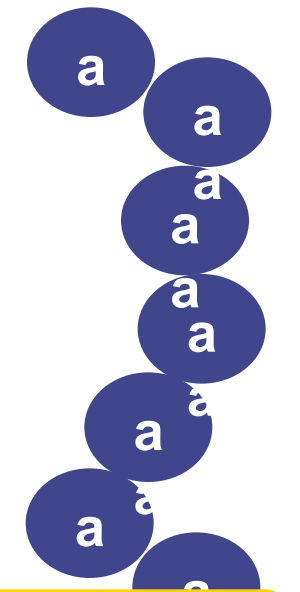
translation

protein

trait

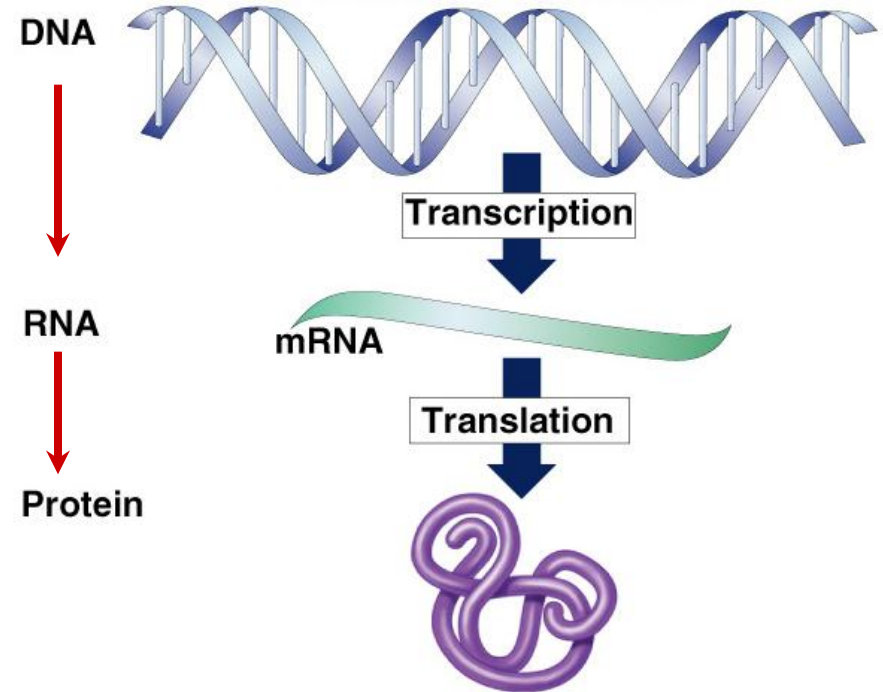


ribosome



Transcription

from
DNA nucleic acid language
to
RNA nucleic acid language



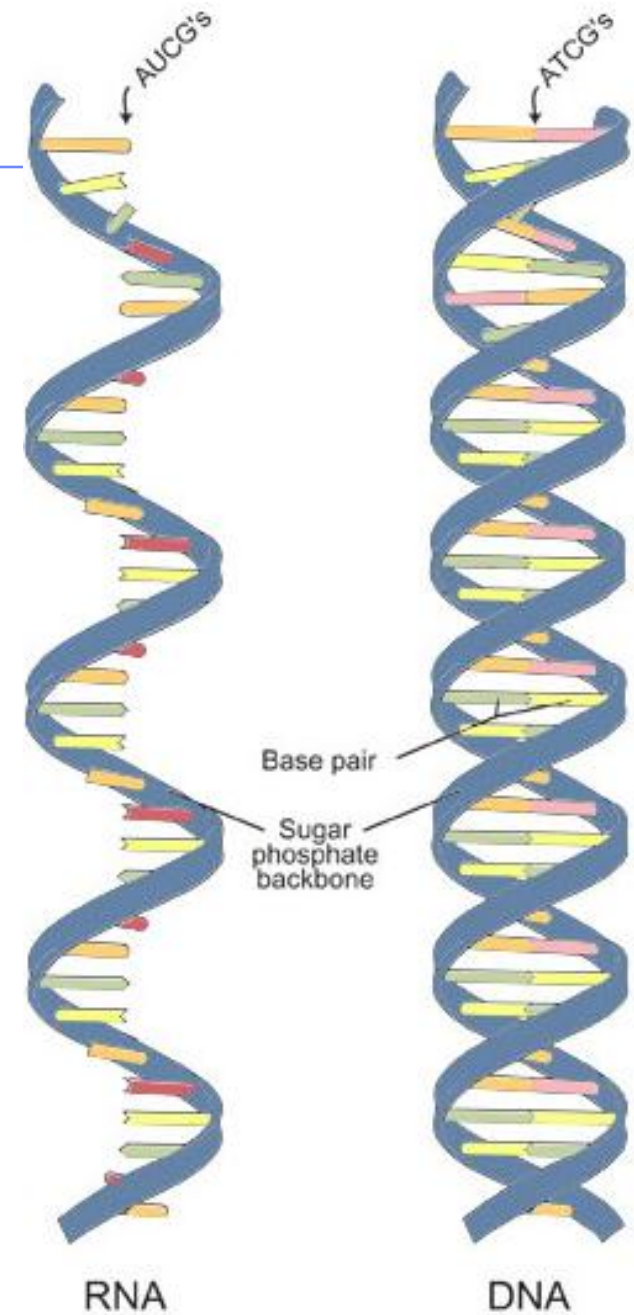
RNA

- ribose sugar
- N-bases
 - ◆ uracil instead of thymine
 - ◆ U : A
 - ◆ C : G
- single stranded
- lots of RNAs
 - ◆ mRNA, tRNA, rRNA, siRNA...

DNA

transcription

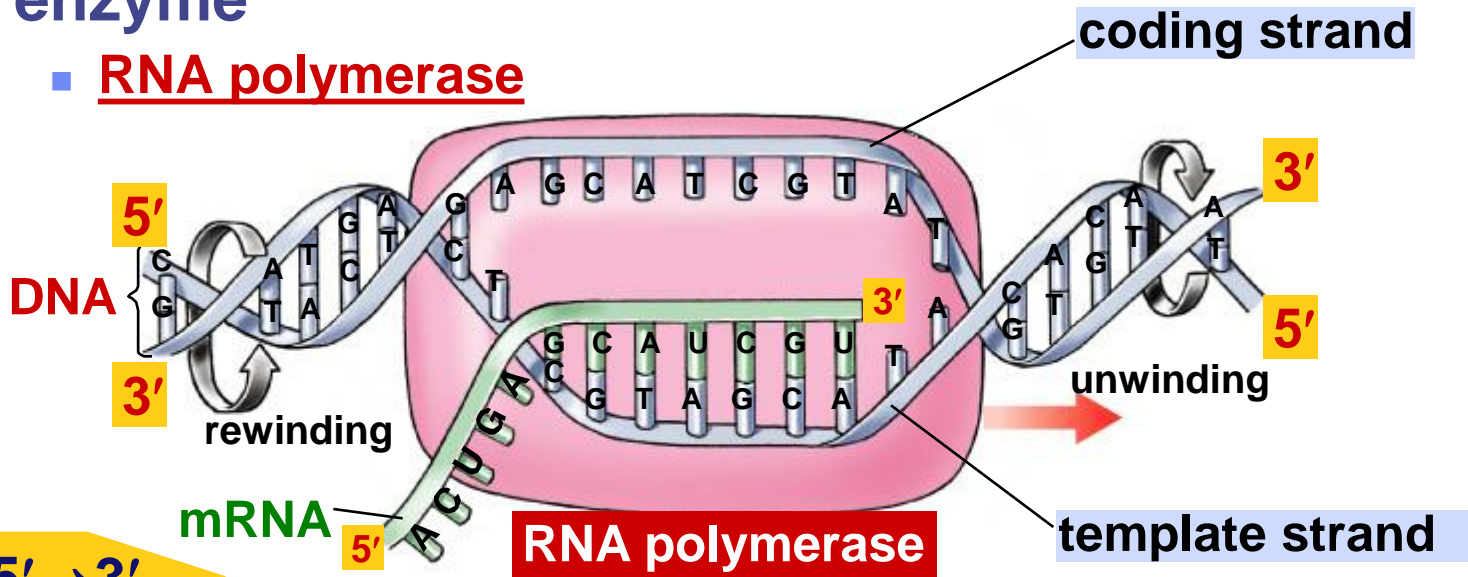
RNA



Transcription

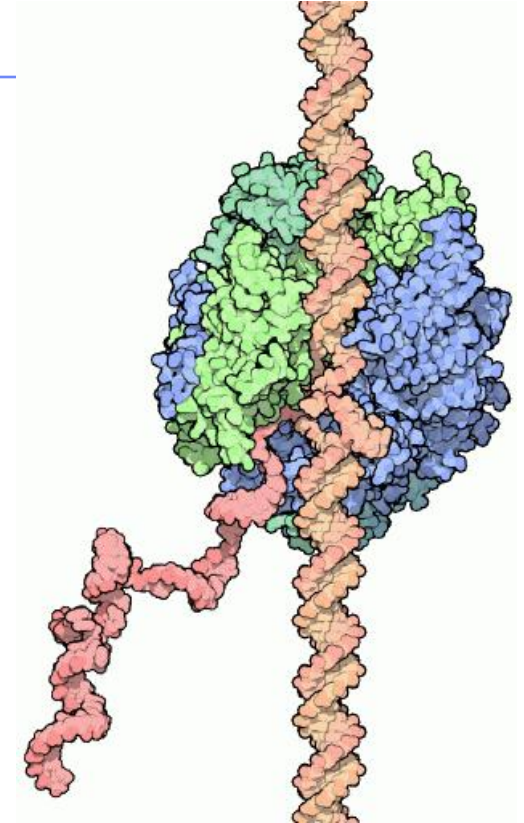
■ Making mRNA

- ◆ transcribed DNA strand = template strand
- ◆ untranscribed DNA strand = coding strand
 - same sequence as RNA
- ◆ synthesis of complementary RNA strand
 - transcription bubble
- ◆ enzyme
 - RNA polymerase



RNA polymerases

- 3 RNA polymerase enzymes
 - ◆ RNA polymerase 1
 - only transcribes rRNA genes
 - makes ribosomes
 - ◆ RNA polymerase 2
 - transcribes genes into mRNA
 - ◆ RNA polymerase 3
 - only transcribes tRNA genes
 - ◆ each has a specific promoter sequence it recognizes



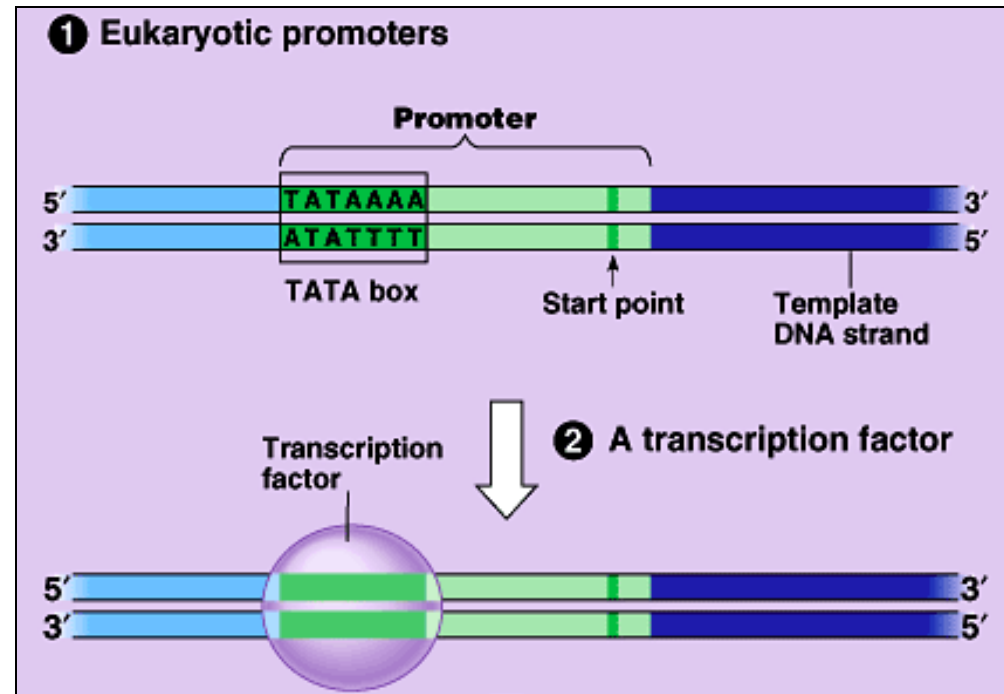
Which gene is read?

■ Promoter region

- ◆ binding site before beginning of gene
- ◆ TATA box binding site
- ◆ binding site for RNA polymerase & transcription factors

■ Enhancer region

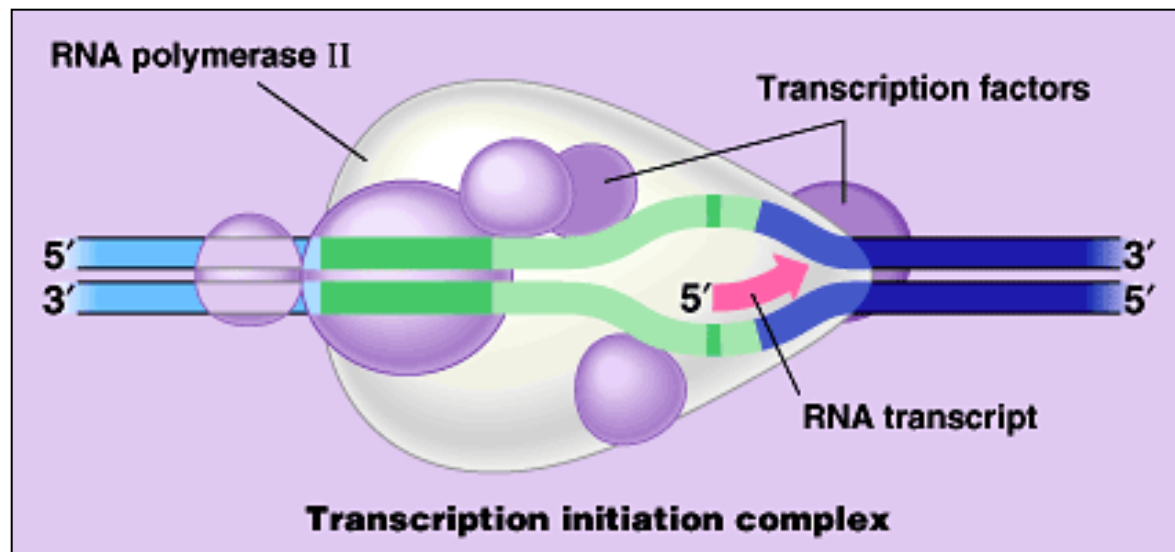
- ◆ binding site far upstream of gene
 - turns transcription on HIGH



Transcription Factors

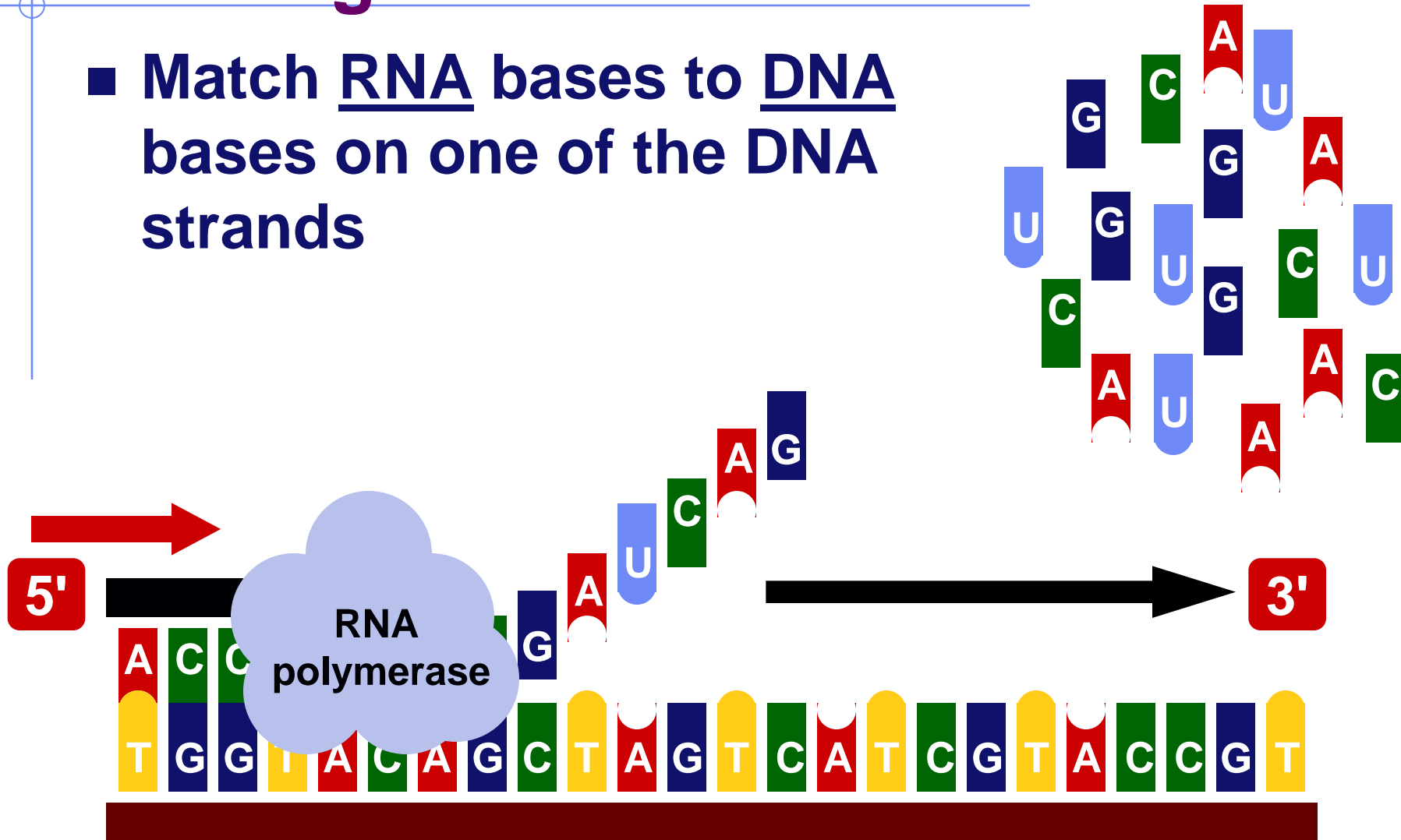
■ Initiation complex

- ◆ transcription factors bind to promoter region
 - suite of proteins which bind to DNA
 - hormones?
 - turn on or off transcription
- ◆ trigger the binding of RNA polymerase to DNA



Matching bases of DNA & RNA

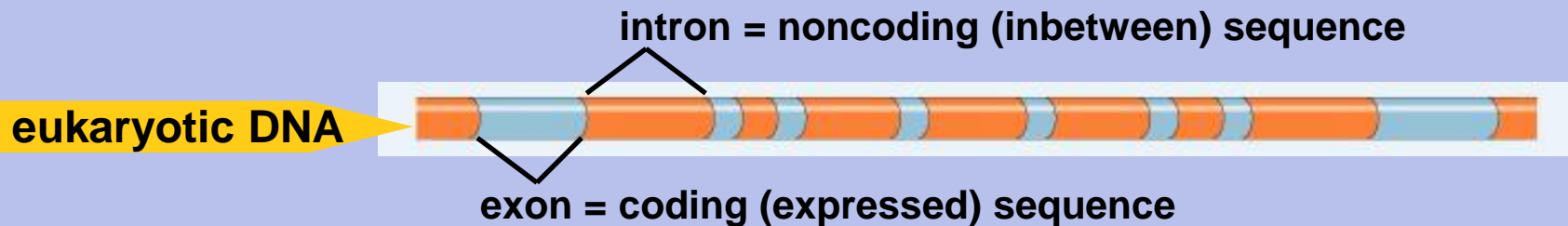
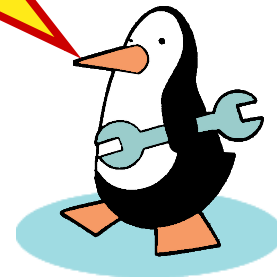
- Match RNA bases to DNA bases on one of the DNA strands



Eukaryotic genes have junk!

- Eukaryotic genes are not continuous
 - ◆ **exons** = the real gene
 - expressed / coding DNA
 - ◆ **introns** = the junk
 - inbetween sequence

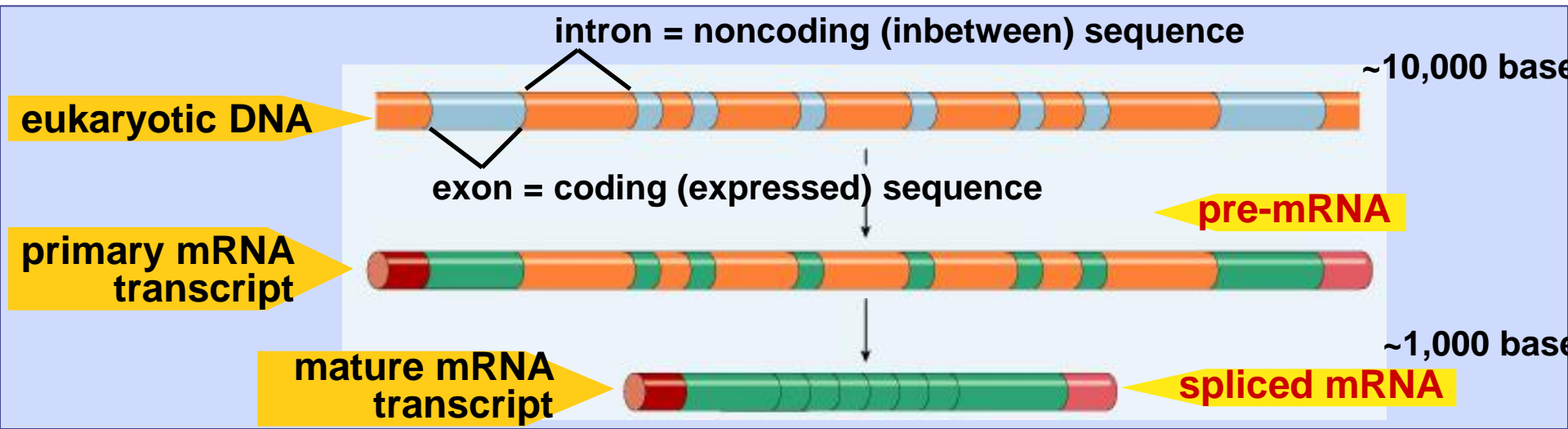
introns
come out!



mRNA splicing

■ Post-transcriptional processing

- ◆ eukaryotic mRNA needs work after transcription
- ◆ primary transcript = pre-mRNA
- ◆ mRNA splicing
 - edit out introns
- ◆ make mature mRNA transcript



Discovery of exons/introns

1977 | 1993



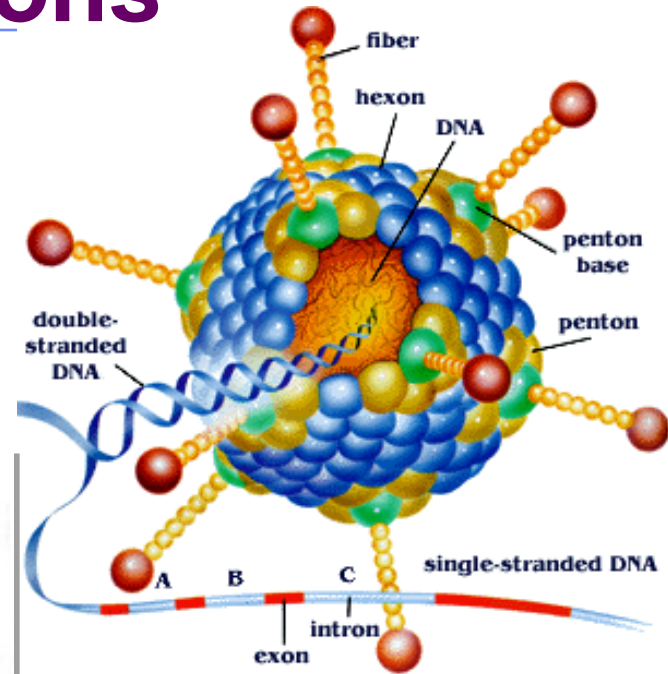
Richard Roberts

CSHL

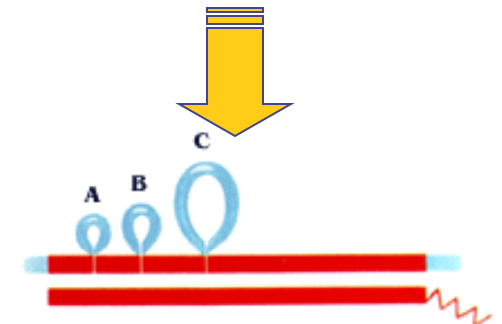
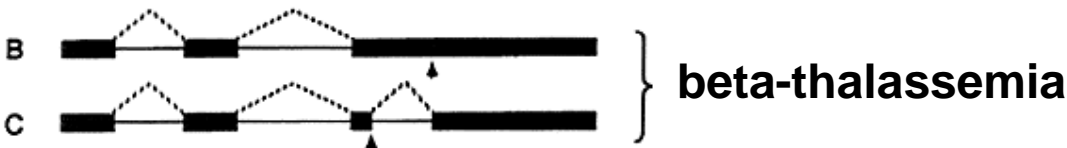


Philip Sharp

MIT



adenovirus
common cold

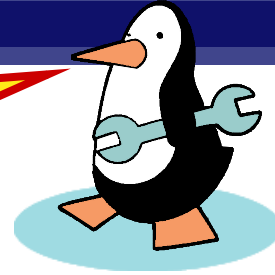


Splicing must be accurate

- No room for mistakes!
 - ◆ a single base added or lost throws off the reading frame

AUGCGGCTATGGGUCCGAUAAGGGCCA
AUGCGGUCCGAUAAGGGCCA
AUG|CGG|UCC|GAU|AAG|GGC|CAU
Met|Arg|Ser|Asp|Lys|Gly|His

AUGCGGCTATGGGUCCGAUAAGGGCCA
AUGCGGGUCCGAUAAGGGCCA
AUG|CGG|GUC|CGA|UAA|GGG|CCA|U
Met|Arg|Val|Arg|STOP|



Whoa! I think we just broke a biological "rule"!

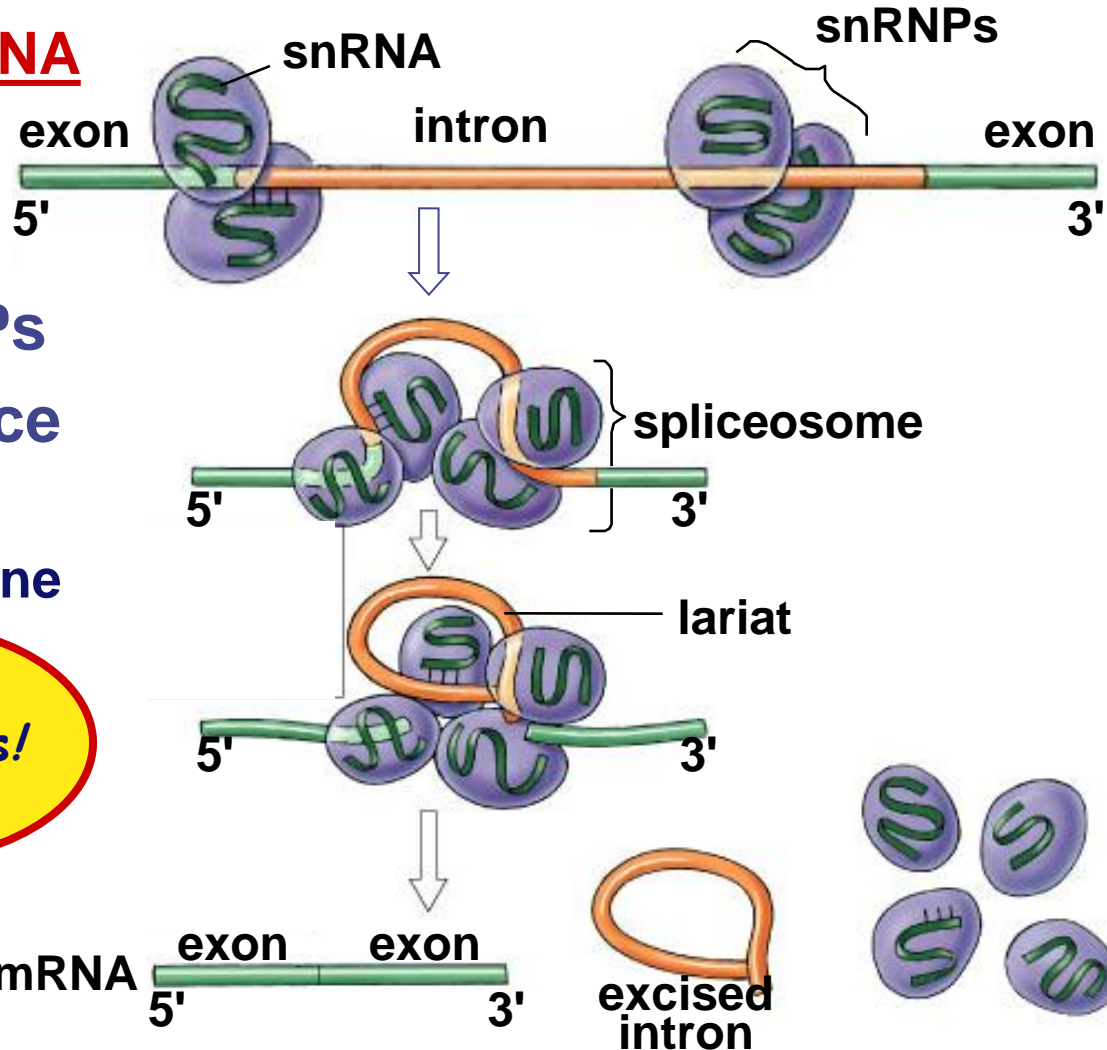
RNA splicing enzymes

■ snRNPs

- ◆ small nuclear RNA
- ◆ proteins

■ Spliceosome

- ◆ several snRNPs
- ◆ recognize splice site sequence
 - cut & paste gene



No, not smurfs! "snurps"

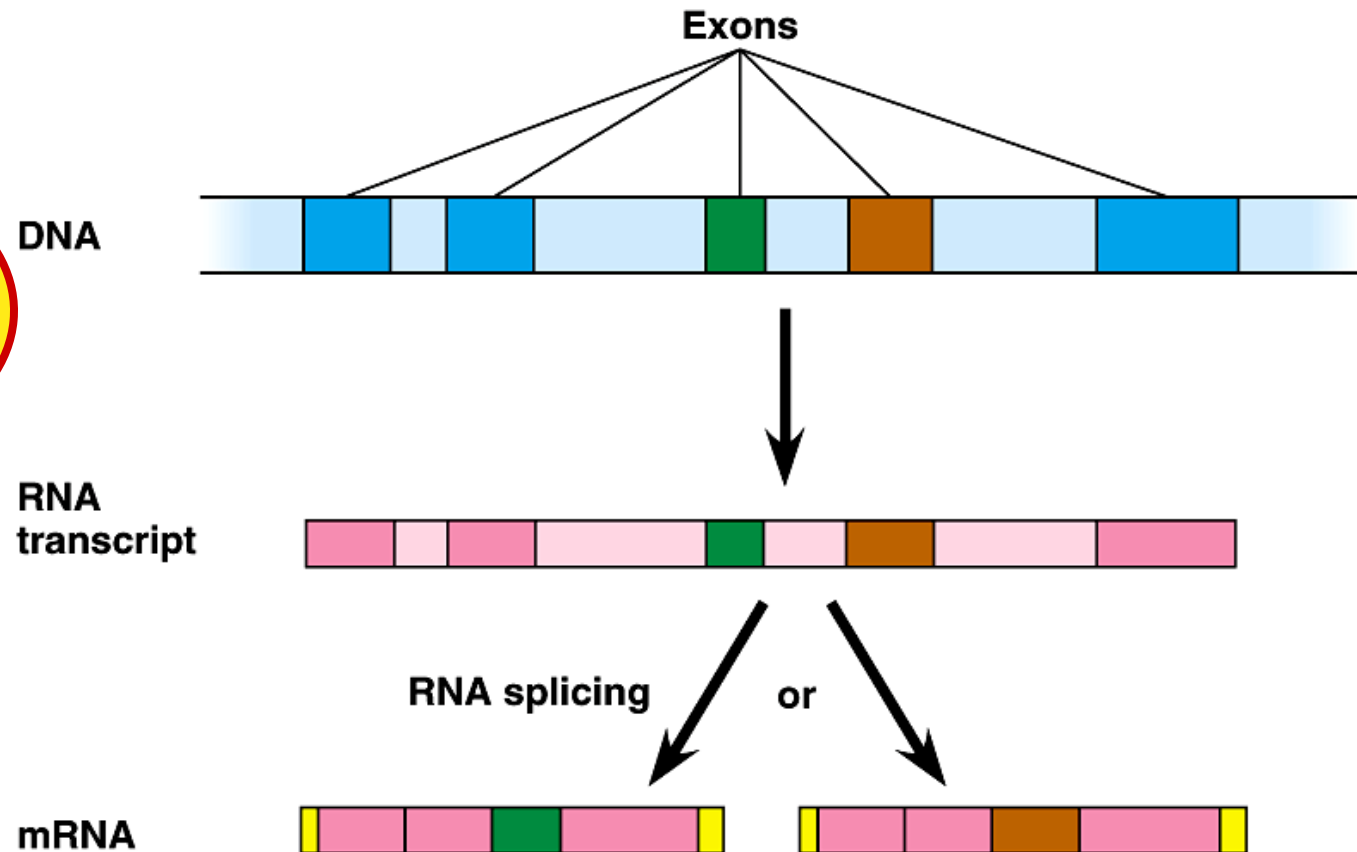
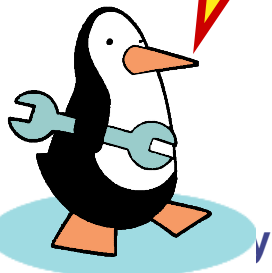
mRNA Splicing

HHMI

Alternative splicing

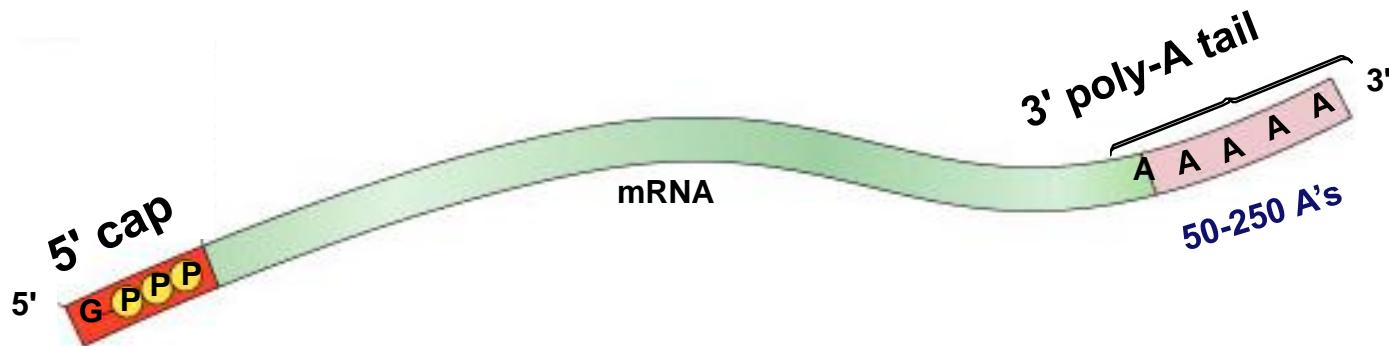
- Alternative mRNAs produced from same gene
 - ◆ when is an intron not an intron...
 - ◆ different segments treated as exons

Starting to get hard to define a gene!



More post-transcriptional processing

- Need to protect mRNA on its trip from nucleus to cytoplasm
 - ◆ enzymes in cytoplasm attack mRNA
 - protect the ends of the molecule
 - add 5' GTP cap
 - add poly-A tail
 - ◆ longer tail, mRNA lasts longer: produces more protein

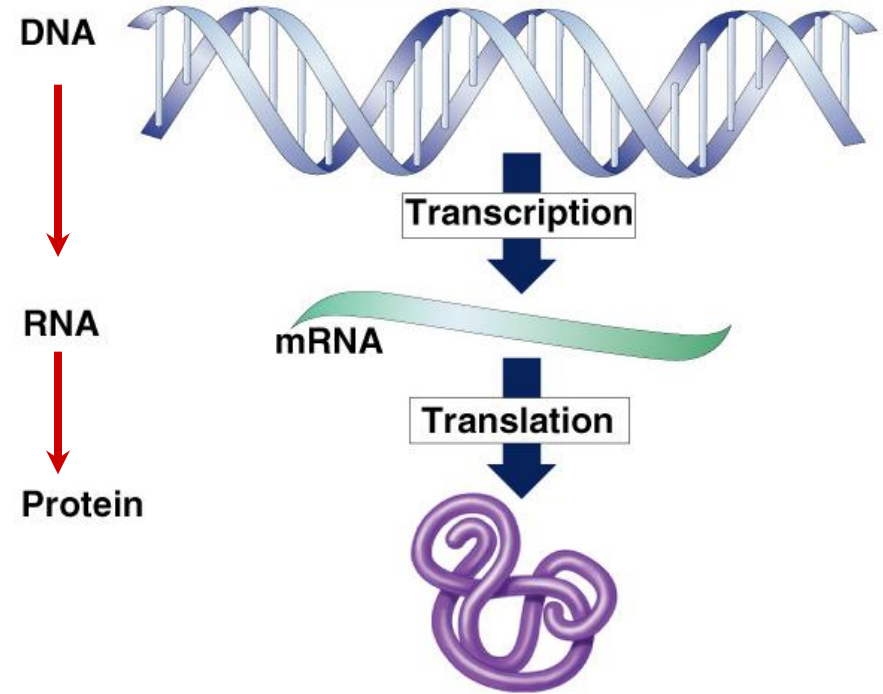


Transcription Review

HHMI

Translation

from
nucleic acid language
to
amino acid language



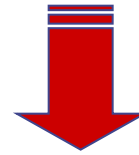
How does mRNA code for proteins?

DNA

TACGCACATTTACGTACGCGG

4

ATCG



mRNA

UGCUGUAAAUGCAUGCGCC

4

AUCG



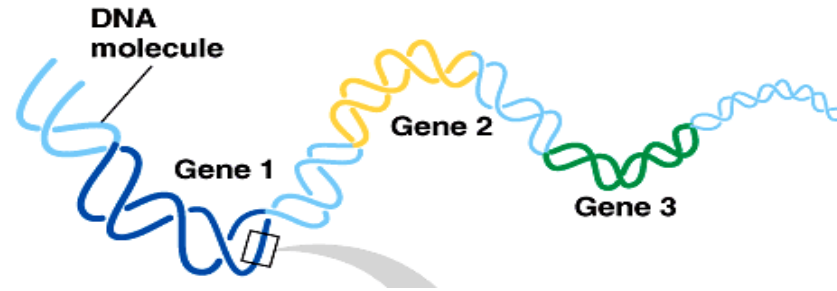
protein

Met Arg Val Asn Ala Cys Ala

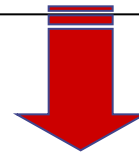
20

How can you code for 20 amino acids with only 4 nucleotide bases (A,U,G,C)?

mRNA codes for proteins in triplets

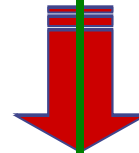


DNA TACGCACATTTACGTACGCGG



codon

mRNA UGCGUGUAAAUGCAUGCGCC



protein Met Arg Val Asn Ala Cys Ala

1960 | 1968

Nirenberg & Khorana

Cracking the code

■ Crick

- ◆ determined 3-letter (triplet) codon system

WHYDIDTHEREDBATEATTHEFATRAT

■ Nirenberg (47) & Khorana (17)

- ◆ determined mRNA–amino acid match
- ◆ added fabricated mRNA to test tube of ribosomes, tRNA & amino acids
 - created artificial UUUUU... mRNA
 - found that UUU coded for phenylalanine

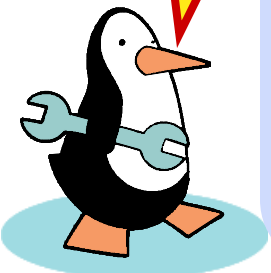
The code

- Code for ALL life!
 - ◆ strongest support for a common origin for all life
- Code is redundant
 - ◆ several codons for each amino acid
 - ◆ 3rd base “wobble”

Why is the wobble good?

- Start codon
 - ◆ AUG
 - ◆ methionine
- Stop codons
 - ◆ UGA, UAA, UAG

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



Triplet Code

HHMI

How are the codons matched to amino acids?

DNA TACGCACATTACGTACGCGG

mRNA UGCGUGUA^{3'}AAUGCAUGCGCC^{5'}

codon

UAC

GCA

CAU

anti-codon

Met

Arg

Val

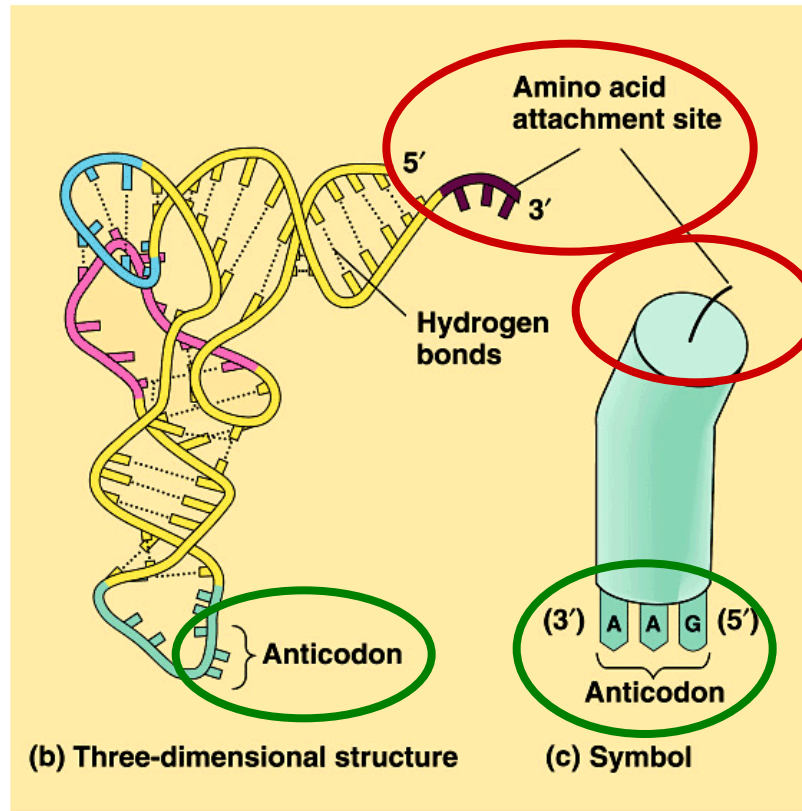
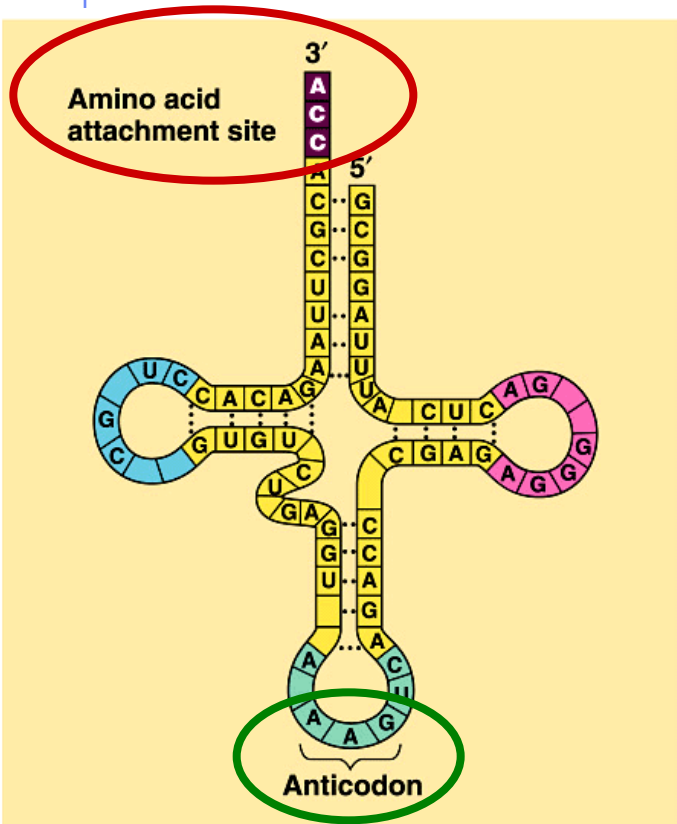
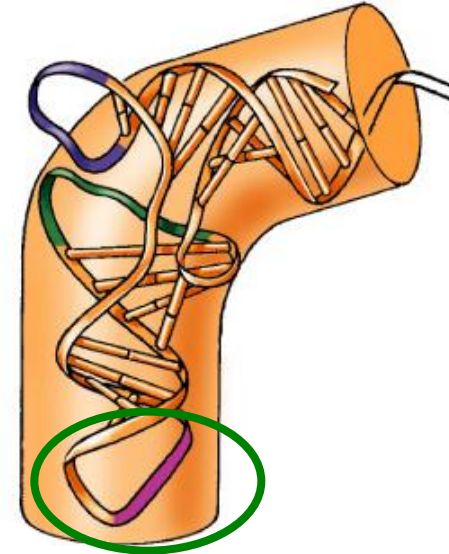
tRNA

amino acid

Transfer RNA structure

■ “Clover leaf” structure

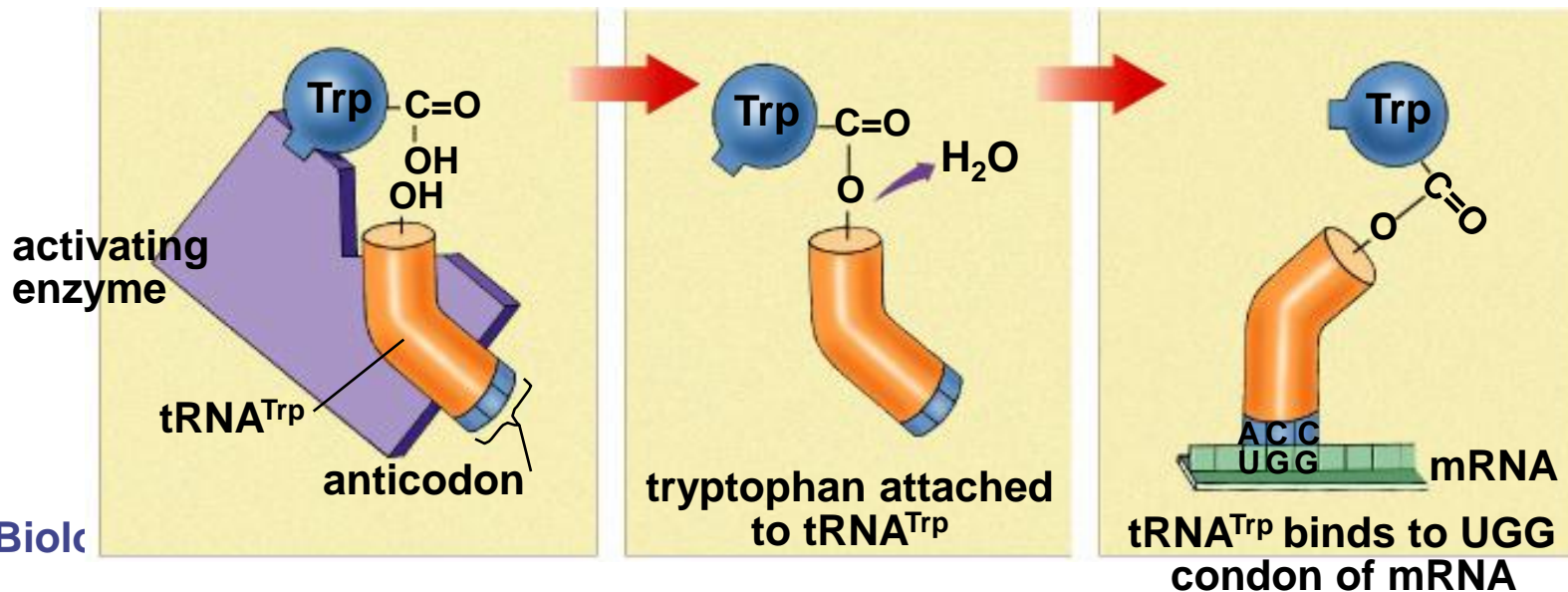
- ◆ anticodon on “clover leaf” end
- ◆ amino acid attached on 3' end



Loading tRNA

■ Aminoacyl tRNA synthetase

- ◆ enzyme which bonds amino acid to tRNA
- ◆ bond requires energy
 - ATP → AMP
 - bond is unstable
 - so it can release amino acid at ribosome easily

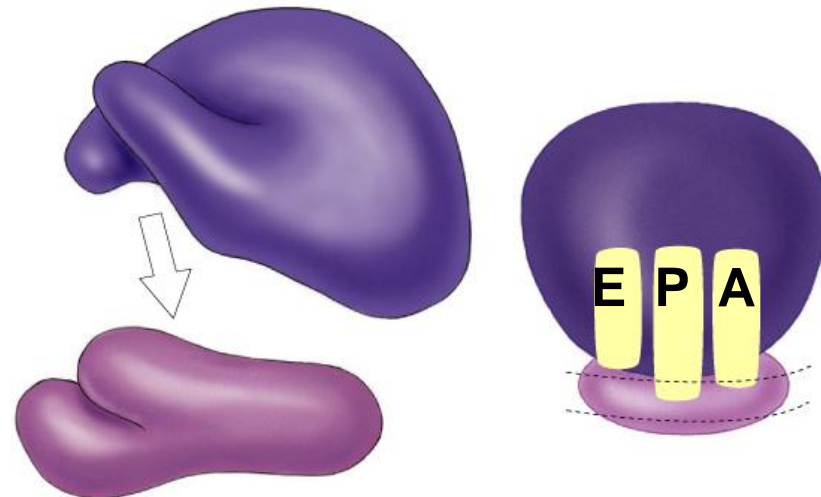
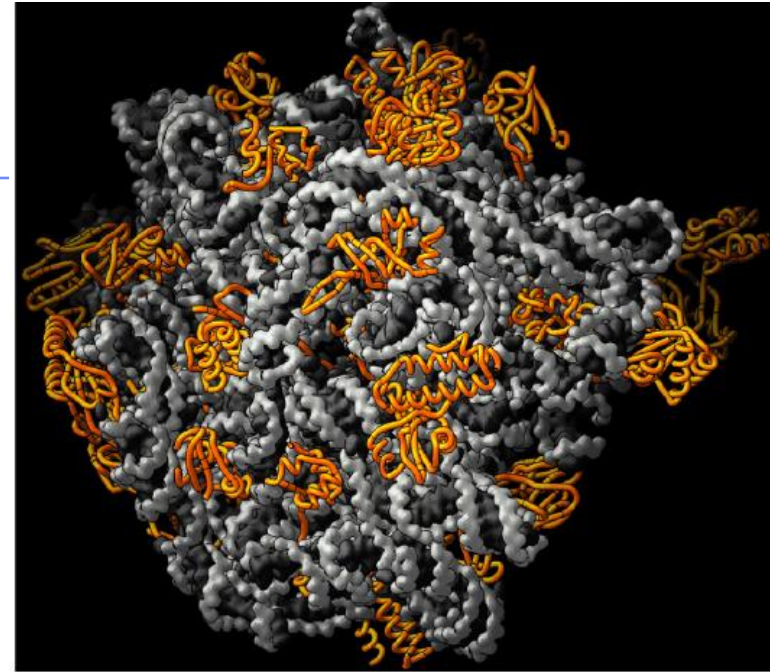


Ribosomes

- Facilitate coupling of tRNA anticodon to mRNA codon
 - ◆ organelle or enzyme?

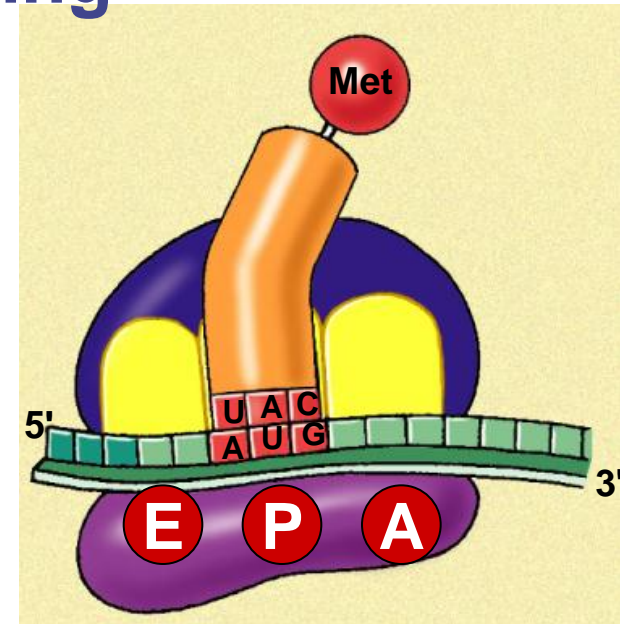
- Structure

- ◆ ribosomal RNA (rRNA) & proteins
- ◆ 2 subunits
 - large
 - small



Ribosomes

- **A site** (aminoacyl-tRNA site)
 - ◆ holds tRNA carrying next **amino acid** to be added to chain
- **P site** (peptidyl-tRNA site)
 - ◆ holds tRNA carrying growing **polypeptide** chain
- **E site** (exit site)
 - ◆ **empty** tRNA leaves ribosome from **exit** site



Building a polypeptide

■ Initiation

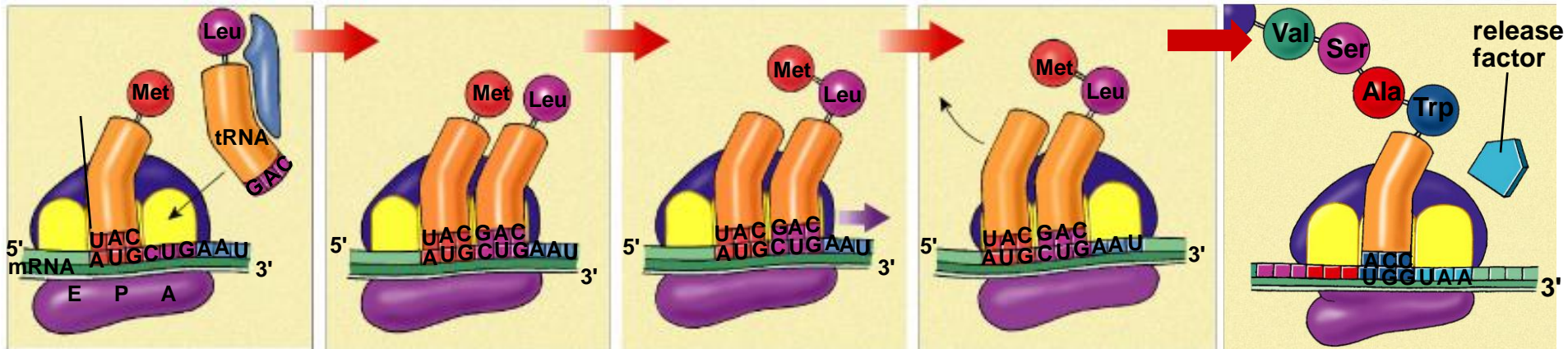
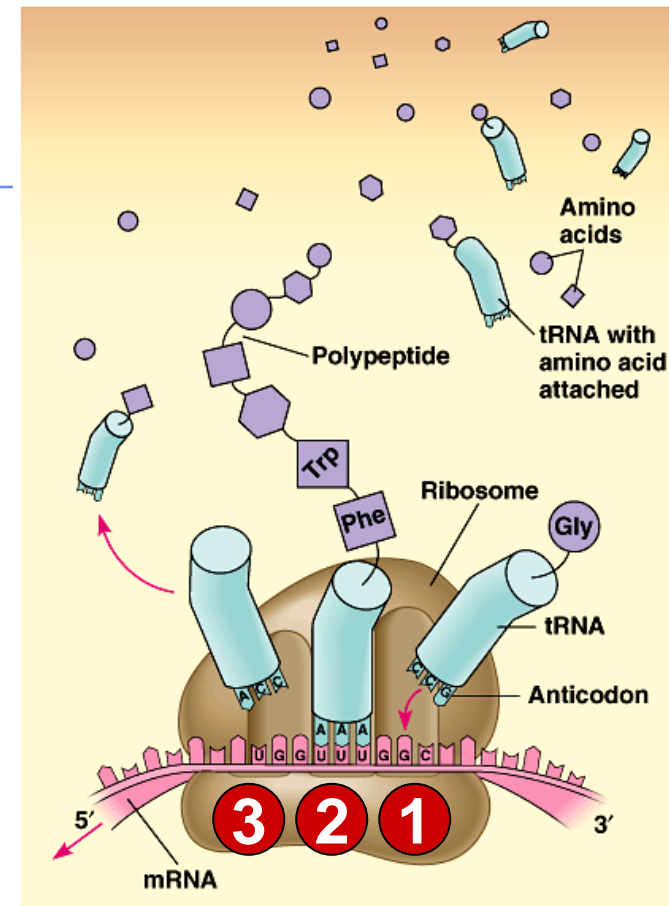
- ◆ brings together mRNA, ribosome subunits, initiator tRNA

■ Elongation

- ◆ adding amino acids based on codon sequence

■ Termination

- ◆ end codon



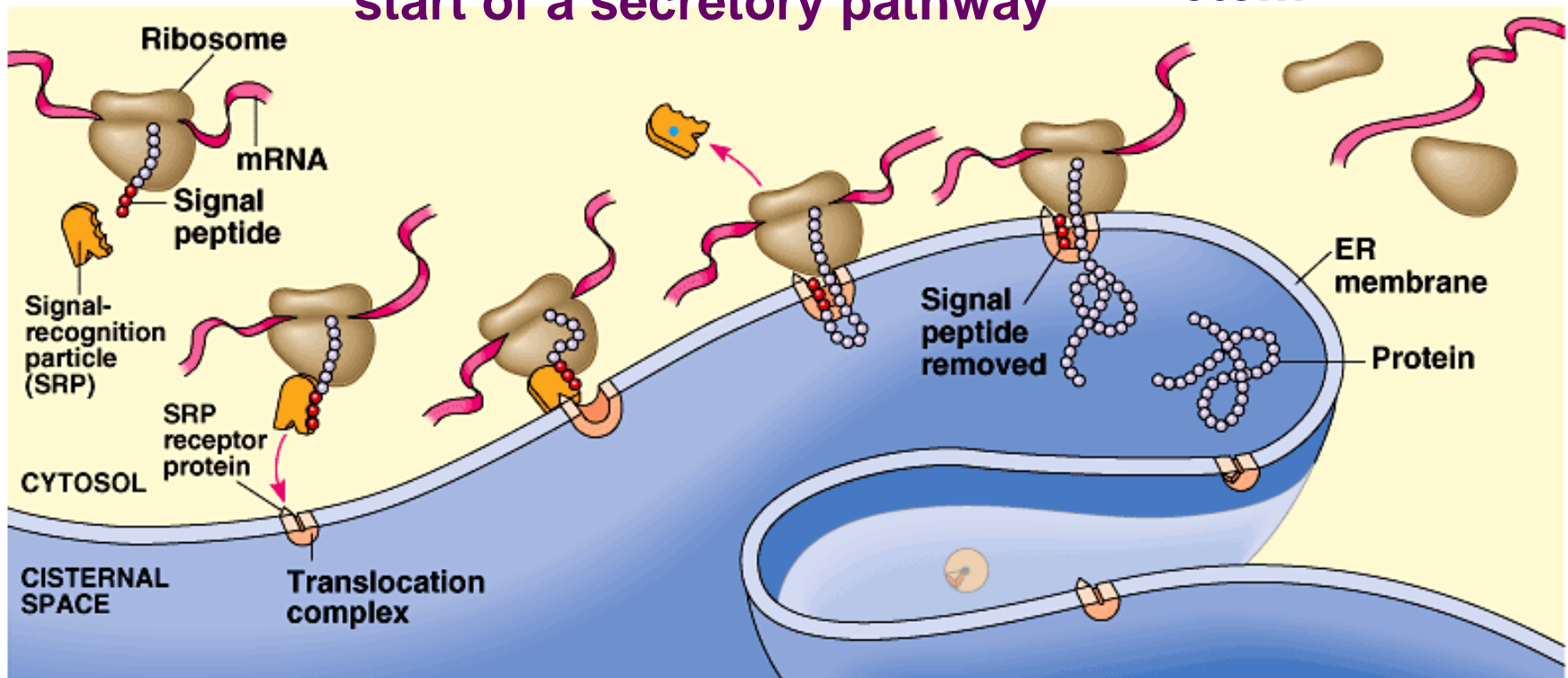
Protein targeting

- **Signal peptide**
 - ◆ address label

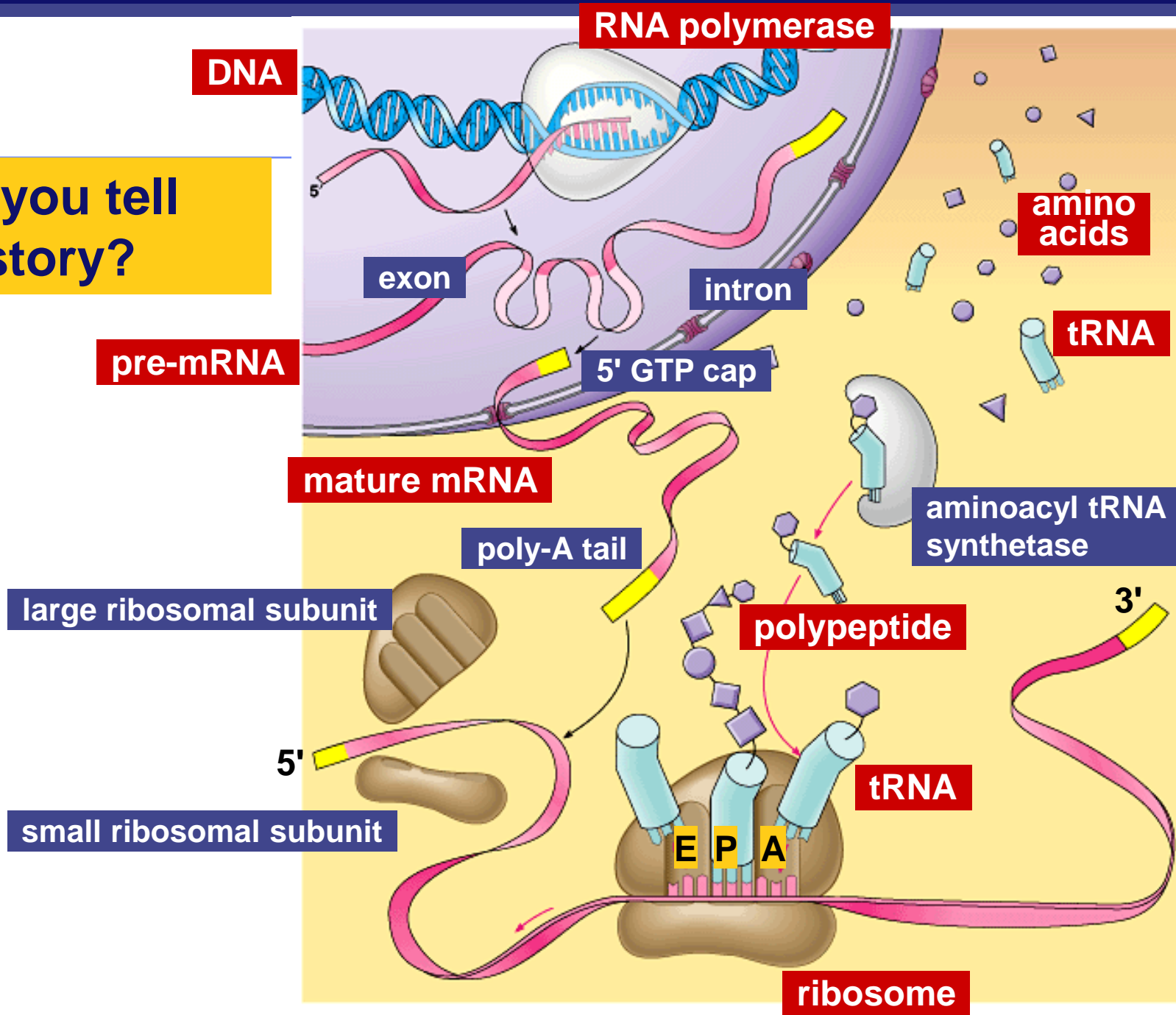
Destinations:

- secretion
- nucleus
- mitochondria
- chloroplasts
- cell membrane
- cytoplasm
- etc...

start of a secretory pathway



Can you tell the story?



Protein Synthesis in Prokaryotes

Psssst...
no nucleus!

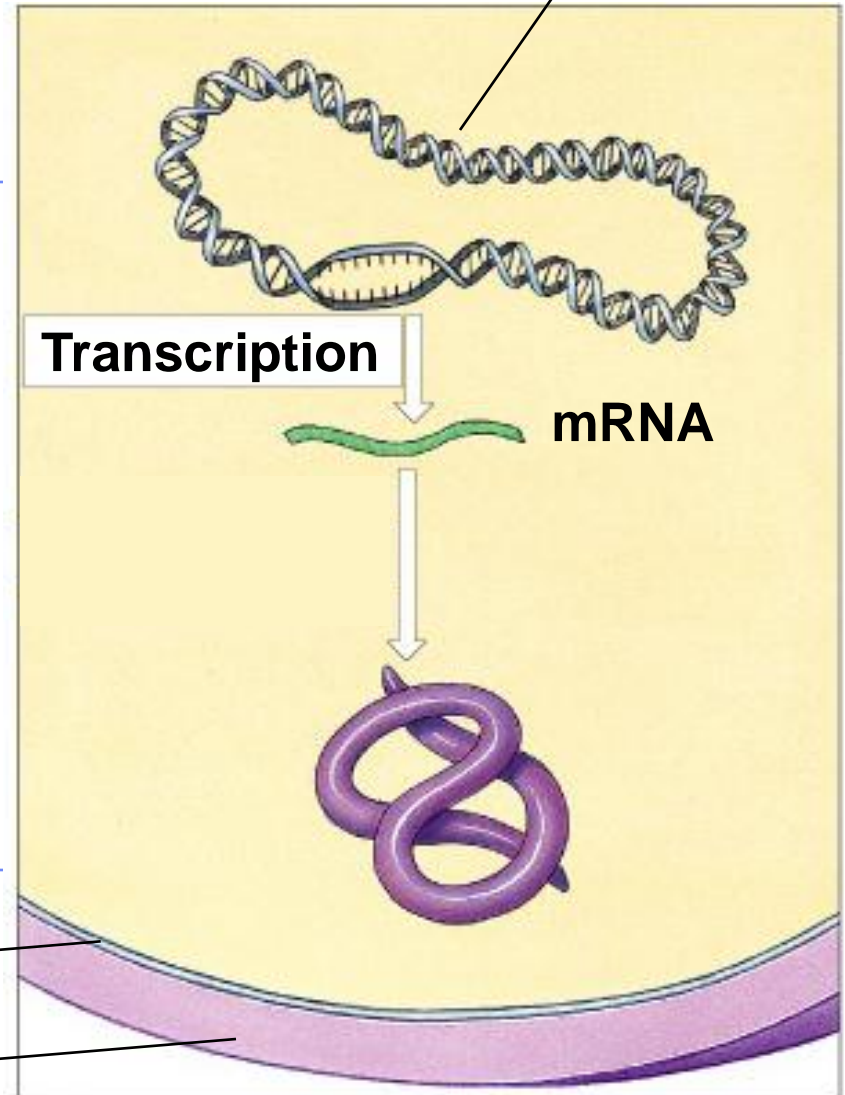
Cell membrane

Cell wall

Bacterial chromosome

Transcription

mRNA



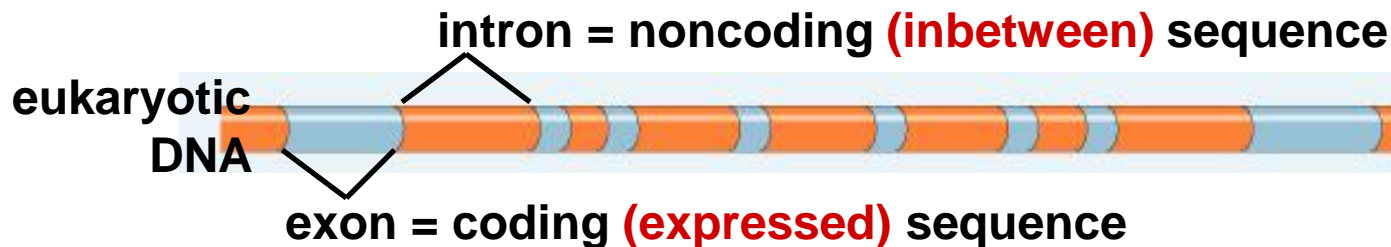
Prokaryote vs. Eukaryote genes

■ Prokaryotes

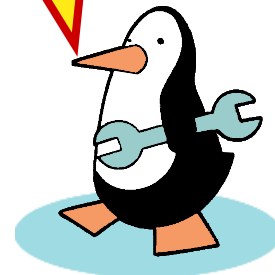
- ◆ DNA in cytoplasm
- ◆ circular chromosome
- ◆ naked DNA
- ◆ no introns

■ Eukaryotes

- ◆ DNA in nucleus
- ◆ linear chromosomes
- ◆ DNA wound on histone proteins
- ◆ introns vs. exons

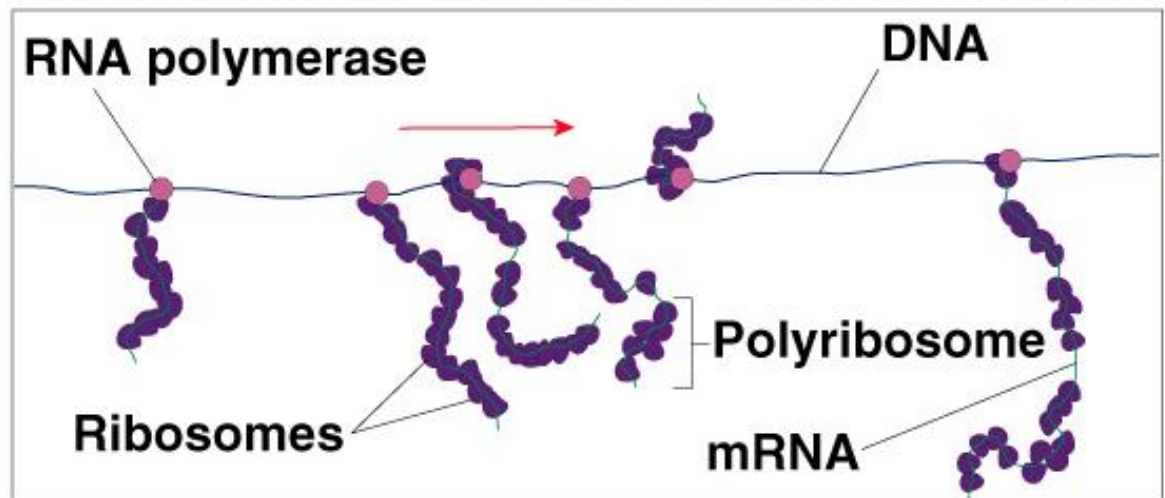
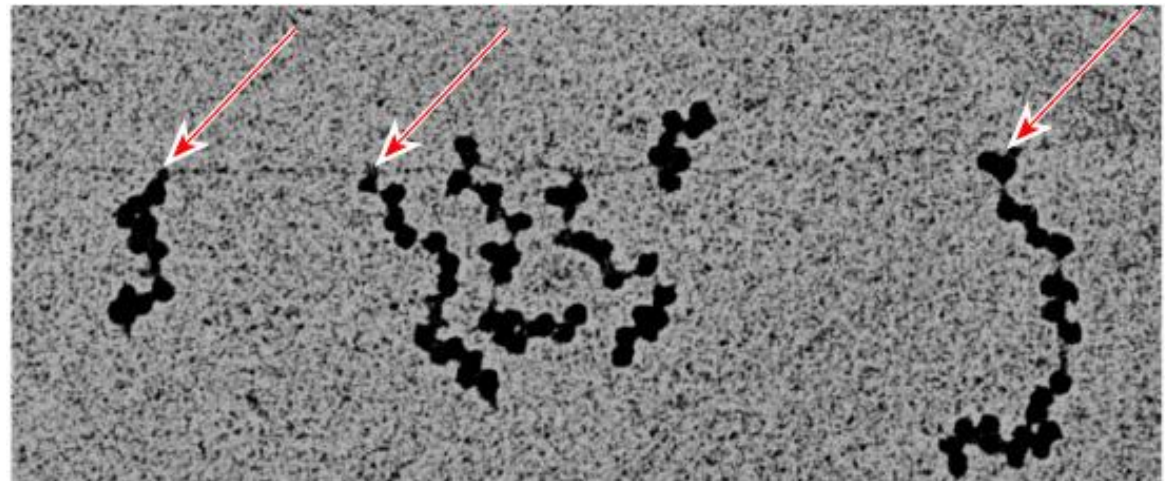


introns
come out!



Translation in Prokaryotes

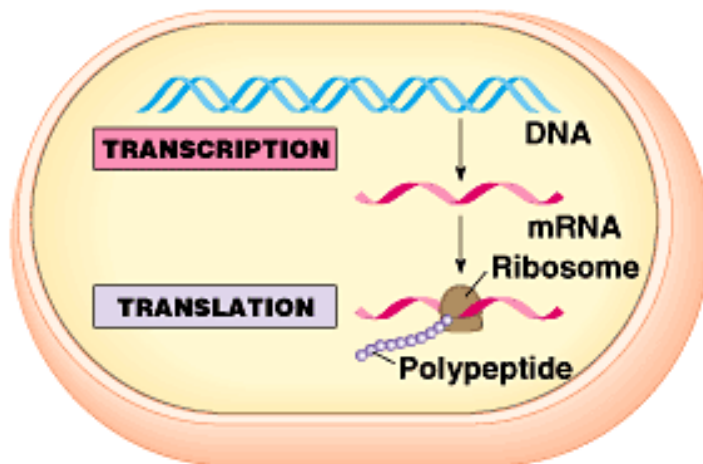
- Transcription & translation are simultaneous in bacteria
 - ◆ DNA is in cytoplasm
 - ◆ no mRNA editing
 - ◆ **ribosomes** read mRNA as it is being transcribed



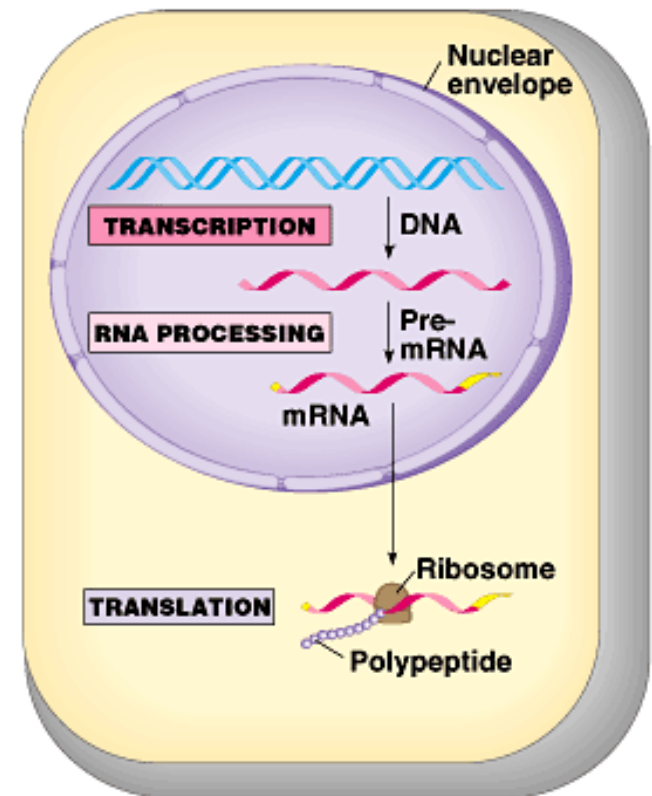
Translation: prokaryotes vs. eukaryotes

■ Differences between prokaryotes & eukaryotes

- ◆ time & physical separation between processes
 - takes eukaryote ~1 hour from DNA to protein
- ◆ no RNA processing



(a) Prokaryotic cell



(b) Eukaryotic cell

Any Questions??

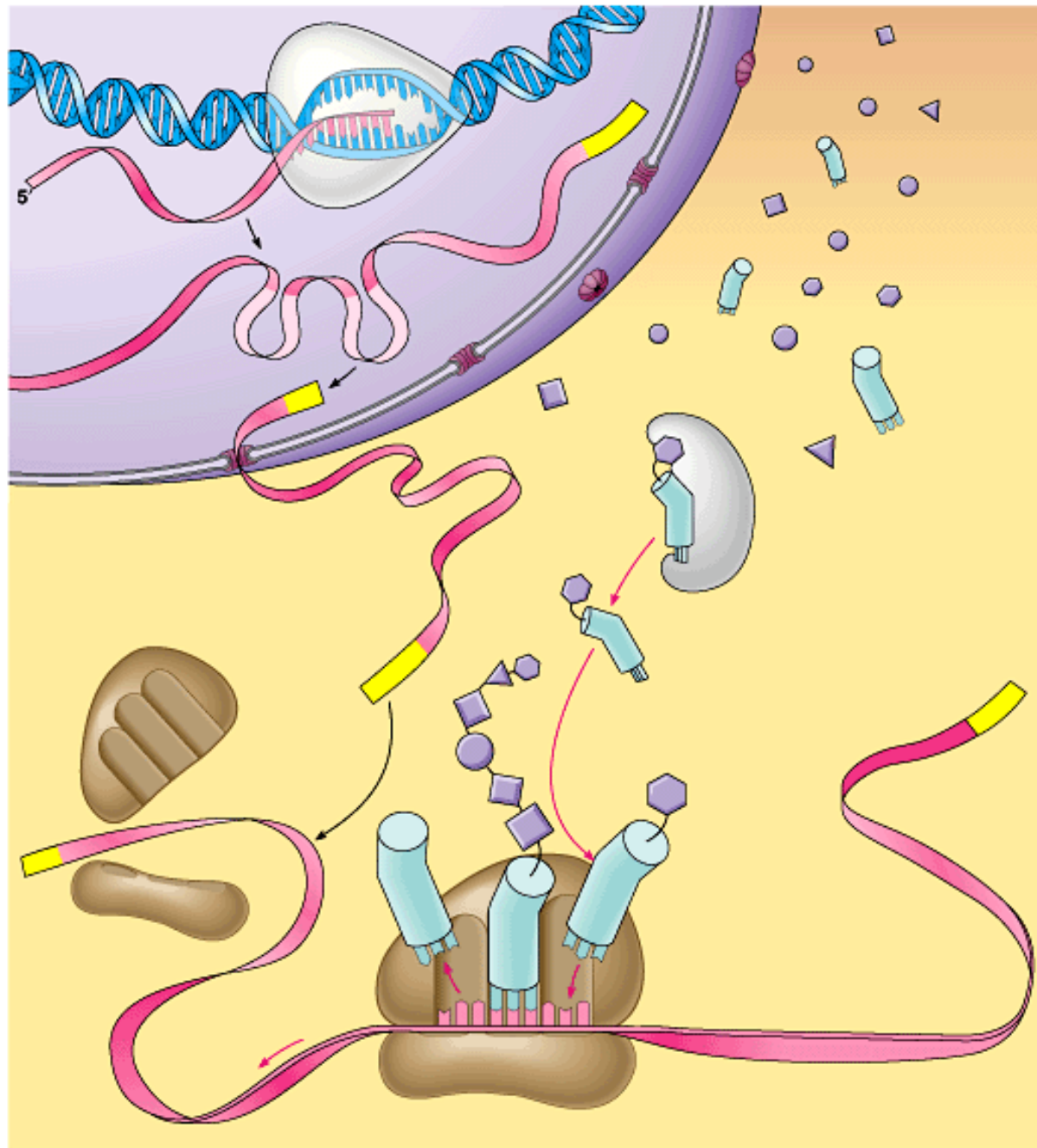
**What color would a smurf turn
if he held his breath?**



A decorative graphic consisting of a horizontal blue line at the top, a vertical blue line on the left, and a horizontal blue line at the bottom. Small white circles with blue outlines are positioned at the top-left and bottom-right corners where the lines meet.

Substitute Slides for Student Print version

Can you tell
the story?



The Transcriptional unit

