



## Topics

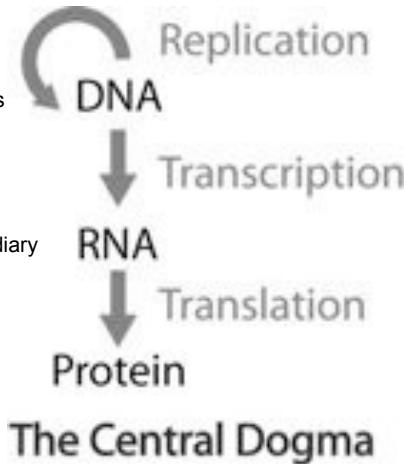
- Gene inheritance
- Genetic disorders
- Transcription
- Mechanism
- RNA processing
- Regulation of Gene Expression

## The flow of information in a cell

The master set of instructions

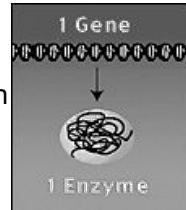
The messenger and intermediary

The product and executor

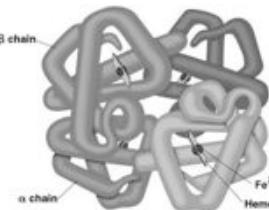


## The flow of information in a cell

Oversimplification



One gene-one polypeptide?

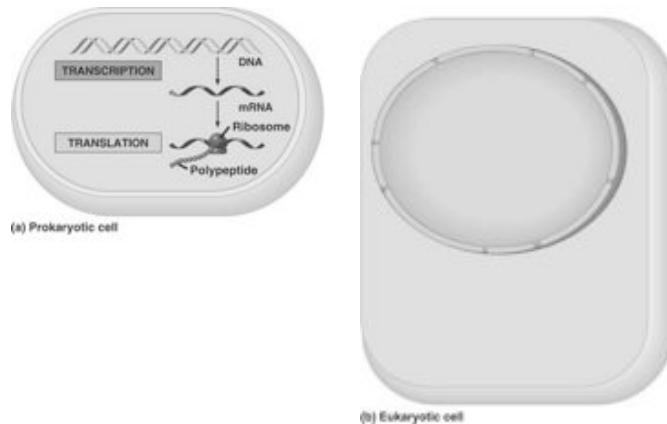


## DNA is Transcribed into RNA

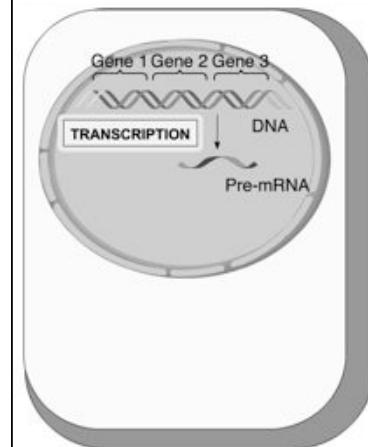
**Q: Why not use DNA directly? Why bother with RNA?**

- **Protection.** There is only **ONE** copy of each allele (gene) in the entire cell.
- **Amplification.** Transcription of one gene can make many RNAs
- **Efficiency.** Each RNA transcript can be repeatedly translated.

## Basic Principles of Transcription and Translation



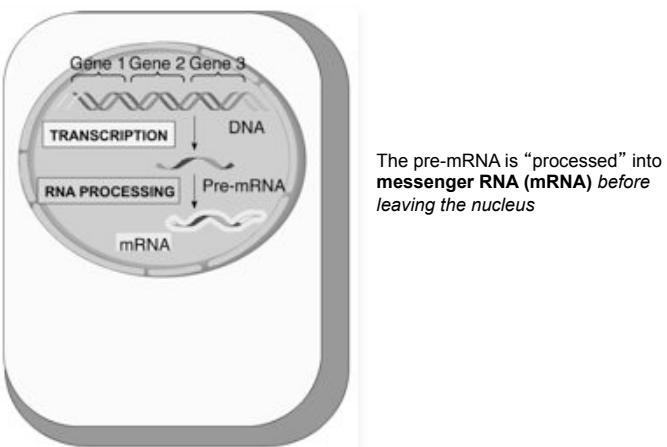
## Gene Expression in Eukaryotes



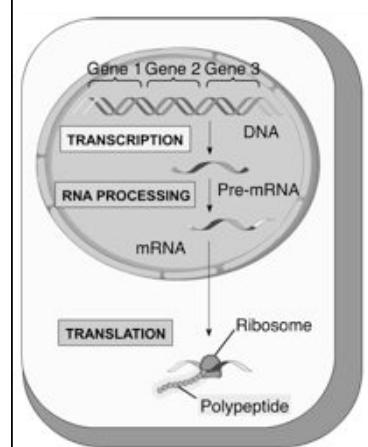
Transcription to a “pre-mRNA” occurs in the nucleus. Also called a **primary transcript**.

This is an exact replica of the gene sequence using **ribonucleotides** instead of deoxyribonucleotides

## Gene Expression in Eukaryotes

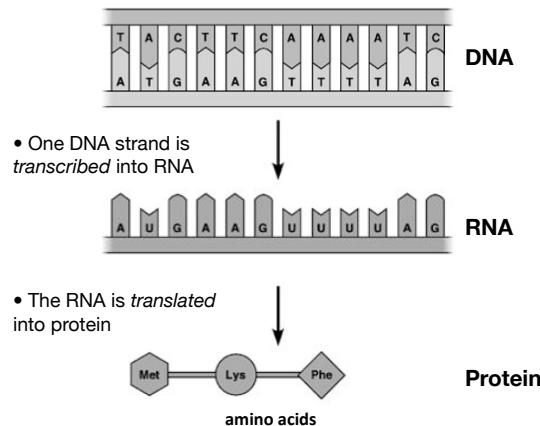


## Gene Expression in Eukaryotes

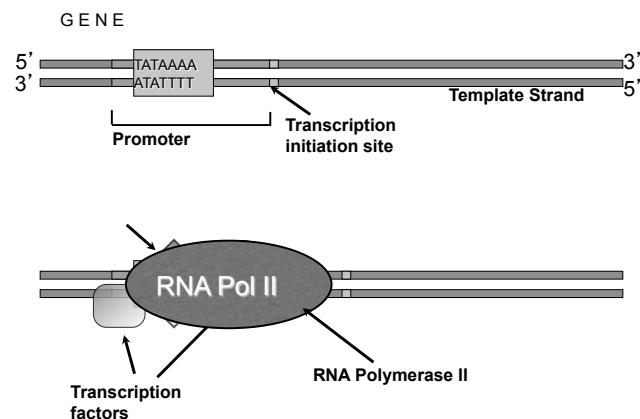


The mRNA sequence of nucleotides is **translated** into a sequence of amino acids in the cytoplasm with the help of **ribosomes**

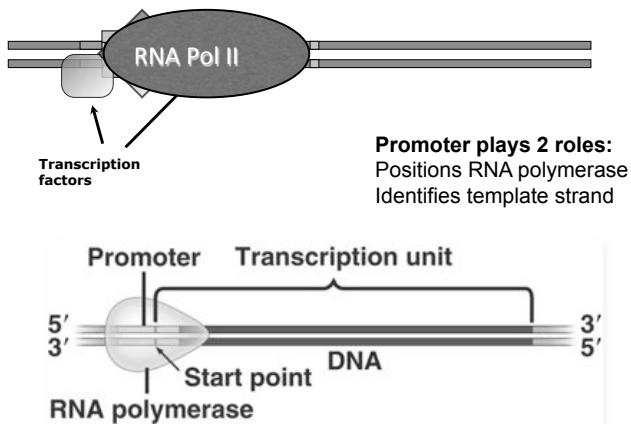
## Transcription/Translation



## Transcription

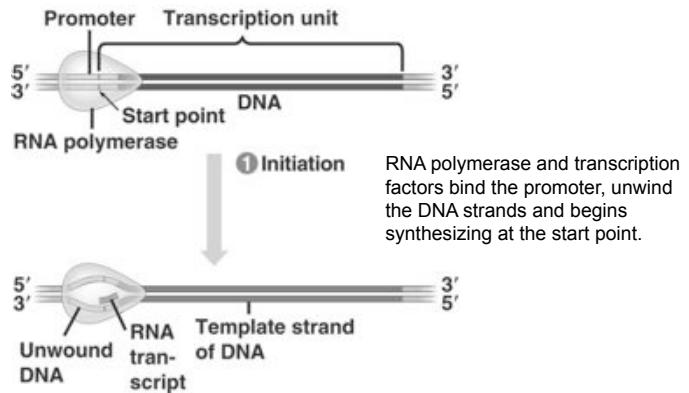


## Transcription



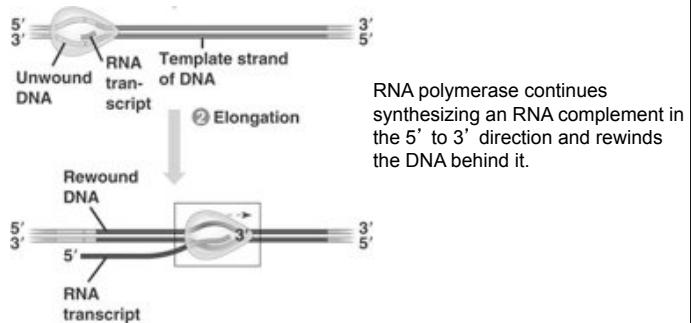
## Transcription- 3 steps

### 1. Initiation



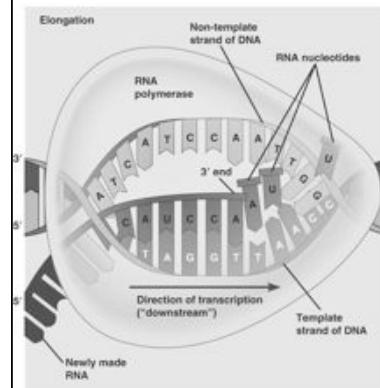
## Transcription- 3 steps

### 2. Elongation



## Transcription- 3 steps

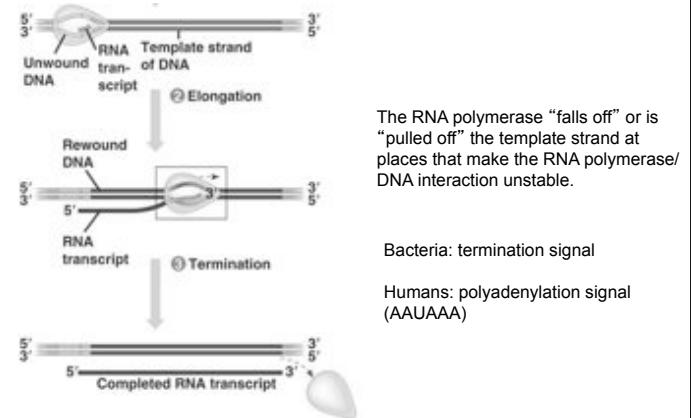
### 2. Elongation



Do you remember how RNA and DNA nucleotides differ?

## Transcription- 3 steps

### 3. Termination



## RNA Processing

The RNA is not yet ready for translation and must be **processed** first

### I. Alteration of RNA ends

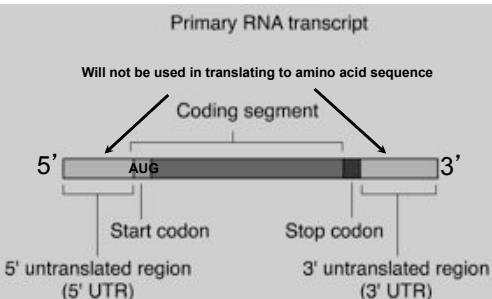
### II. RNA splicing

Does 1 gene = >1 protein?

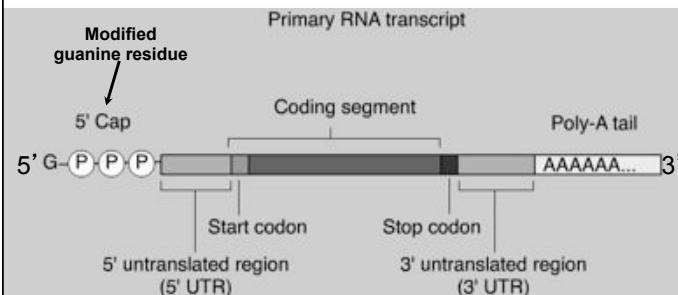
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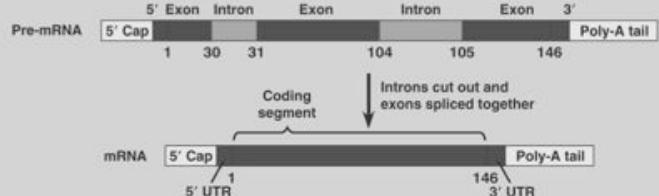


## RNA Processing



The **5' cap** and **3' poly-A tail** help protect the RNA from being degraded

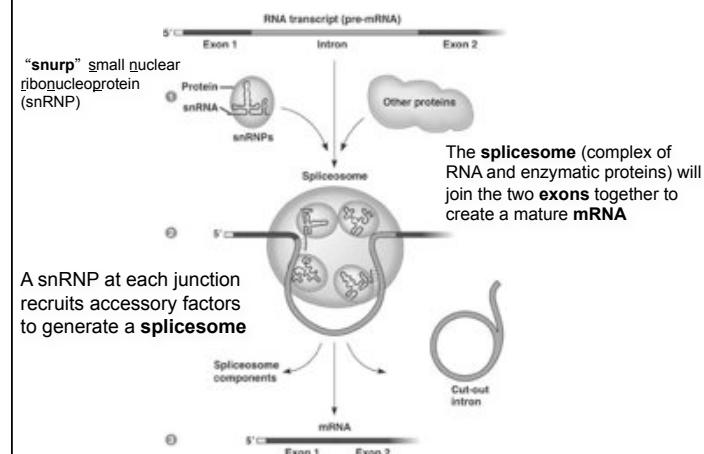
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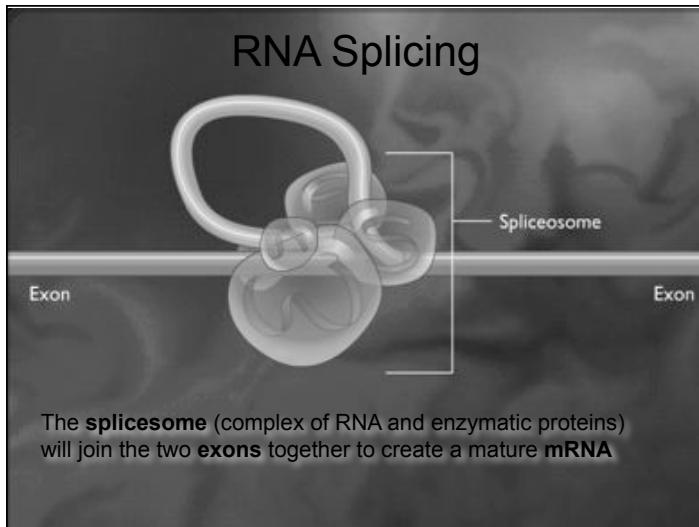


DNA in eukaryotes contains both **exons** and **introns**.

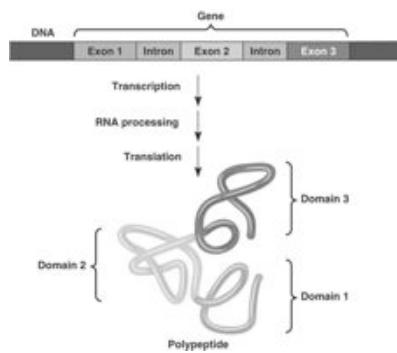
Only **exon** sequences end up in the coding sequence of the protein.

### How are introns removed?





### What's the Deal with Introns?



Depending on how the cell treats them, introns can be exons, and vice-versa. Called **alternative splicing**.

This would create diversity in protein sequence. Different **isoforms** of a protein.

### Differences between Prokaryotic and Eukaryotic Gene Expression

Characteristic	Prokaryotic	Eukaryotic
Transcription location	cytoplasm	nucleus
Translation location	Linked to transcription	cytoplasm
promoter	Shared by several genes	One gene = one promoter

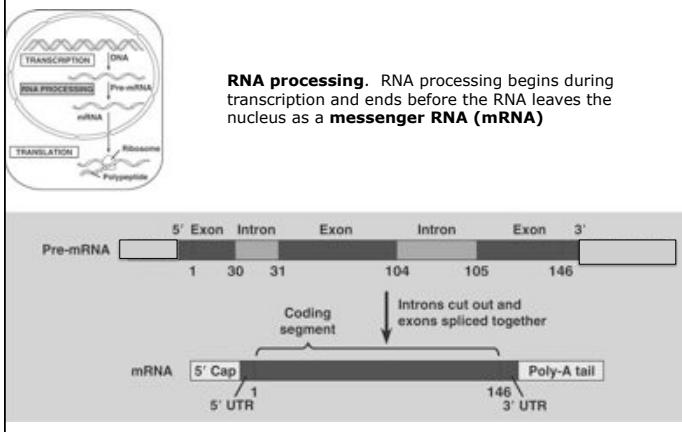
### Differences between Prokaryotic and Eukaryotic Gene Expression

Characteristic	Prokaryotic	Eukaryotic
Transcription location	cytoplasm	nucleus
Translation location	Linked to transcription	cytoplasm
promoter	Shared by several genes	One gene = one promoter
Gene structure	Not interrupted by introns	Transcribed regions often interrupted by non-coding introns
Post-transcriptional regulation	none	Introns removed 5' cap and 3' poly A tail added

## The Forms and Functions of RNA

Form	Function
Primary Transcript	The precursor to ALL RNAs. mRNAs begin as primary transcripts and are then processed to begin mature mRNAs.
messenger RNA (mRNA)	Carries information specifying amino acid sequences of proteins from DNA to ribosomes

## RNA Processing



## The Forms and Functions of RNA

Form	Function
Primary Transcript	The precursor to mRNA, rRNA or tRNA before being processed by splicing or cleavage.
messenger RNA (mRNA)	Carries information specifying amino acid sequences of proteins from DNA to ribosomes
snRNA	Plays structural and catalytic roles in spliceosomes, the complexes of protein and RNA that splice pre-mRNA
Transfer RNA (tRNA)	Serves as adapter molecule in protein synthesis; translates mRNA codons into amino acids
Ribosomal RNA (rRNA)	Plays catalytic (ribozyme) roles and structural roles in ribosomes

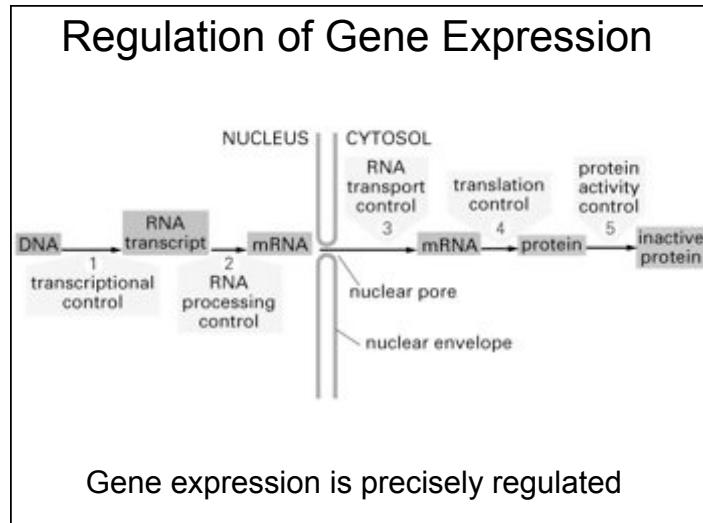
## The Many Functions of RNA

### What makes RNA so versatile?

1. RNA is single-stranded. It can form hydrogen bonds with its bases to DNA, proteins or other RNAs.
2. RNA can develop base pairing with its own strand to create varied forms of 3-dimensional structure.
3. RNA's varied structure and functional groups can allow it to act as a type of enzyme... called **ribozymes**

**Some RNAs are 'self-splicing'!**

## Regulation of Gene Expression

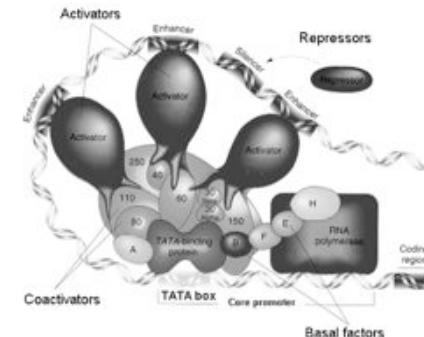


## Transcriptional Regulation in Eukaryotes

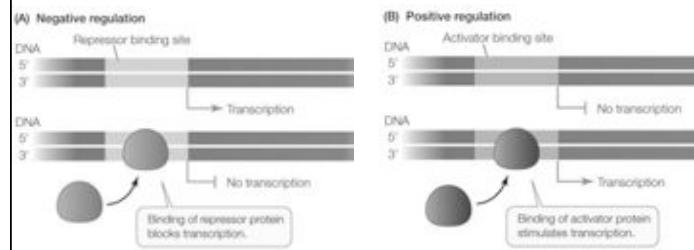
**TATA-box:** core promoter sequence bound by RNA pol, in 50-70% of promoters

**General transcription factors:** interact with most promoters, help position the RNA polymerase

**Activators/Repressors:** highly gene specific, allows varied gene expression



## Positive vs. Negative Gene Regulation



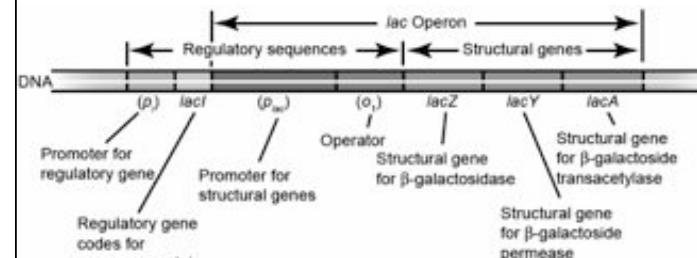
**Inducible** gene expression: expressed only when needed by the cell, involves repressors and activators

**Constitutive** gene expression: actively expressed at all times

## Transcriptional Regulation in Prokaryotes

**Operon:** cluster of genes regulated by a single promoter

**Operator:** repressor binding site that helps regulate expression



## Why regulate gene expression?

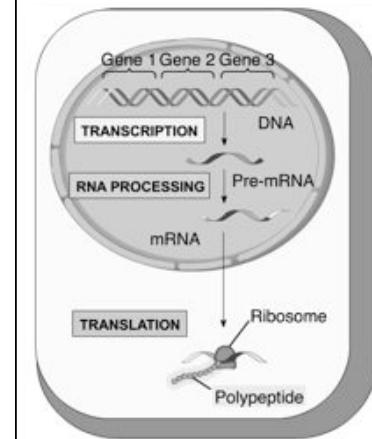
Conserve energy

regulating at level of transcription is most efficient  
protein synthesis is highly endergonic

To be able to respond to the environment

Allows greater variety of proteins made as needed

## Gene Expression in Eukaryotes



The mRNA sequence of nucleotides is **translated** into a sequence of amino acids in the cytoplasm with the help of **ribosomes**

## The Code has a Reading Frame

Every three bases codes for an amino acid

Each triplet is called a **codon**

*with three letter words representing codons*

Example in frame:

“The red dog ate the cat”

Frameshifted by one:

“T her edd oga tet hec at”

## The Nucleotide Sequence is a Code

AUGACCAAAACCGAGUUGA

Cracking the Code

RNA  
UUUUUUUUUUUUUUUUUUUUUU...



Protein

Phe-Phe-Phe-Phe-Phe...

AUG ACC AAA CCG AGU UGA

First position (5' end)	U	C	A	G	Third position (3' end)
U	Phe	Ser	Tyr	Cys	U
	Phe	Ser	Tyr	Cys	C
	Leu	Ser	STOP	STOP	A
	Leu	Ser	STOP	Trp	G
C	Leu	Pro	His	Arg	U
	Leu	Pro	His	Arg	C
	Leu	Pro	Gln	Arg	A
	Leu	Pro	Gln	Arg	G
A	Ile	Thr	Asn	Ser	U
	Ile	Thr	Asn	Ser	C
	Ile	Thr	Lys	Arg	A
	Met	Thr	Lys	Arg	G
G	Val	Ala	Asp	Gly	U
	Val	Ala	Asp	Gly	C
	Val	Ala	Glu	Gly	A
	Val	Ala	Glu	Gly	G

## The Code is Redundant

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

AUG (Met): always the starting codon

There are multiple codons for most amino acids

3rd base “wobble”

There is NO ambiguity...

## Overview of amino acids

