



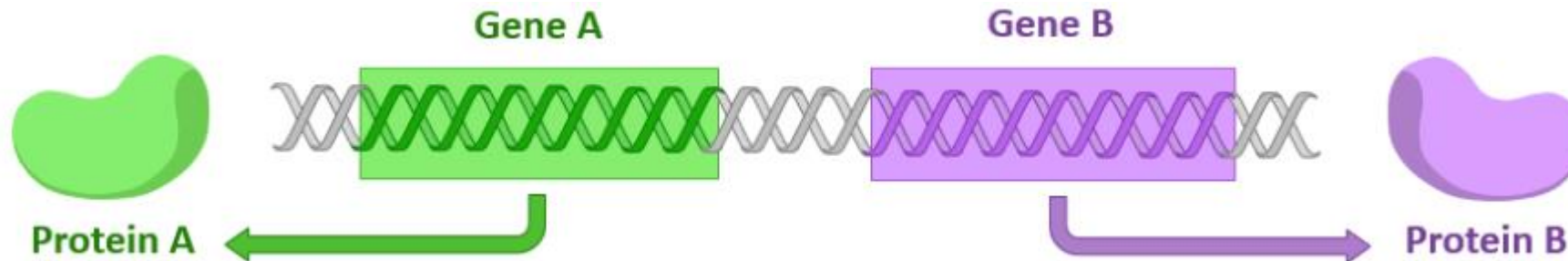
3.1

Genes

BASIC IDEA

DNA = genetic blueprint → codes for & determines an organism's characteristics
it encodes proteins, which carry out cellular processes

A specific length / sequence of DNA that encodes for a particular protein = **gene**
→ heritable factor that influences a specific characteristic



COMPARING NUMBERS OF GENES

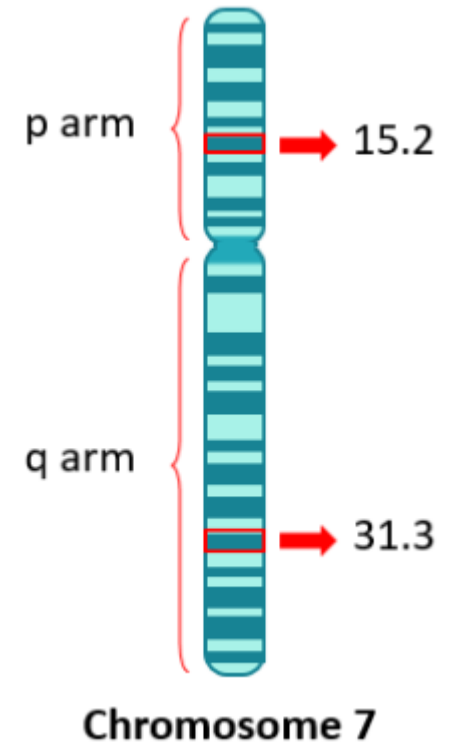
IB companion: p.142 orange box

GENE LOCI

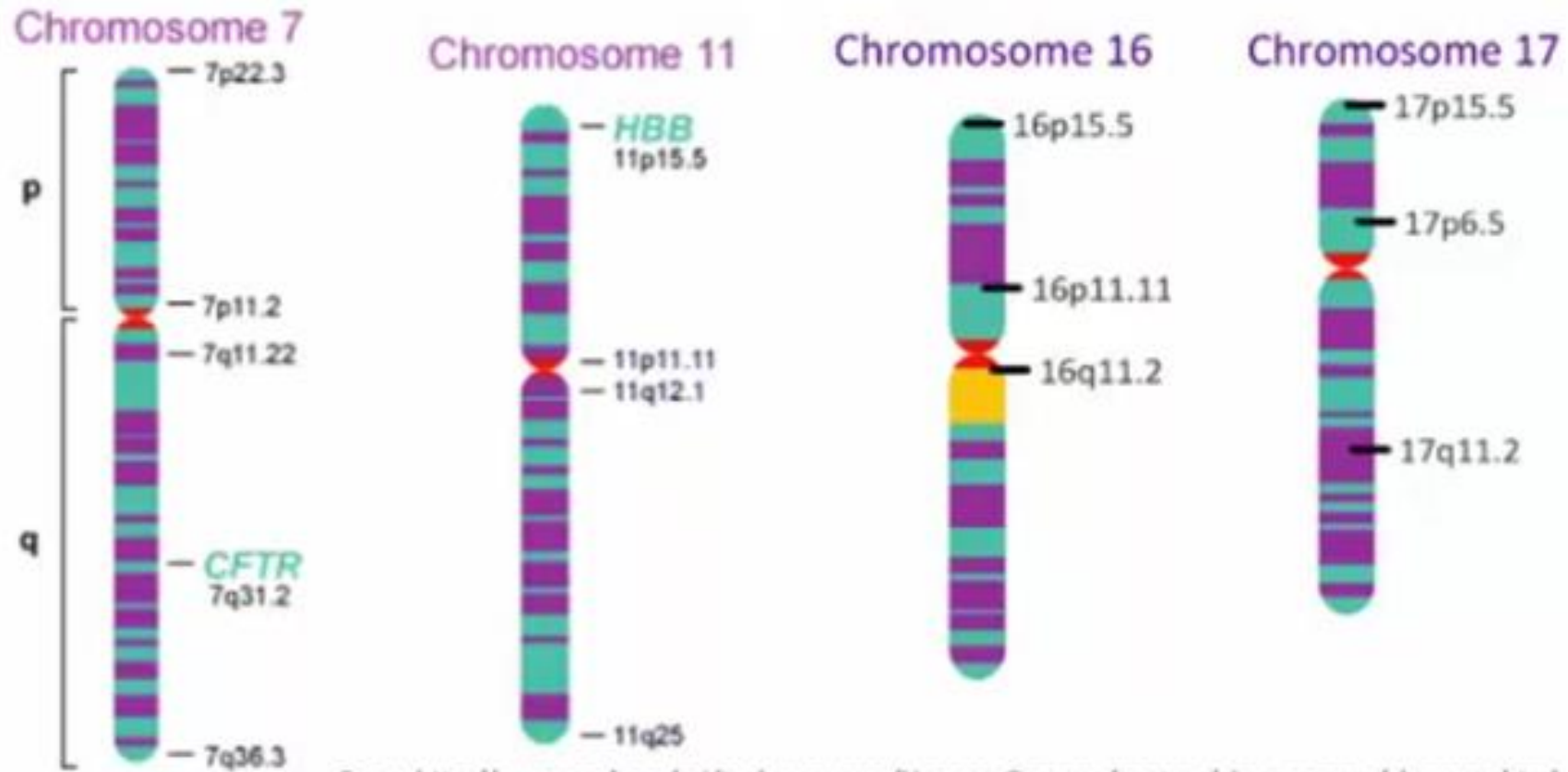
Position of a gene on a particular chromosome = **locus**
(plural = loci)

Eukaryotic gene loci identified to following features:

- chromosome number
- arm (p = short ; q = long)
- region and band



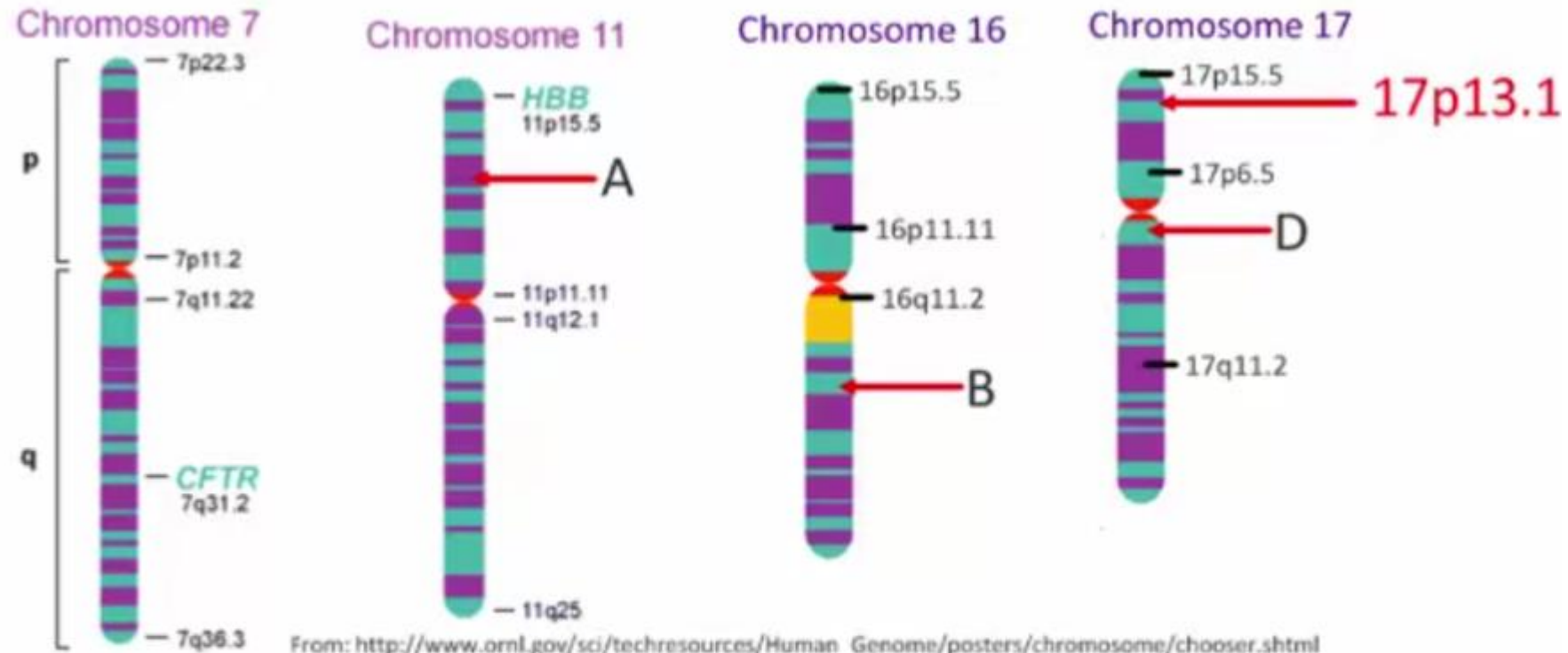
GENE LOCI: EXAMPLES



From: http://www.ornl.gov/sci/techresources/Human_Genome/posters/chromosome/chooser.shtml

The gene **TP53** (prevents tumours) is found at **17p13.1**.

GENE LOCI: EXAMPLES



Match the gene loci with the letters

16q13.2

11p13.1

17q1.1

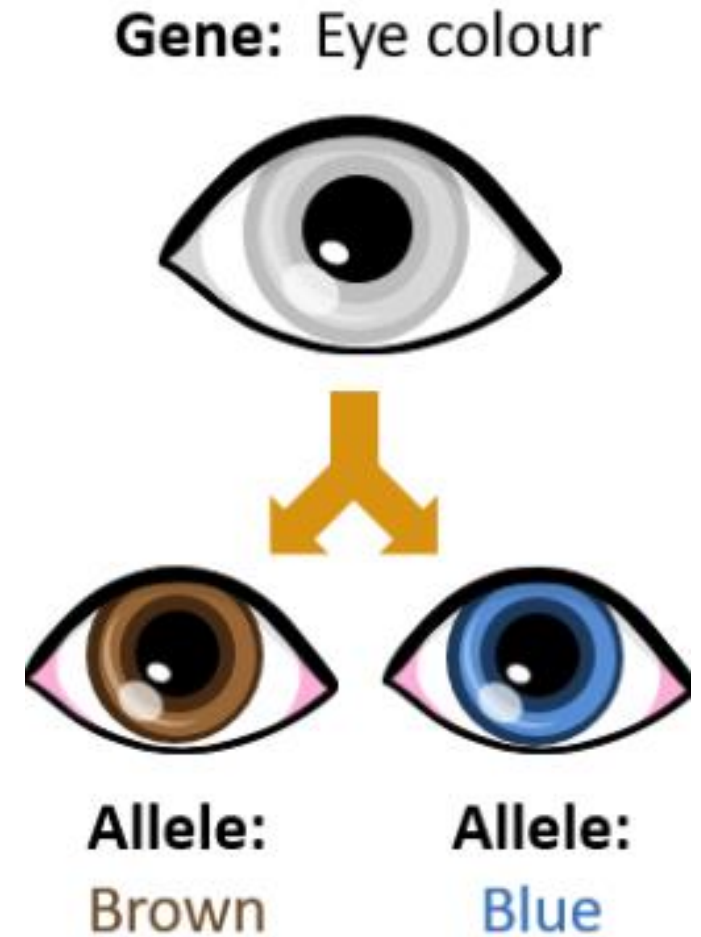
ALLELES

Alleles = different possible variations of a gene











Genes encode a general characteristic: Eye color

Alleles encode for a specific alternative: Blue eyes

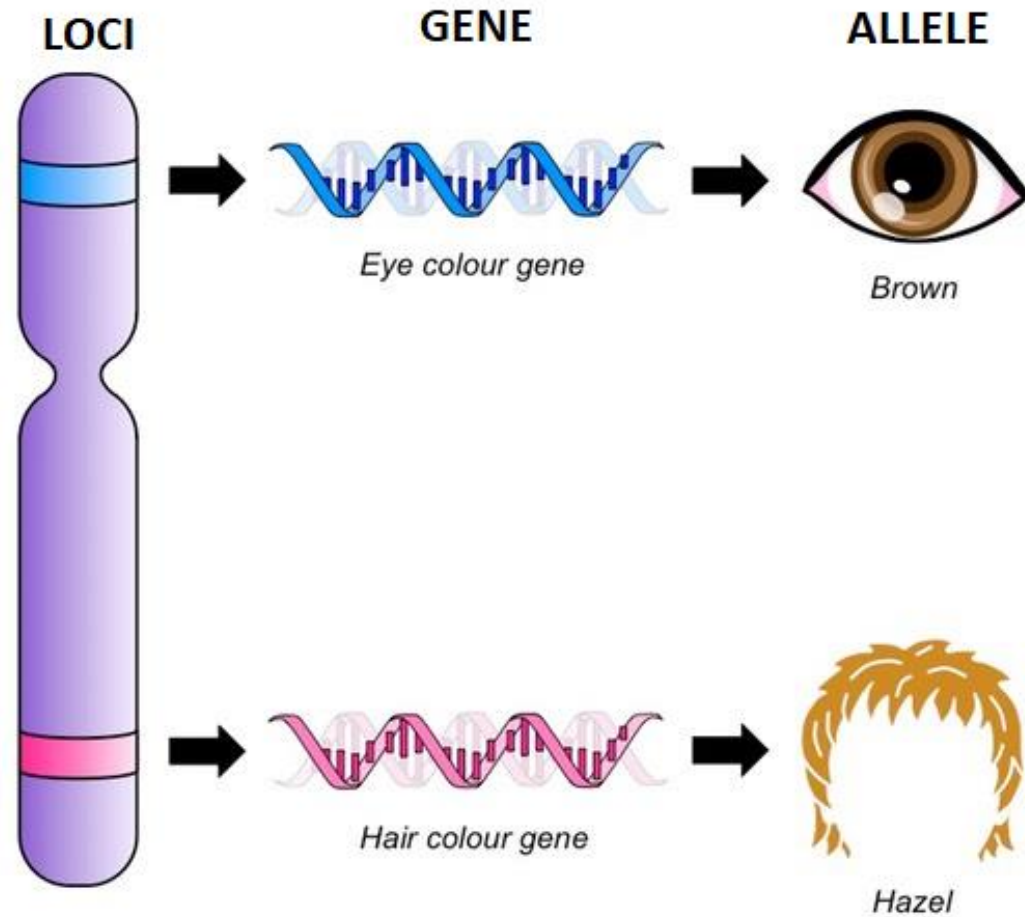
Alleles only differ from each other by one or few bases



ALLELES

Gene	Alternative Alleles			
 <i>Eye colour</i>	 <i>Brown</i>	 <i>Blue</i>	 <i>Emerald</i>	 <i>Grey</i>
 <i>Hair colour</i>	 <i>Blonde</i>	 <i>Red</i>	 <i>Brown</i>	 <i>Black</i>

LOCUS VS GENE VS ALLELE



COX-2, SMOKING AND STOMACH CANCER

Data-based questions page 145

MUTATIONS

A **gene mutation** is a change in the nucleotide sequence of a section of DNA coding for a specific trait → may change a protein's function

Gene mutations can either be:

- **Somatic** – occur in body cells → not inherited
- **Germline** – occur in gametes → can be passed on to offspring



MUTATIONS








Gene mutations can be:

- Beneficial: new variations of a trait
- Detrimental: cannot carry out normal function
- Neutral: no effect on functioning



MUTAGEN

A mutagen is an agent that changes (mutates) the genetic material of an organism

Physical		Chemical			Biological	
						
UV <i>(from sun)</i>	X-rays <i>(medical)</i>	Carcinogen <i>(cigarettes)</i>	Processed foods	Cleaning products	Viruses <i>(HPV)</i>	Bacteria <i>(Helicobacter)</i>

TYPES OF MUTATIONS

Point mutations → modification of a single nucleotide within a sequence

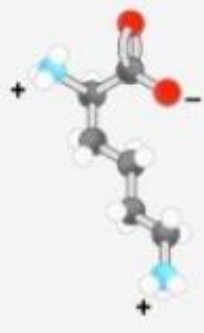
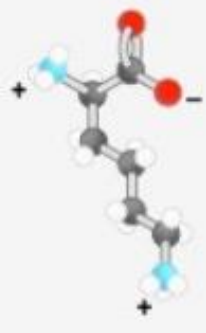

- **substitutions, insertions or deletions**

Substitution mutations can either cause:

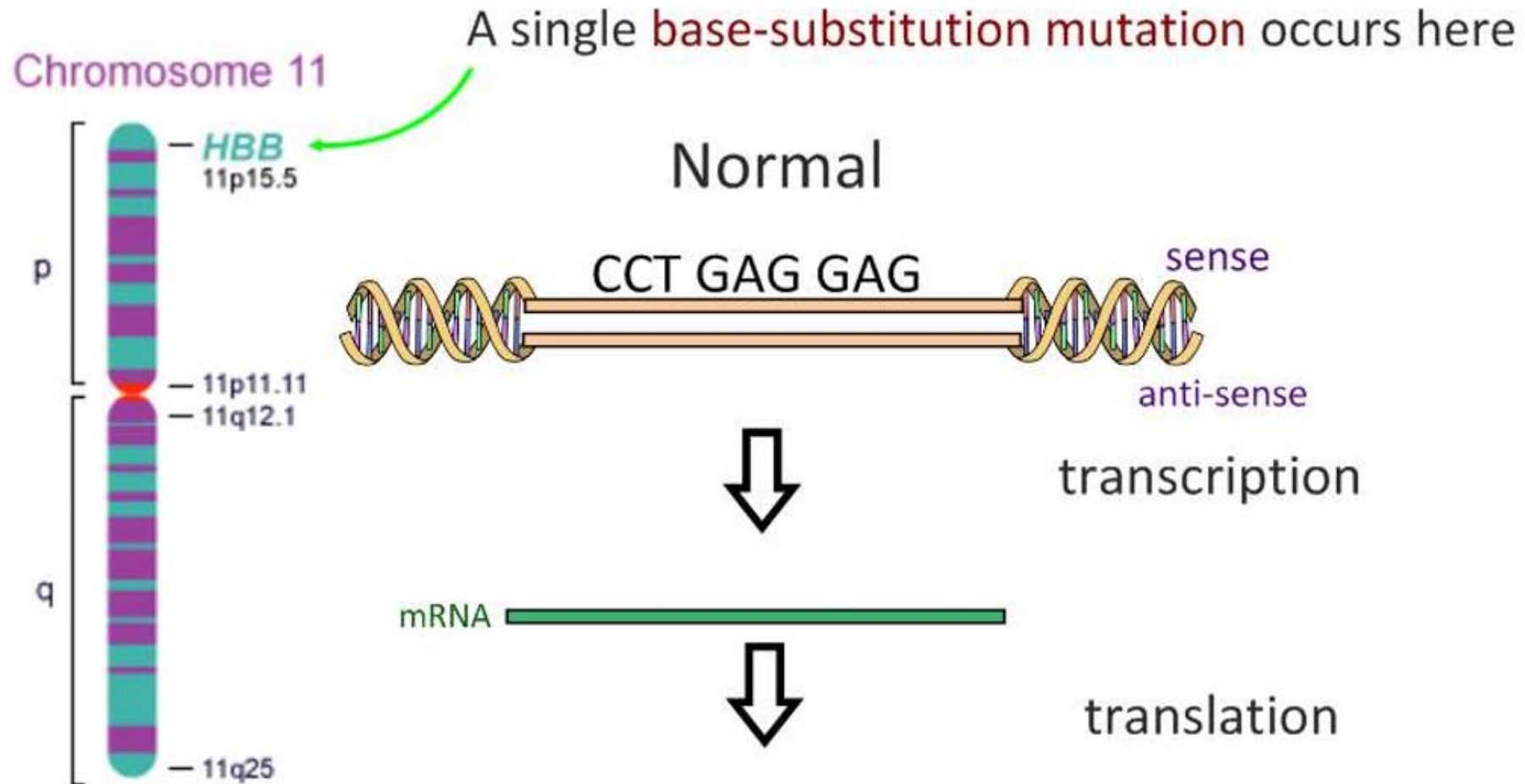
- *Silent mutations* – no change to sequence (*due to degeneracy*)
- *Missense mutations* – a single amino acid is changed in the polypeptide
- *Nonsense mutations* – a stop codon is created, malfunction of polypeptide

Insertions and deletions cause frameshift mutations (*changes reading frame*)

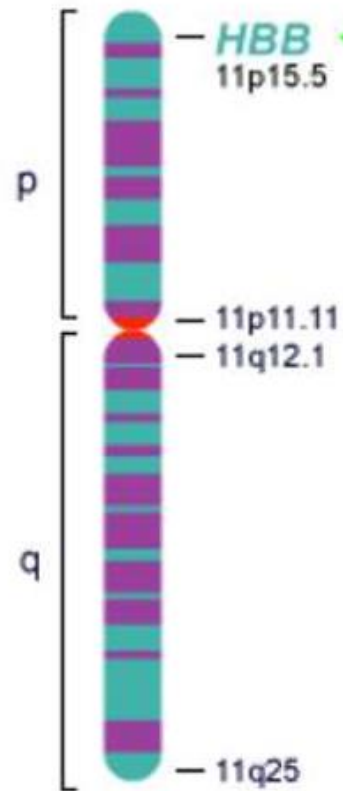
TYPES OF MUTATIONS:

	No mutation	Point mutations		
		Silent	Missense	Nonsense
DNA	TTC	TTT	TCC	ATC
mRNA	AAG	AAA	AGG	UAG
Protein	Lys	Lys	Arg	STOP
				

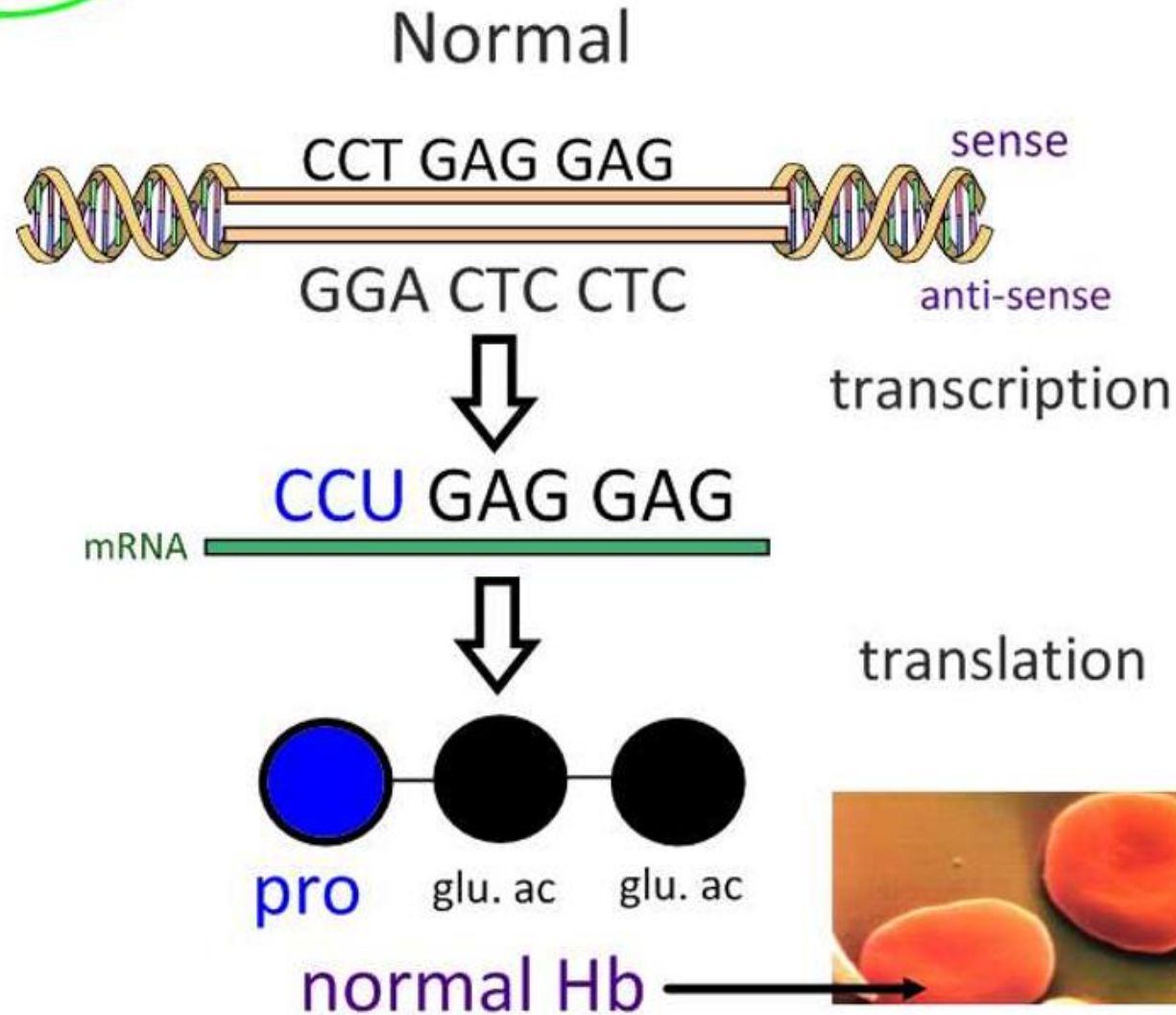
MUTATION EXAMPLE: SICKLE CELL



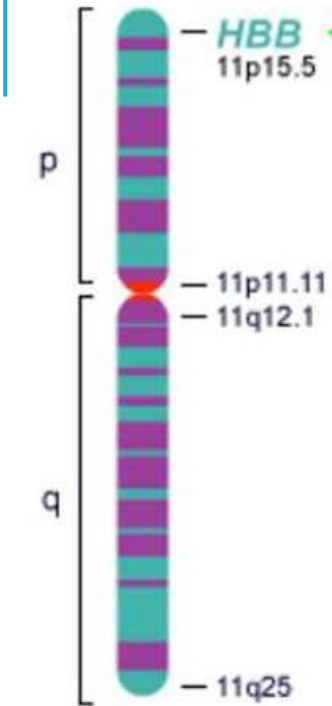
Chromosome 11



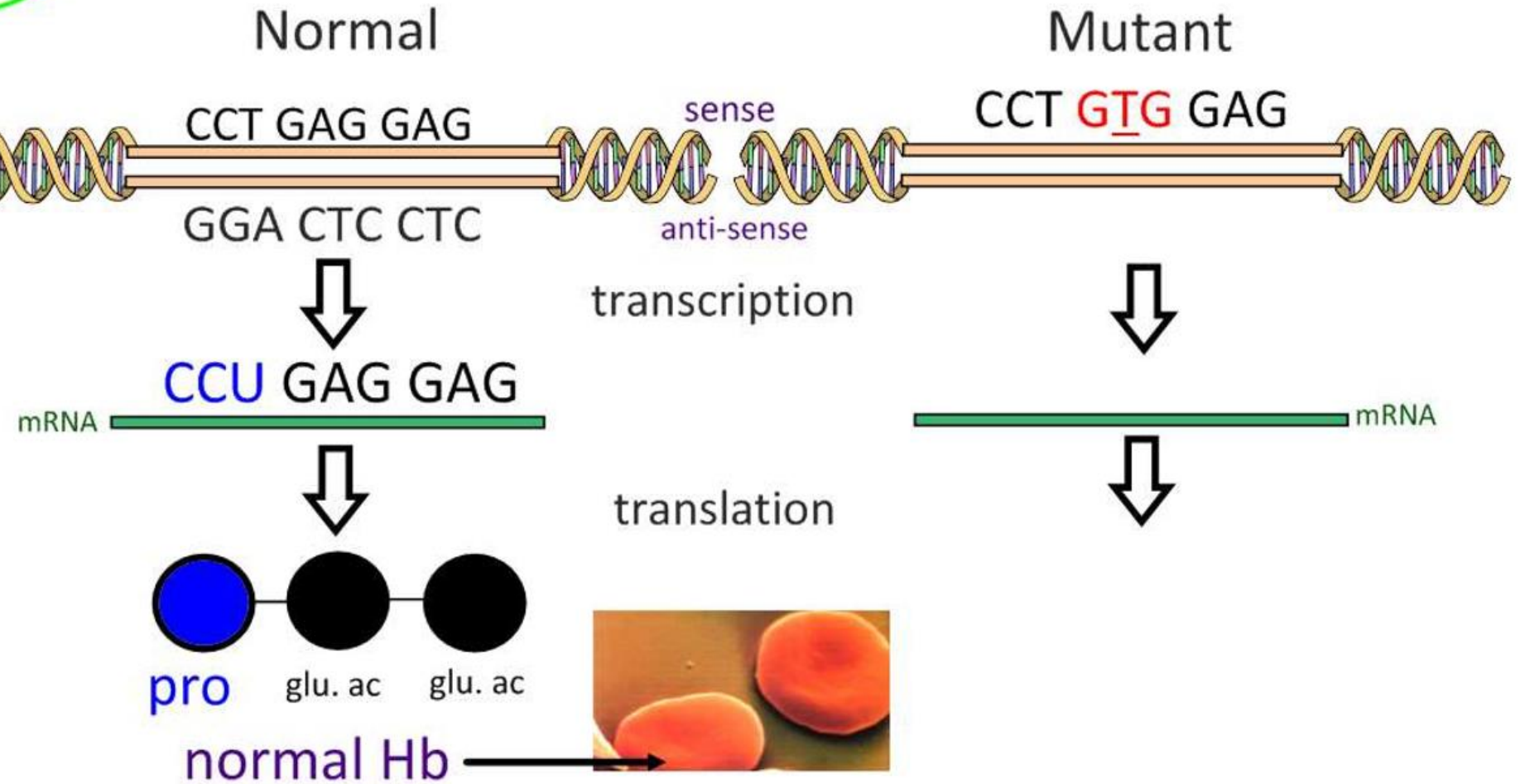
A single **base-substitution mutation** occurs here



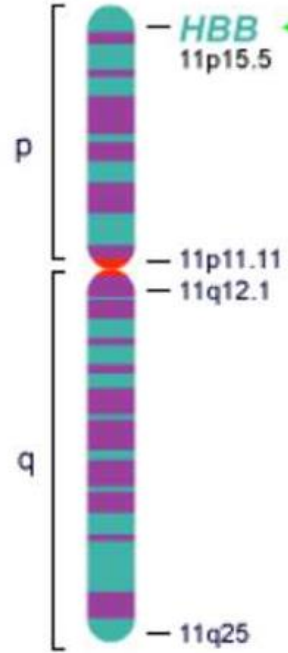
Chromosome 11



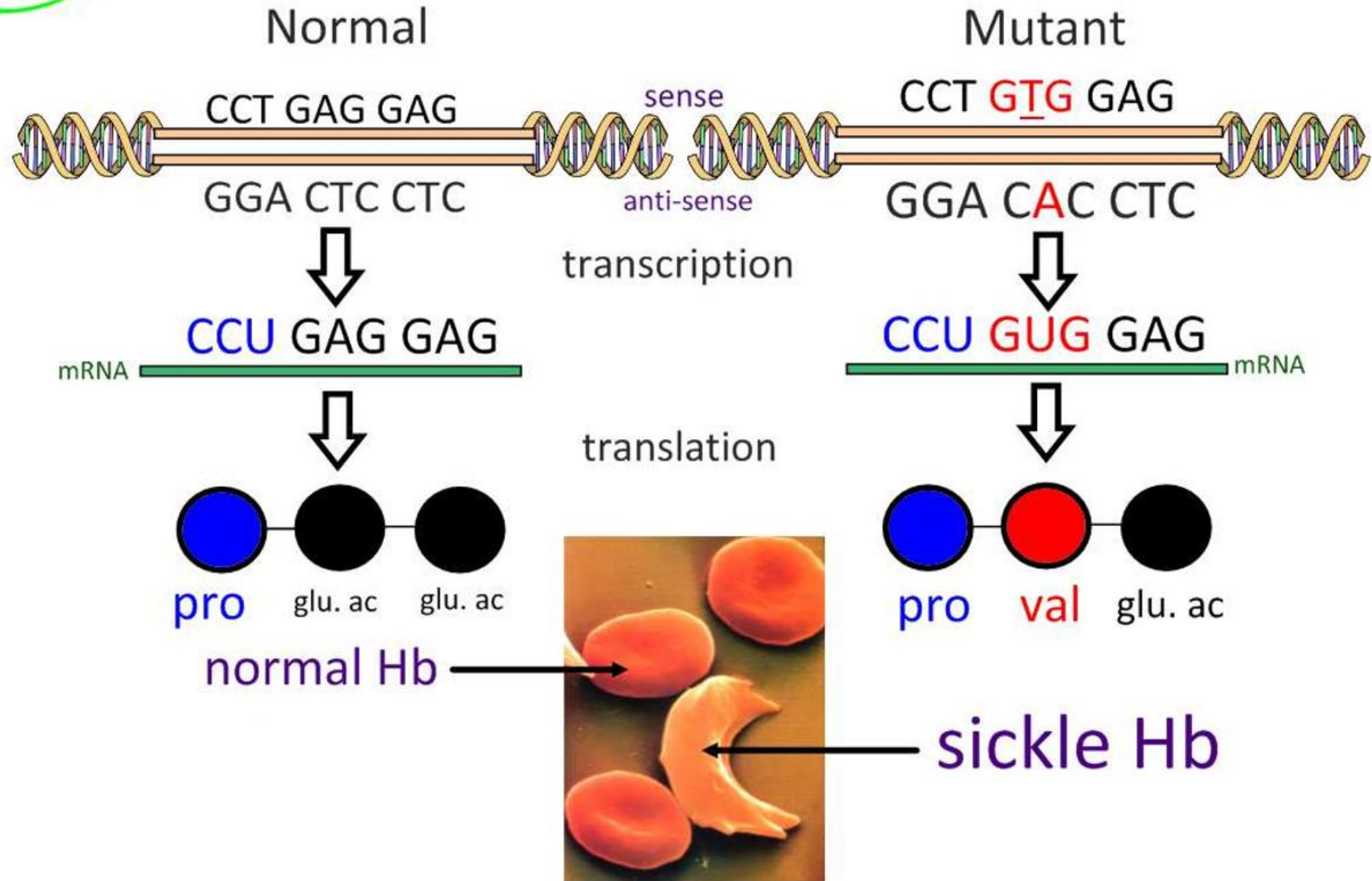
A single **base-substitution mutation** occurs here



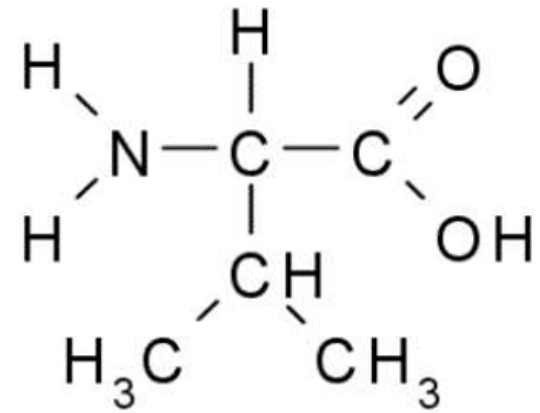
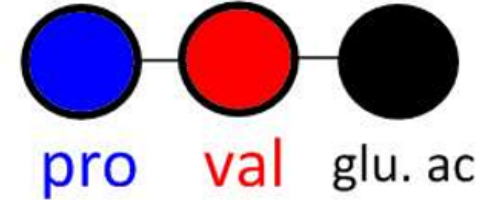
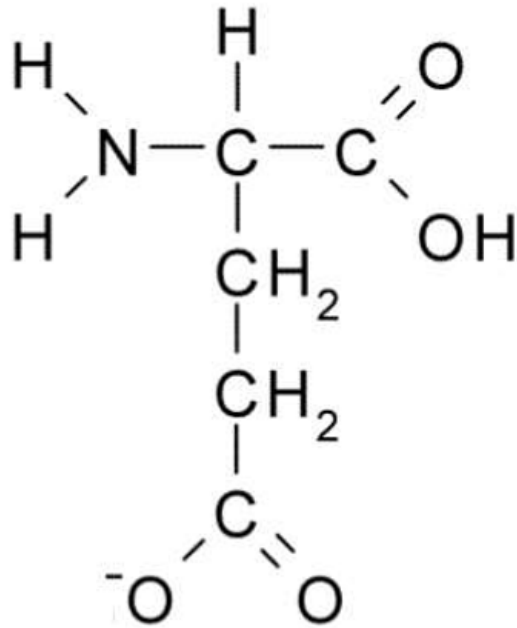
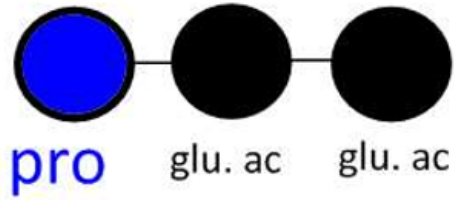
Chromosome 11



A single **base-substitution mutation** occurs here



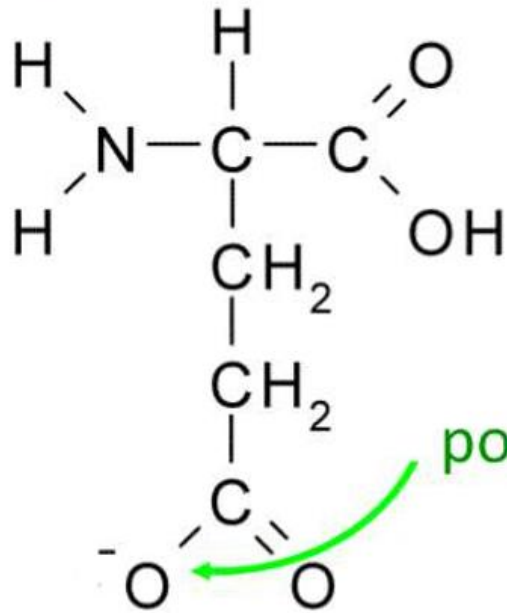
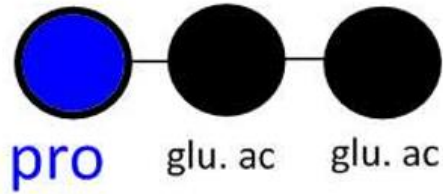
Comparing glutamic acid and valine



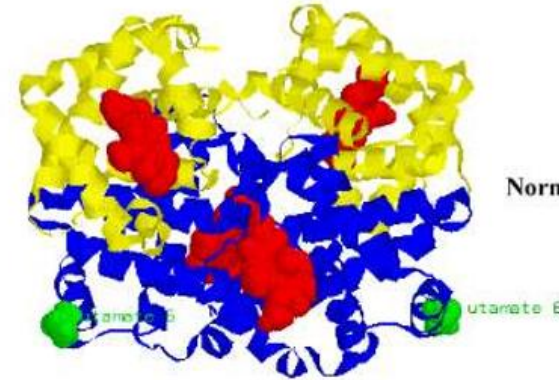
What differences in **properties** can we deduce from the structures?

How might this change the quaternary protein structure?

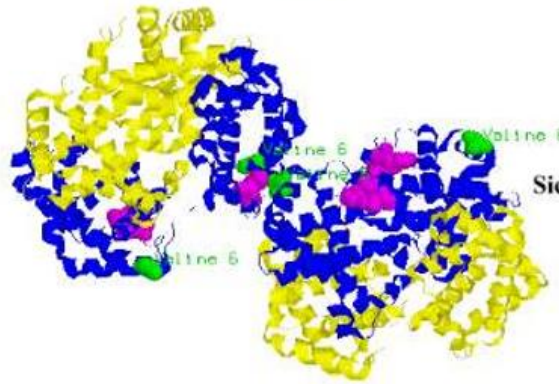
Comparing glutamic acid and valine



polar

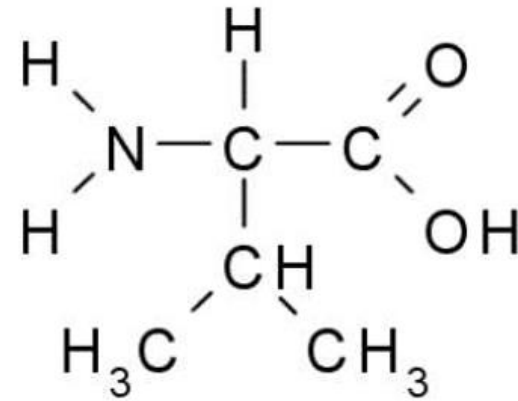
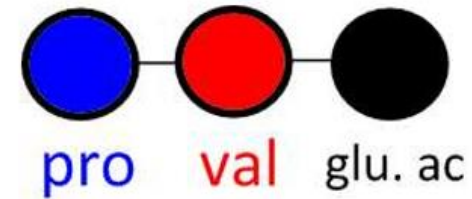


Normal Hemoglobin



Sickle Hemoglobin

Note: The Sickle hemoglobin image is drawn at 50% of the size of the Normal hemoglobin



non-polar

What differences in **properties** can we deduce from the structures?

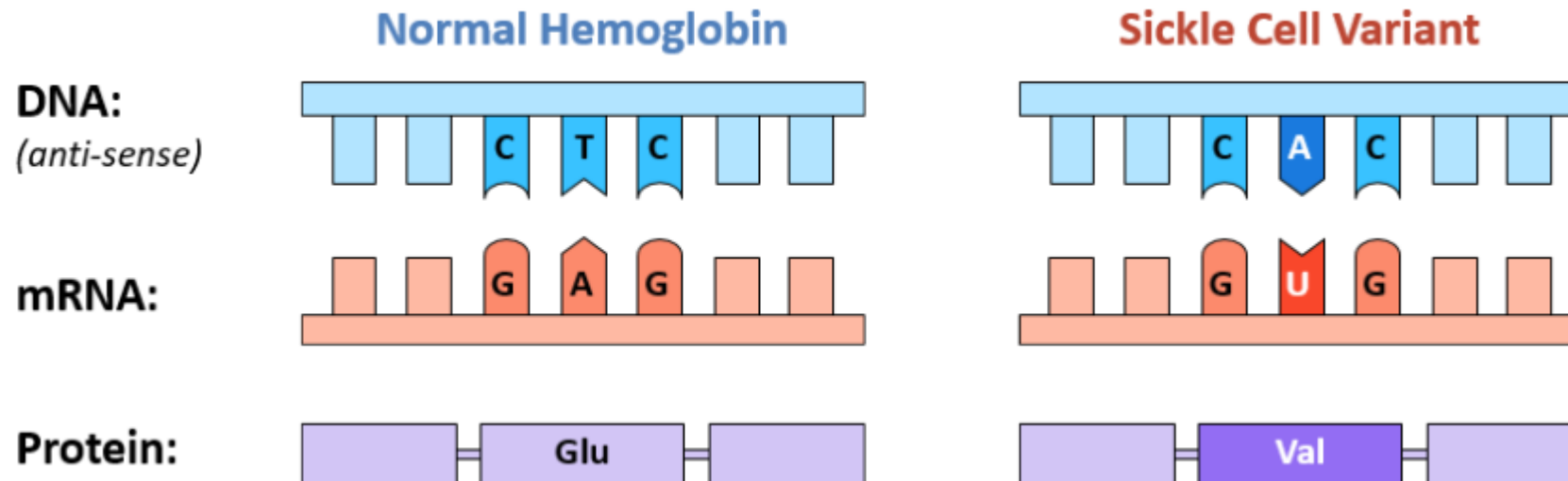
How might this change the quaternary protein structure?

SUMMARY: SICKLE CELLS

Sickle cell anemia is caused by a base substitution in hemoglobin

- 6th codon (*sense strand*): **GAG → GTG**

- Amino acid: **Glu → Val**



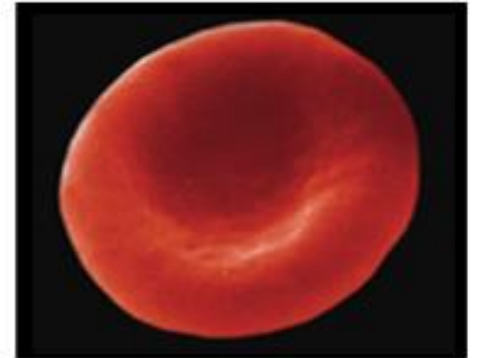
MUTATION CONSEQUENCE: ANAEMIA

The amino acid change alters structure → hemoglobin forms insoluble strands

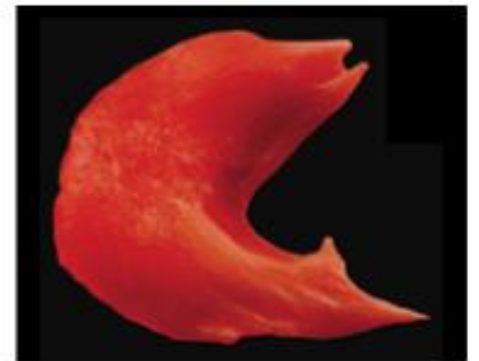
These insoluble strands change the shape (*sickle shape*)

The red blood cells cannot carry oxygen effectively and are destroyed at a higher rate – leading to **anemia**

Sickle cells are more likely to form clots

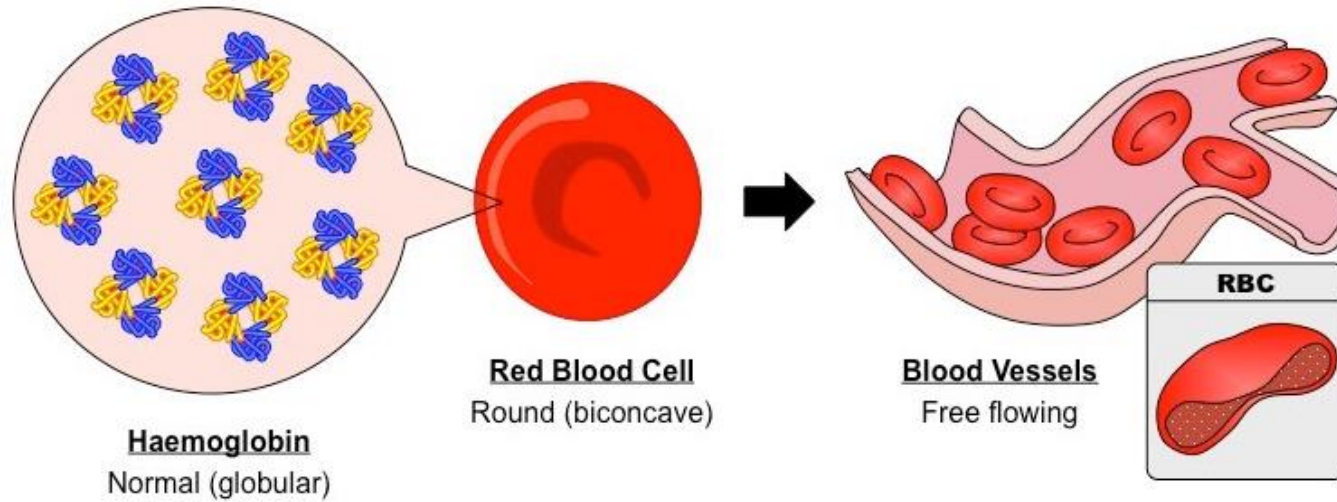


Normal Blood Cell

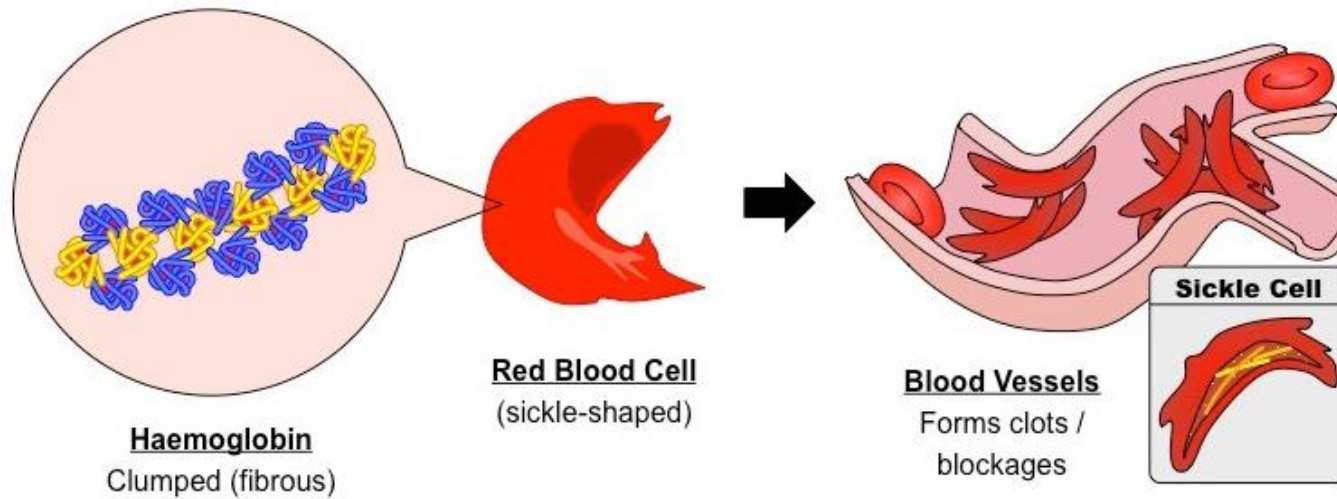


Sickle Blood Cell

Normal (Wild-Type) Haemoglobin

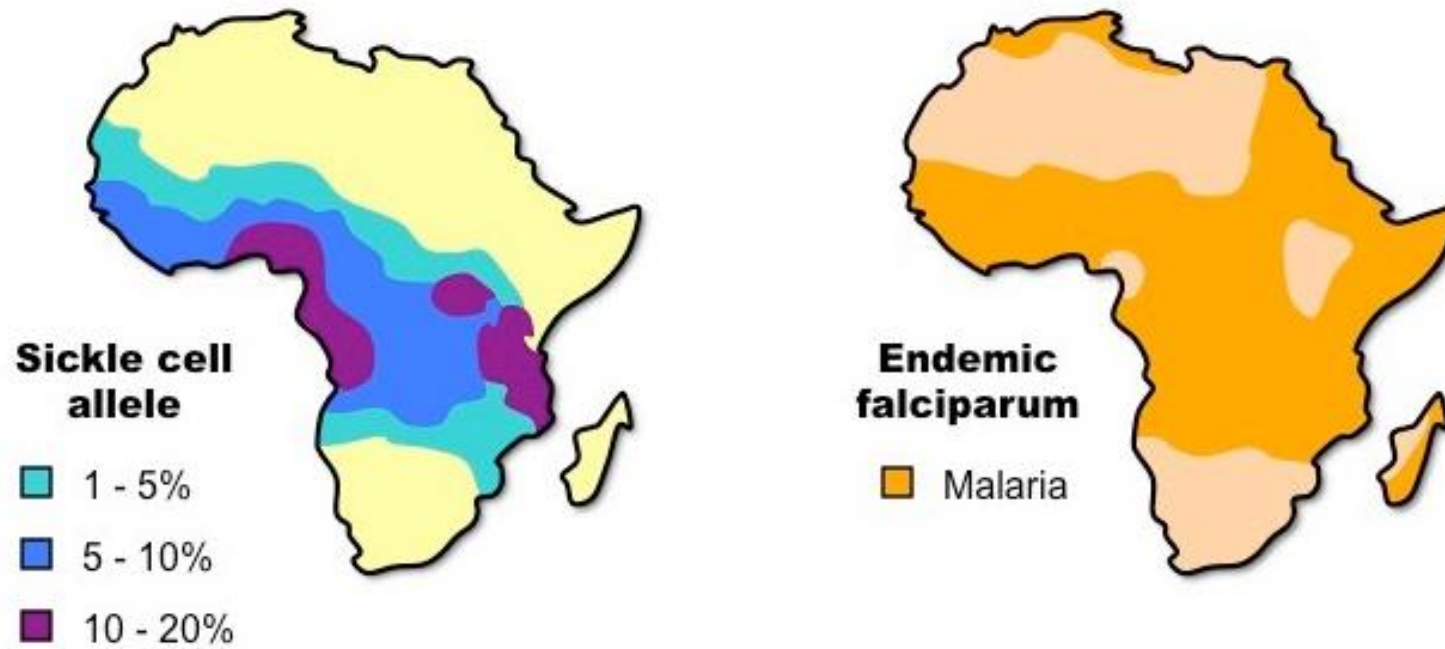


'Sickle Cell' Haemoglobin



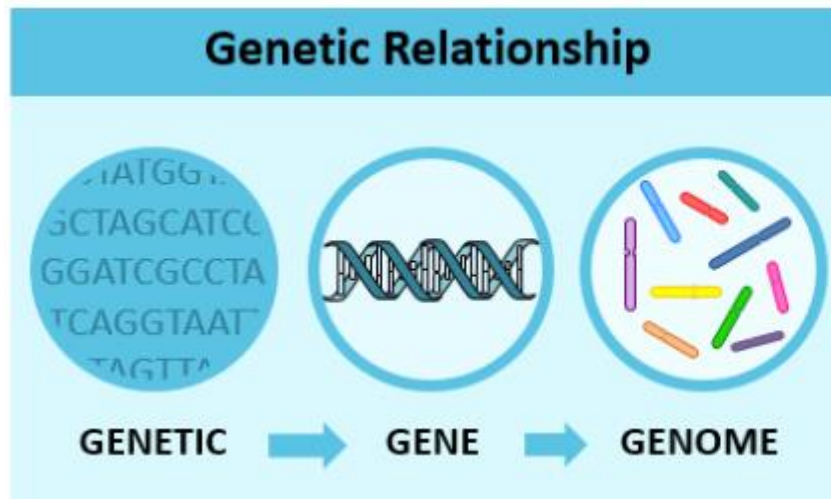
SICKLE CELL VERSUS MALARIA

Correlation between Prevalence of Sickle Cell Allele and Malaria



GENOMES

Genome = totality of genetic information in a cell or organism (coding + non-coding sequences)



HUMAN GENOME

The Human Genome Project (HGP) → international cooperative venture → sequence entirety of human genome.

- 46 chromosomes
- ~3 billion base pairs
- ~21 000 genes

GENETICS COMPARISON






Species differ in genome size, number of chromosomes and number of genes

There is **no** clear correlation between these factors → but other trends






- viruses and prokaryotes tend to have a smaller genome + fewer genes
- eukaryotes have huge variation in genome size and gene numbers

Overall **no** consistent or reliable pattern.






GENOME SIZE

Species	<i>T2 phage</i>	<i>Escherichia coli</i>	<i>Drosophila melanogaster</i>	<i>Homo sapiens</i>	<i>Paris japonica</i>
Genome Size	170,000 bp	4.6 million bp	130 million bp	3.2 billion bp	150 billion bp
Common Name	 Virus	 Bacteria	 Fruit fly	 Human	 Canopy Plant

CHROMOSOME NUMBER

Species	<i>Parascaris equorum</i>	<i>Oryza sativa</i>	<i>Homo sapiens</i>	<i>Pan troglodytes</i>	<i>Canis familiaris</i>
Chromosomes	4	24	46	48	78
Common Name	 Roundworm	 Rice	 Human	 Chimpanzee	 Dog

GENE NUMBERS

Species	<i>Escherichia coli</i>	<i>Gallus gallus</i>	<i>Homo sapiens</i>	<i>Daphnia pulex</i>	<i>Oryza sativa</i>
Genome Size	~4,200	~17,000	~21,000	~31,000	~38,000
Common Name	 Bacteria	 Chicken	 Human	 Water flea	 Rice