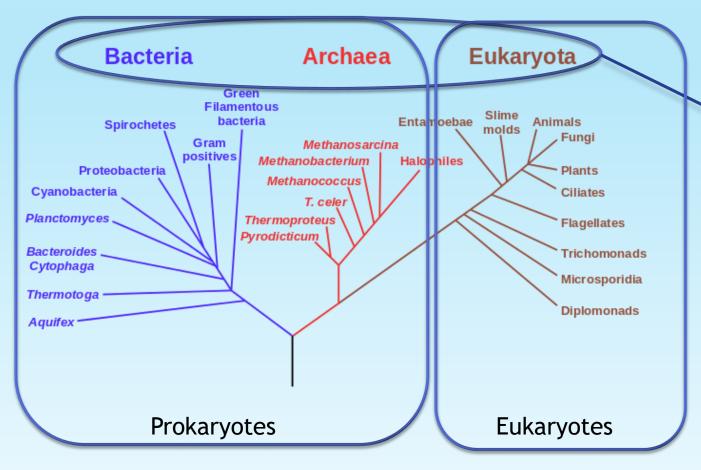
Biology Bioinformaticians

Gregor Gilfillan

Topics:

- 1. Tree of life
- 2. Building blocks of life
- 3. Structure (and differences) of DNA and RNA
- 4. DNA makes RNA makes protein
- 5. Genomes and genomic features
- 6. Genetics
- 7. Epigenetics

1. Phylogenetic tree of life



Tree assembled by comparison of rDNA sequences.

Viruses?

Hierarchical classification of life Life **Domain** Kingdom **Phylum** Class Order Family Genus

Species

Prokaryotes vs Eukaryotes

Prokaryote





Eukaryote



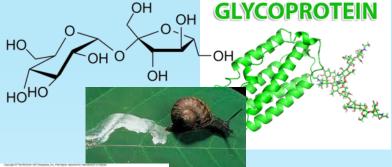
Size
Ribosome type
Chromosomes
Histone proteins
Membranes with steroid content

Defining differences = Eukaryotes:

- Contain DNA in a nucleus
- Contain organelles (mitochondria, also chloroplasts in some)

2. Building blocks of life

Carbohydrates



Energy / Structure / Identification / Lubrication / cometabolites

Lipids (fats)

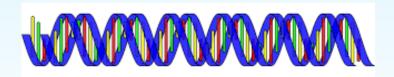
Proteins

Fat section Fat cells (adipocytes)

Energy storage / membranes / signaling / insulation

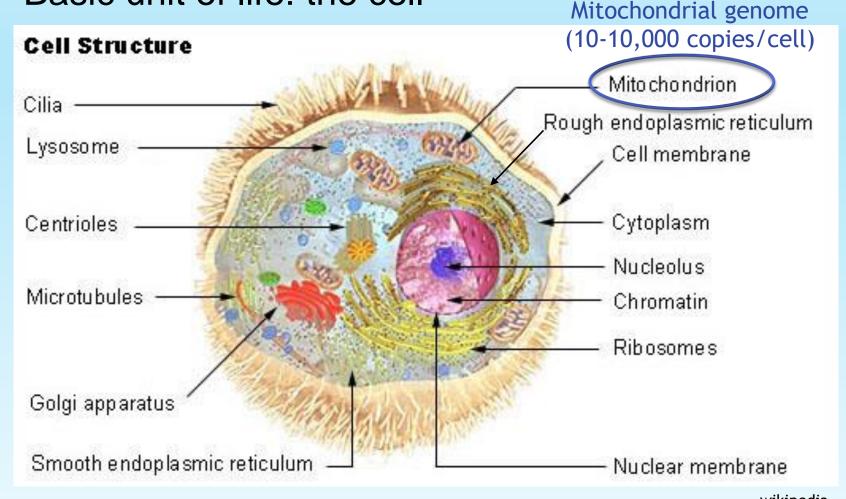
Structures / Enzymes / signaling

Nucleic acids



formation storage and transfer / ribozymes

Basic unit of life: the cell

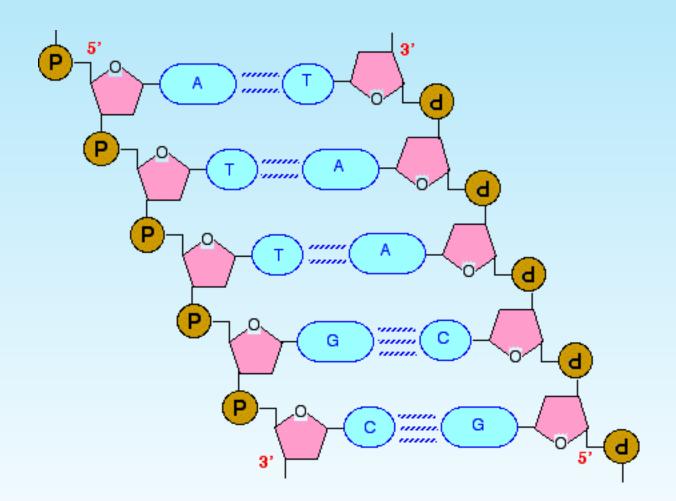


Average human = 5×10^{13} cells Typical eukaryotic cell ca. $25 \mu m$ diameter / Typical bacterium ca. $1 \mu m$ Smallest known cell (mycobacteria) = $0.1 \mu m$ Largest (ostrich egg) = 20 cm

Topics:

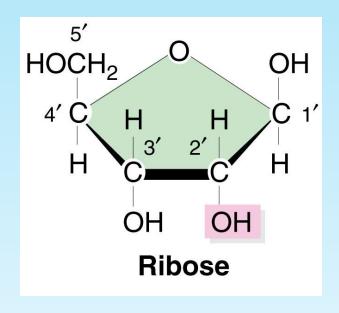
- 1. Tree of life
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3. Structure and features of DNA and RNA

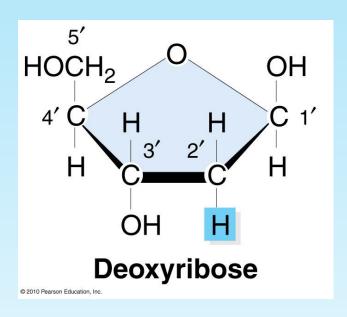


chemguide.co.uk

Sugar-phosphate backbone, bases and nucleotides



Ribose (found in RNA)



Deoxyribose (found in DNA)

- Planar sugar molecules
- Carbon atoms 1-5 (numbered clockwise, from Oxygen atom)
- Written 1' (you say 1-prime) etc to distinguish these carbon atoms from those in the bases AGCT (which you don't need to worry about)

(Deoxy)ribose-phosphate

Sugar phosphorylated at C5 = sugar-phosphate backbone of DNA / RNA

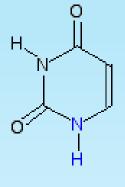
(Deoxy)ribose-phosphate + base = Nucleotide

The bases:

Pyrimidines NH2
NH2
Cytosine (C)

CH₃

thymine (T)



uracil (U)
= replaces T
in RNA

Called bases because they are alkaline in chemical terms. But this is not of importance to their function in DNA. Just call them bases!

Purines

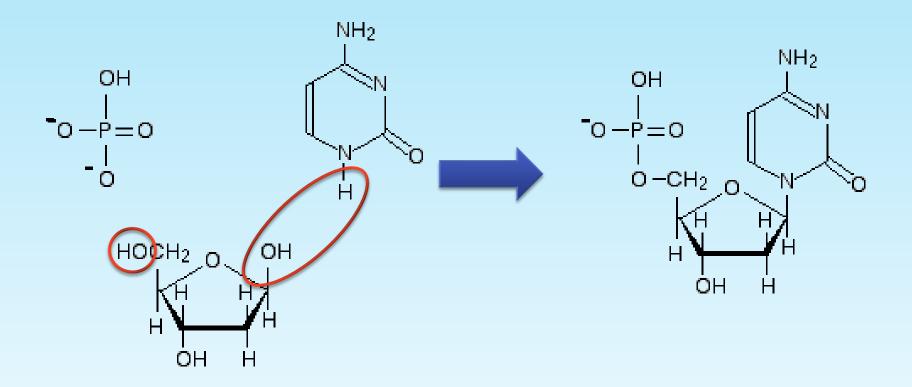
adenine (A) gi

guanine (G)

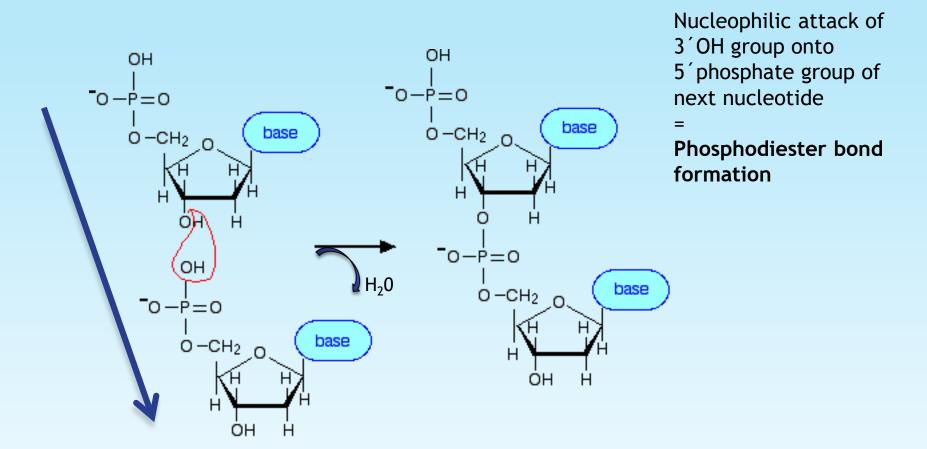
(Nitrogen atoms in blue = attach to C1´ of ribose in a nucleotide)

Phosphoric acid + sugar + base = nucleotide

(The basic repeating unit of nucleic acid)

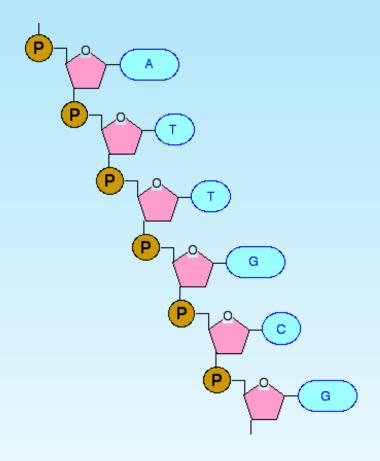


Joining nucleotides into a polynucleotide strand



IMPORTANT: Biological reaction can ONLY proceed in 5 'to 3 'direction!

Simplified view of 1 strand....



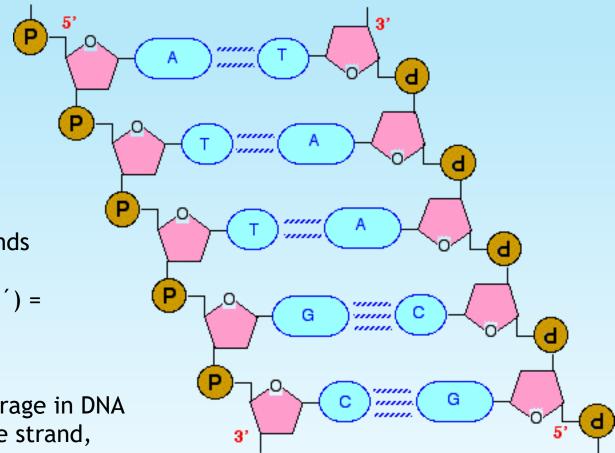
...but DNA is double stranded...

Base-pairing

- Based on hydrogen bonds (weak attractions, rather than strong chemical bonds)
- G always binds to C, A always binds to T
- G:C = 3 hydrogen bonds / A:T = only 2
- Always a purine and a pyrimidine in a pair

DNA with high GC content is more difficult to "melt" (separate the strands)

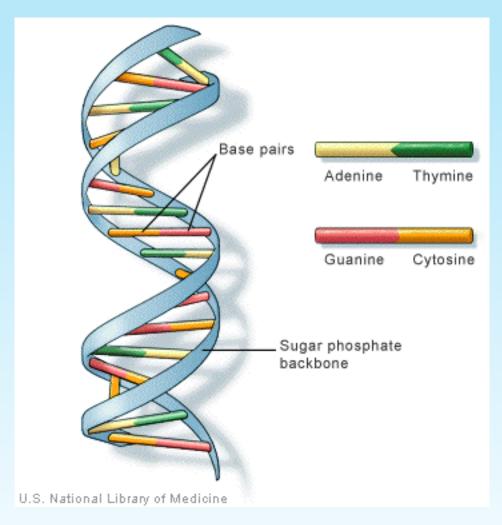
Structure of DNA

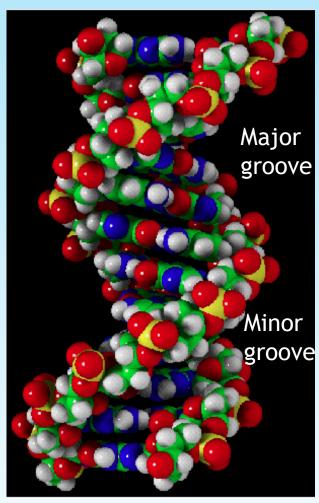


 Notice that strands go opposite directions (5´- 3´) = antiparallel.

Basis of data storage in DNA
 if you have one strand,
 you have all the
 information, and can copy
 it to double-stranded form.

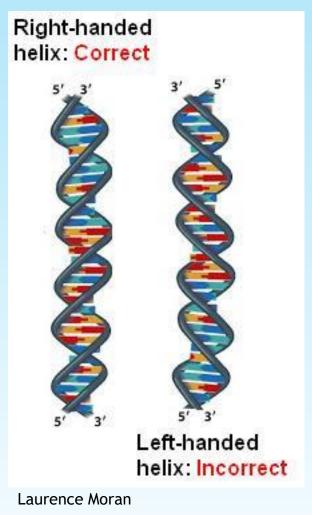
3D Structure of DNA = double helix



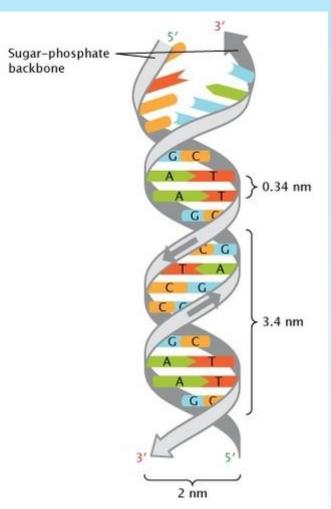


Some final details on DNA 3D structure:

It forms a right-handed helix...



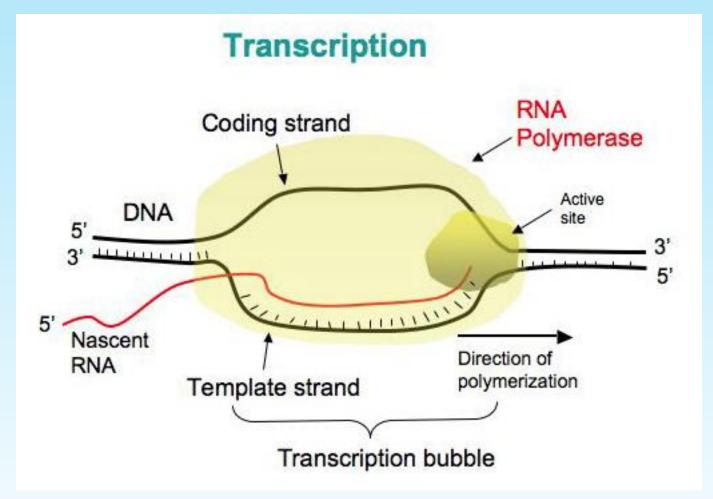
of (fairly) regular dimensions



10-10.5 bases per turn

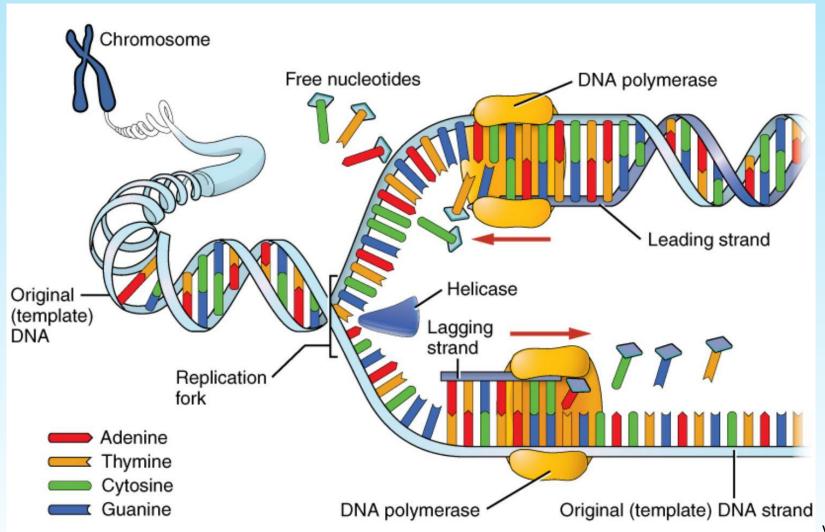
Transcription of RNA

By DNA-dependent RNA polymerase

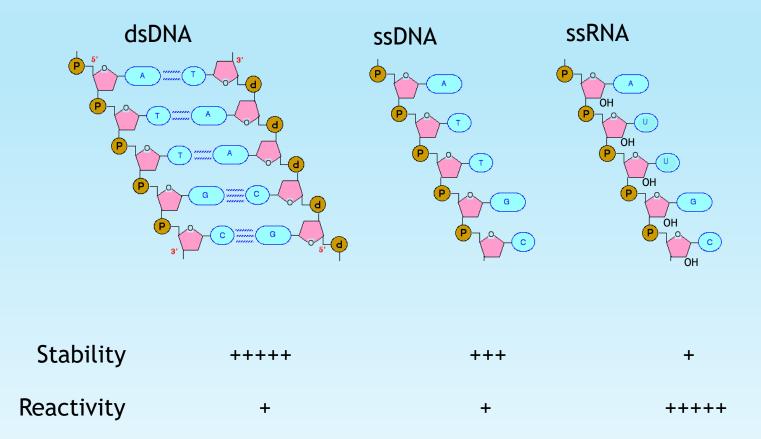


Replication of DNA

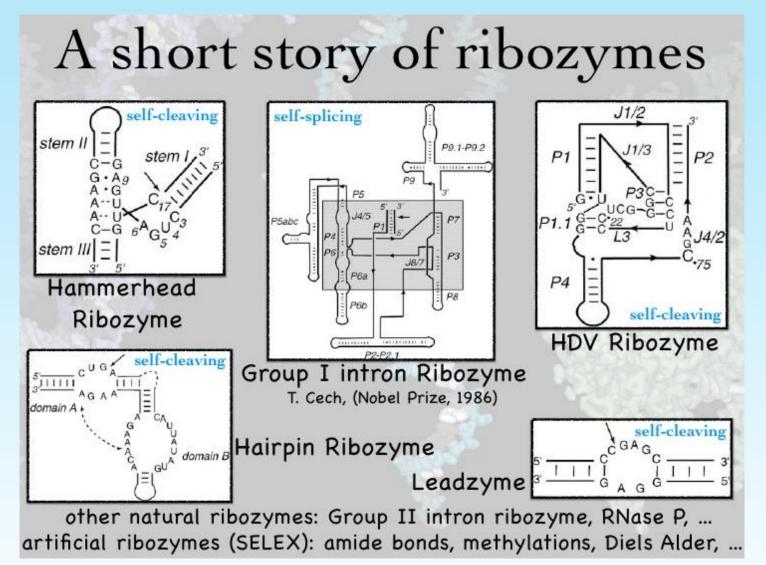
By DNA-dependent DNA polymerases

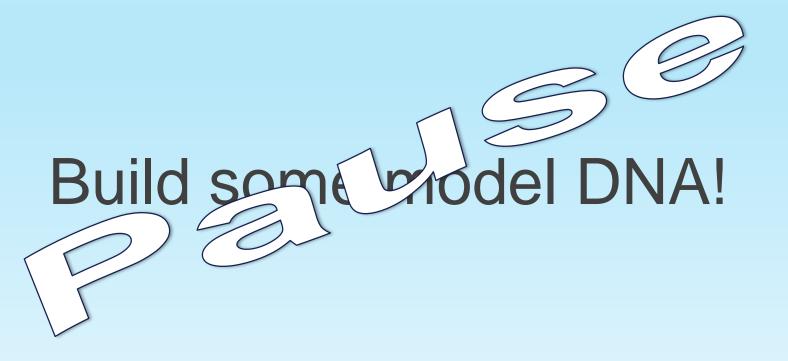


Stability / reactivity of DNA and RNA



Reactive RNA: ribozymes (i.e. RNA enzymes)





Video summary of DNA structure

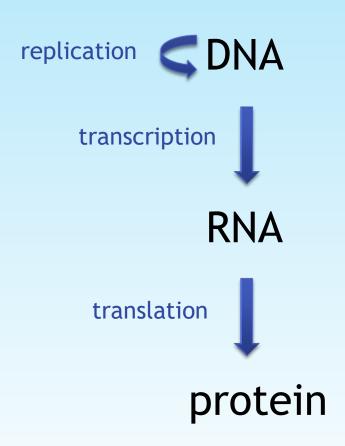
https://www.youtube.com/watch?v=o_-6JXLYS-k

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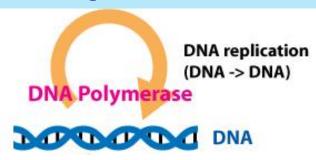
4. DNA makes RNA makes protein

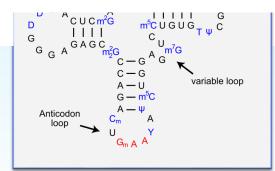
"The central dogma of molecular biology" (Attributed to Francis Crick, although not the words he used)



4. DNA makes RNA makes protein (cont.)

The central dogma in some more detail





wikipedia

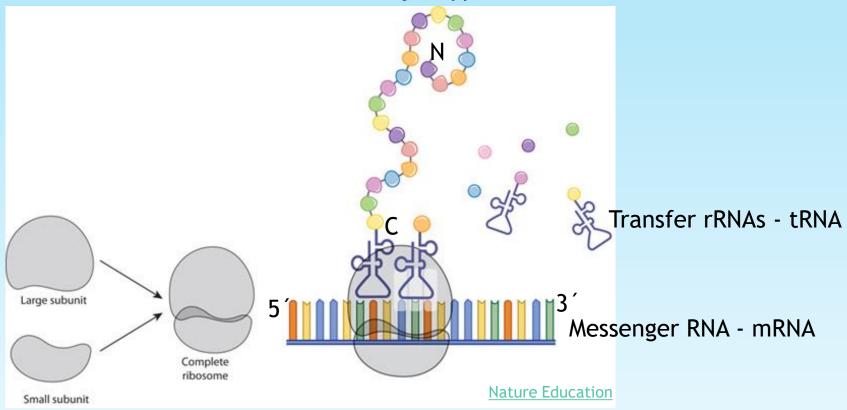
The genetic code

- Genessuse a Briefetter word (called a codon) to director protein synthesis
- There are 20 amino acids that make up all natural proteins

-	- 13-letter code with an alphabet 20fo24 letters (AGCT) = 43 = 64 po									
	codons ^u		С		Α		G		base	
_			Applettersabed	UCU	(Ser/S) Serine	UAU	(Tyr/Y) Tyrosine	UGU	(Cys/C) Cysteine	U
		UUC		UCC		UAC		UGC		С
		UUA		UCA		UAA ^[B]	Stop (Ochre)	UGA ^[B]	Stop (Opal)	A
		UUG		UCG		UAG ^[B]	Stop (Amber)	UGG	(Trp/W) Tryptophan	G
	С	CUU	(Leu/L) Leucine	CCU	(Pro/P) Proline	CAU	(His/H) Histidine	CGU	(Arg/R) Arginine	U
		CUC		ccc		CAC		CGC		С
		CUA		CCA		CAA	(Gln/Q) Glutamine	CGA		A
		CUG		CCG		CAG		CGG		G
	AUU			ACU	ACU	AAU	(Asn/N) Asparagine	AGU	(Ser/S) Serine	U
Start codon 3/1) Isoleucine				ACC	(Thr/T) Threonine	AAC	(iointy / ioparagino	AGC	(delive) delivide	С
		ALIA	(Met/M) Methionine	ACA	(TIII/T) TINGGIIIIG	AAA	(Lys/K) Lysine	AGA	(Arg/R) Arginine	A
		AUG ^[A]		ACG		AAG		AGG		G
		CUU	(Val/V) Valine G	GCU	(Ala/A) Alanine	GAU	(Asp/D) Aspartic acid	GGU	(Gly/G) Glycine	U
	G	GUC		GCC		GAC		GGC		С
	•	GUA		GCA		GAA	(Glu/E) Glutamic acid	GGA		A
		GUG		GCG		GAG		GGG		G

Translation

Occurs on the ribosome. All three major types of cell RNA involved.

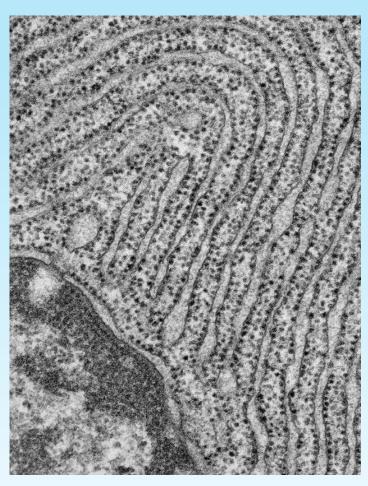


Contains ribosomal RNA - rRNA (28S, 18S and 5S)

In a typical cell, >90% of all RNA is rRNA!

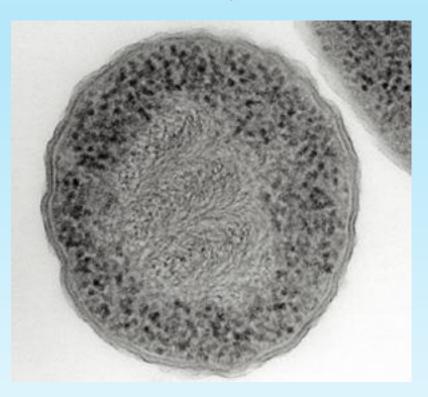
mRNA is read 5 to 3 Protein is synthesis N-terminus to C-terminus

Ribosomes



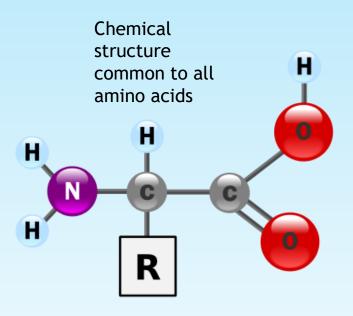
Electron micrograph of eukaryotic cell showing endoplasmic reticulum studded with ribosomes.

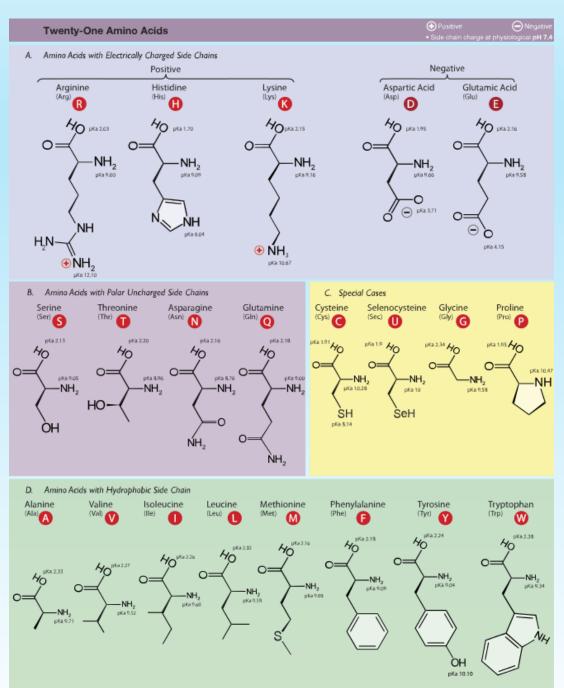
The ribosome is a ribozyme!



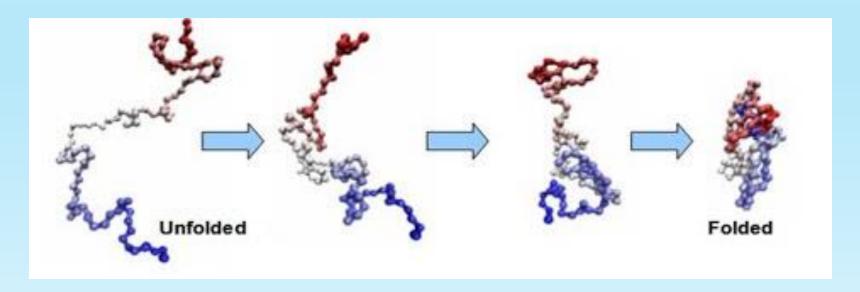
Electron micrograph of prokaryotic cell showing DNA in center and ribosomes on periphery.

Amino acids and their properties





Protein folding



Factors driving / affecting protein folding:

- Hydrophobic interactions
- Ionic interactions (+ and charges)
- Hydrogen bonds
- Metal ion binding
- Chaperone protein interactions
- Effects of salt and pH

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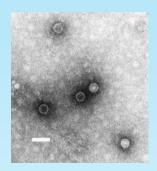
5. Genomes and genomic features

Not all genomes are of double-stranded DNA:

ssRNA viruses (e.g. Poliovirus)

dsRNA viruses (e.g. Rotavirus)

ssDNA (e.g. bacteriophage PhiX 174)



Poliovirus. Scale bar = 50 nM (credit: US EPA)

Viruses tend to have high mutation rates because:

- RNA polymerases do not have proofreading (spell checking) activity
- DNA is more stable in double-stranded form. Rearrangement and mutations more likely in ssDNA.

Genomes of dsDNA:

Ploidy (the number of sets / copies of chromosomes in a cell or organism):

Haploid = 1 copy of dsDNA genome

e.g. bacteria (typically circular genomes)*

e.g. yeast (16 linear chromosomes)

e.g. male bees, ants, wasps

Diploid = 2 copies of genome (one from mum, one from dad)

e.g. us (note that 3.2 Gb genome = haploid size)

e.g. yeast

Triploid = 3 copies

e.g. wheat

Tetraploid = 4 copies

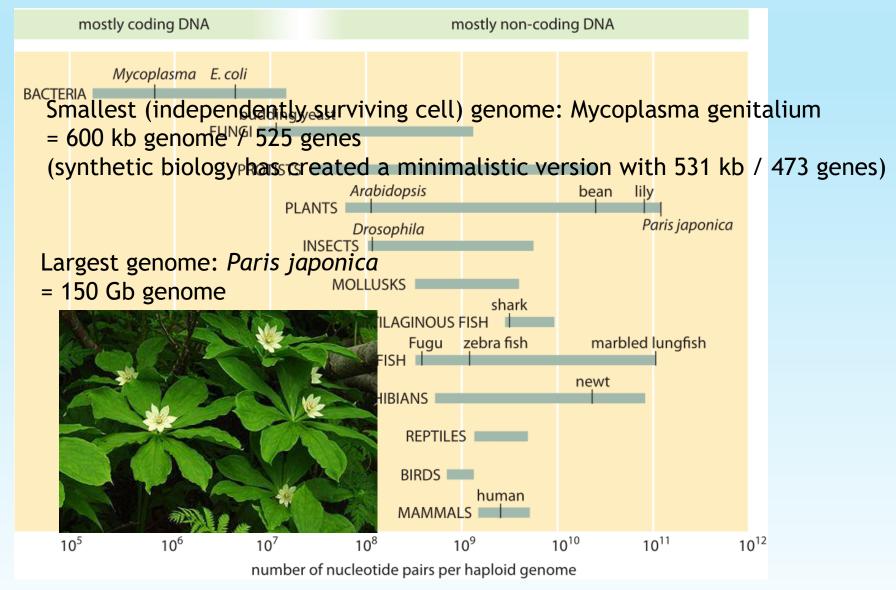
e.g. potato, Salmon

1048576-ploidy has been measured in silk gland of silkworm!

* Commonly also contain extra circles of DNA called plasmids, that can be transferred between bacteria (also between different species). Common mechanism for the spread of antibiotic resistance. Some plasmids can be present in multiple copies.

Polyploidy

Genome Sizes



http://book.bionumbers.org/how-big-are-genomes/)

Genome Sizes

For organisms that have not had their genome sequenced / have poor assemblies / lots of repetitive DNA, genome size estimated using "C number" = amount of DNA (in pg) in a haploid genome (eg a sperm cell).

1 pg = 978 Mb (ca. 1 Gb)

Database of eukaryotic genome sizes: http://www.genomesize.com/

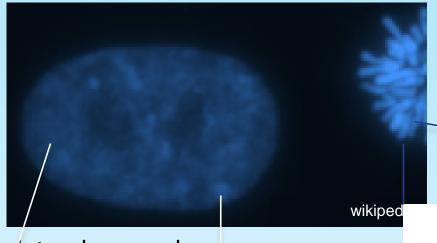
NOTE:

Large Genome ≠ many genes!

(generally means lots of repetitive and non-coding DNA)

Genomic structure and features

Cells stained with DNA-dye DAPI



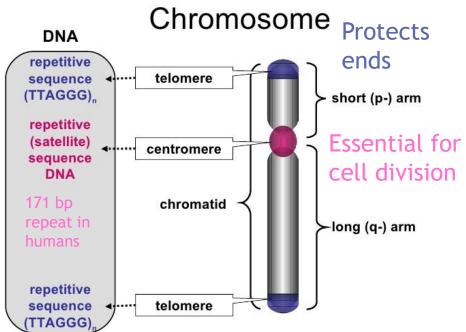
Nucleus undergoing mitosis (cell division)

Interphase nucleus (how nuclei look most of the time)

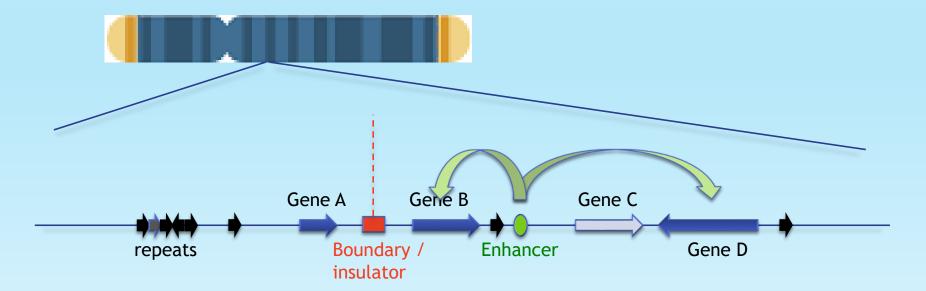
Euchromatin (open, expressed)

Heterochromatin (dense, repressed)

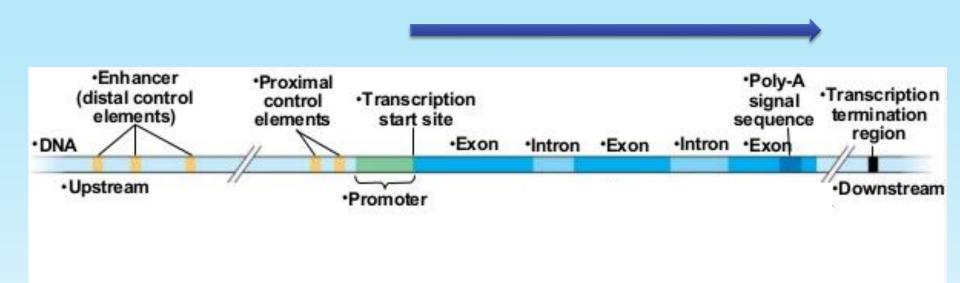
DNA in cells = protein bound = Chromatin



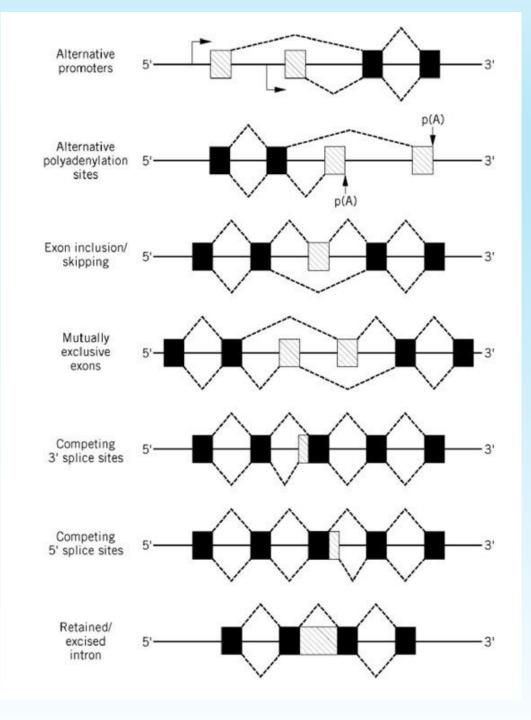
Genomic structure and features



Genes in more detail

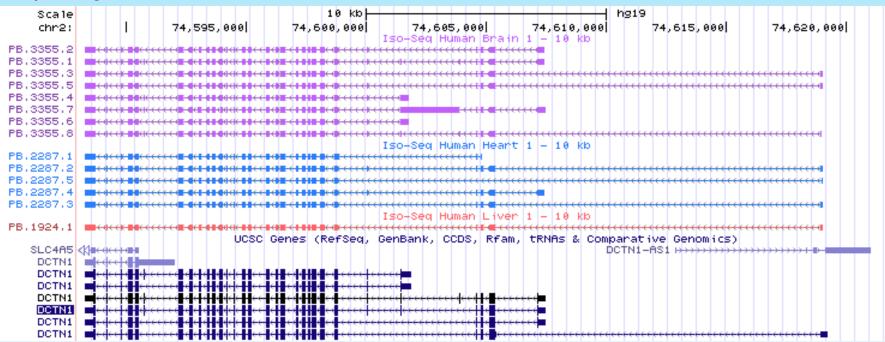


Splicing in more detail



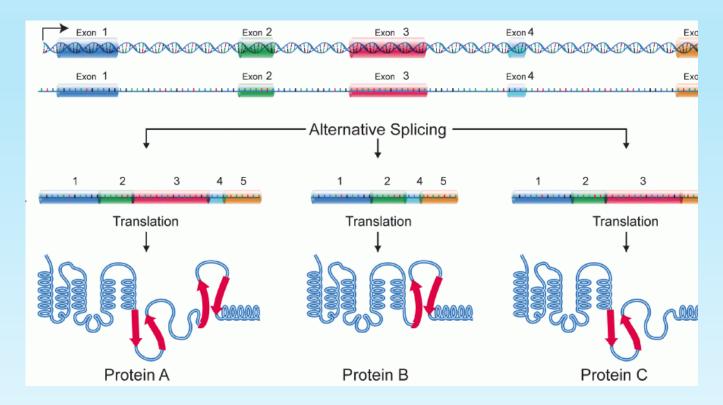
Splicing – one example

https://genome.ucsc.edu/



Note: Only ca. 1.5% of the human genome codes for protein!

Splicing: one gene = many proteins

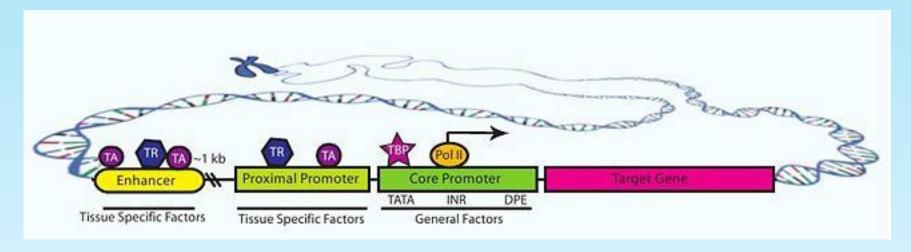


Average human gene = 8.8 exons Average exon size ~160 bp

To add to the complexity of the proteome: Most proteins can be post-translationally modified (acetylated, phosphorylated etc.).

Promoters

(of protein-coding genes = Pol II transcribed)*



ca. 1000 bp ca. 200 bp

Promoter

TATA = sequence typically found in ca. 50% promoters (binds TBP; TATA binding protein)

INR = initiator element, where Polymerase binds

DPE = downstream promoter element

*

Pol I = transcribes the two large rRNAs (18S and 28S) Pol III = transcribes small rRNA (5S),

tRNAs, and other small RNAs.

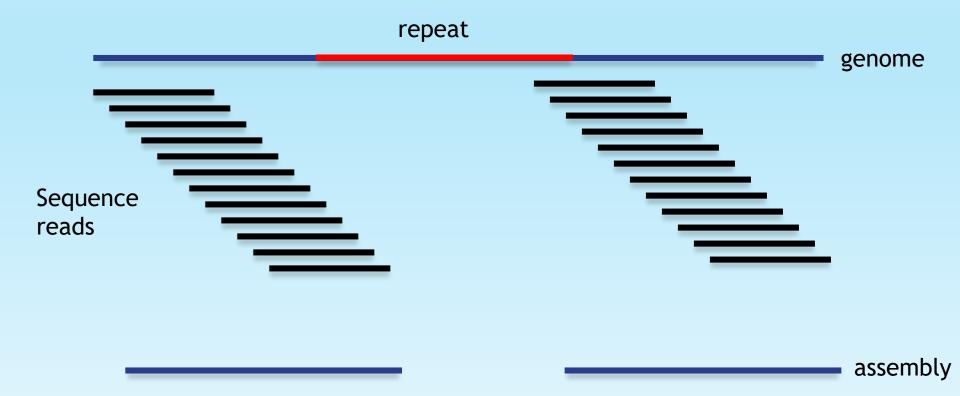
Repeats

- 1. "Microsatellites". Very short, < 10 bp, (particularly dinucleotide, trinucleotide and telomeric (6-8 bp)
- 2. "Minisatellites". 10-50 bp (eg centromeric repeats)

Also called SSRs, STRs and VNTRs. Used in forensics.

- 3. Transposons and retrotransposons.
- "Copy and paste" genetic elements
- Retro = Transcribed to RNA -> copied back into DNA at new location
- May encode enzymes to do so (many copies are dysfunctional remnants)
 e.g. SINE and LINE elements
- 42% of the human genome is derived from retrotransposons!

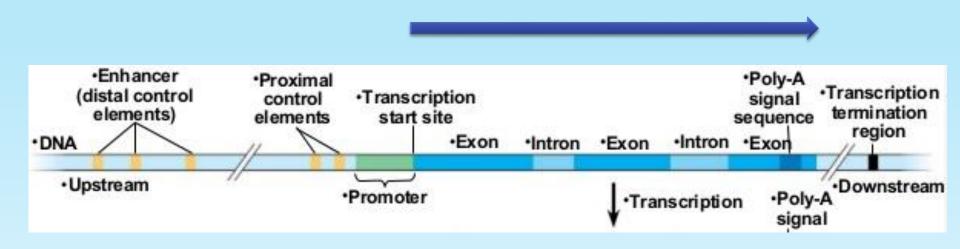
e.g. problem of repeats in genome assembly



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6. Genetics: What is a gene?



- You can also have non-coding genes (RNAs that do not contain a reading frame encoding protein) eg. rRNA, antisense transcripts, Xist
- miRNAs sequences are often transcribed in UTR sequences
- Poly-cistronic RNAs (common in bacteria) = multiple gene sequences in one mRNA.

What is a locus?

Locus = place in the genome where a particular genetic feature resides.

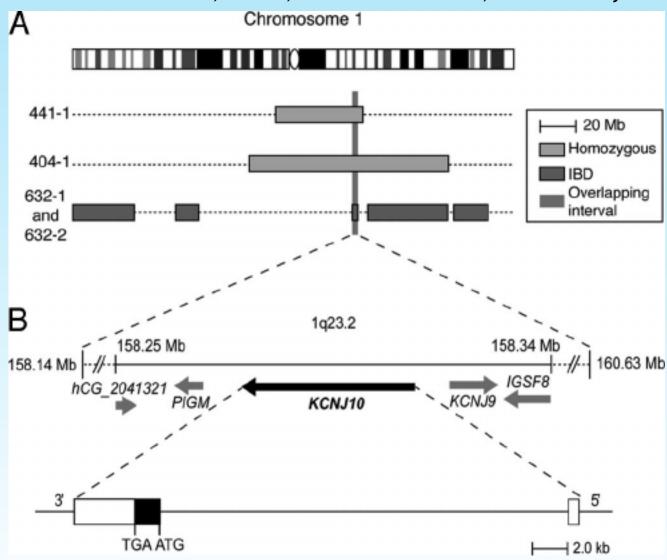
If the gene responsible for a particular trait (eg a disease) has not been identified, we may still know the locus resides on a particular area of a chromosome.

A locus can contain more than one gene, and there can be more than one locus for a kind of gene in the genome. e.g.

- the rDNA loci = in humans, 5 chromosomes carry the rDNA each contains many copies of the rRNA gene (about 200 copies in total).
- HLA locus (important in immunology) = on chromosome 6, >200 genes
- The locus for Down's syndrome = entire chromsome 21

E.g. SESAME* syndrome locus eventually mapped to *KCNJ10* gene

*seizures, sensorineural deafness, ataxia, mental retardation, and electrolyte imbalance



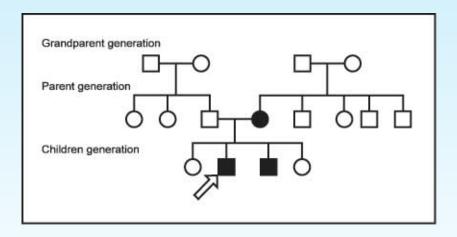
What is an allele?

= A variant of any one gene.

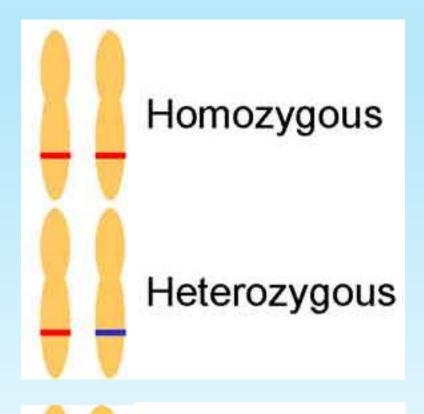
Eg a gene where one variant encodes blue eyes, the second variant encodes brown eyes, has two alleles.

The variants can be single nucleotide polymorphisms (SNPs), larger differences in sequence, copy number variants (CNVs), splice variants, insertions / deletions (indels).

Often discussed in terms of heredity



Zygosity



Both alleles the same

Two different alleles

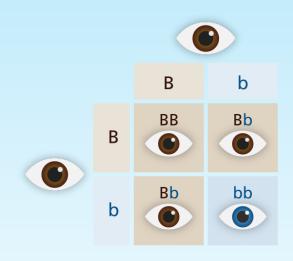


Only one allele possible

Dominant and recessive alleles

Since we have 2 copies of our genes (diploid):

- If both alleles at a given locus contribute to **phenotype** = co-dominance
- If only one allele is displayed in phenotype = dominant allele
- For a recessive allele to display phenotype, both copies must be the recessive allele.



Genome Research Limited

B - dominant brown eye allele
b - recessive blue eye allele
BB ● brown eyes
Bb ● brown eyes
bb ● blue eyes

Mutations, variants and polymorphisms

Molecular genetics

AGCTTGCATTGAATCG

Process by which variation arises = mutation

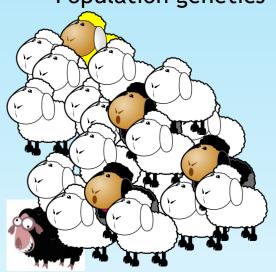
AGCTTGCAATGAATCG

Benign, no disease = variant

AGCTTGCAGTGAATCG

Disease-causing = mutation

Population genetics

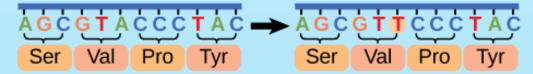


Black coat variant (arose by mutation several generations ago) is a polymorphism **inherited** by a minority of individuals. This flock also contains two mutants that have arisen spontaneously. Only one looks damaging.

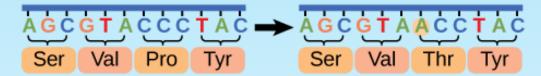
Silent, missense and nonsense mutations

Point Mutations

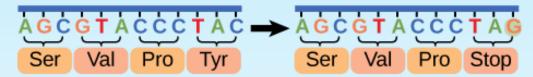
Silent: has no effect on the protein sequence



Missense: results in an amino acid substitution



Nonsense: substitutes a stop codon for an amino acid



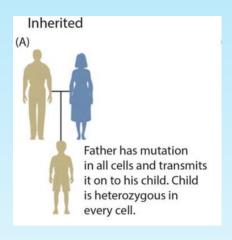
Frameshift Mutations

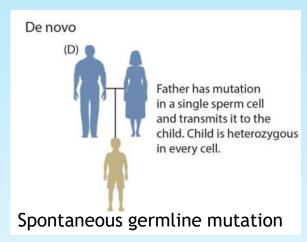
Insertions or deletions of nucleotides may result in a shift in the reading frame or insertion of a stop codon.

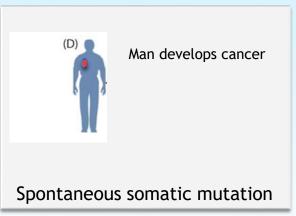


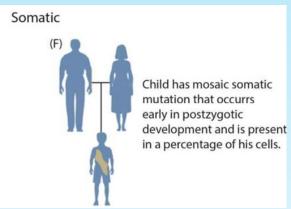
Somatic vs Germline mutations, and mosaicism

Germline = testes, ovaries (sperm and eggs) Soma = all other tissues





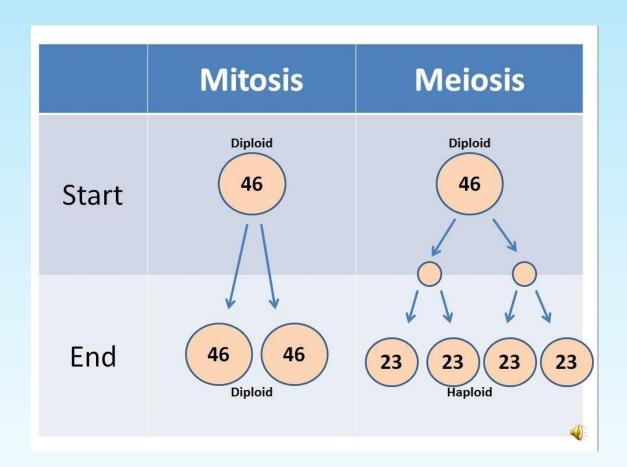




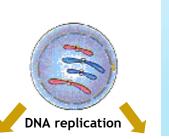
Mosaic individual.

If mutation present in germ cells, can be inherited by next generation

Difference between somatic cell division and DNA replication (mitosis) and germ-line (meiosis)

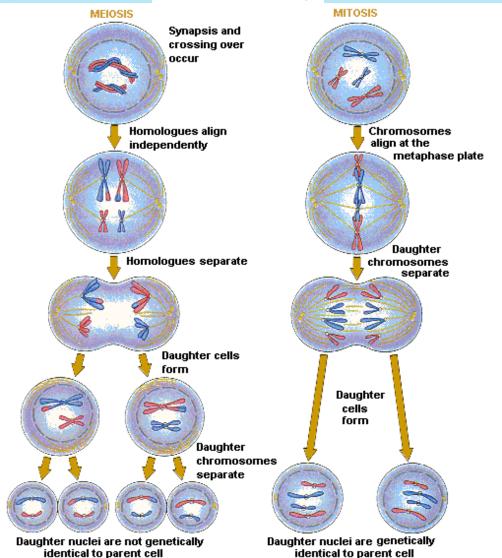


Difference between somatic cell division and DNA replication (mitosis) and germ-line (meiosis)

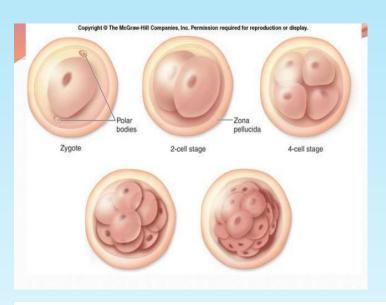


Diploid cell (n=2)

One chromosome set from mum
One chromosome set from dad



Every cell in your body contains the same DNA (almost)



Exceptions:

- Red blood cells (lose their nuclei)
- B-cells (undergo genomic rearrangement to produce different antibodies)
- Sperm / egg and precursors = undergo meiotic recombination
- Cancer cells
- Somatic mutations that accumulate as you develop and age.

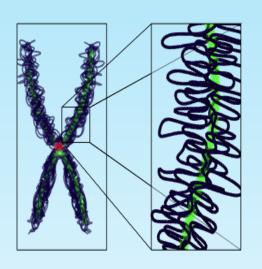


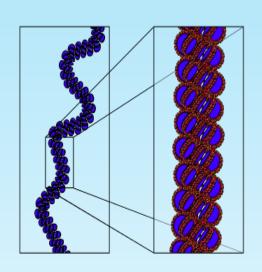
Which genes are turned on / off at a given time or in a tissue determine cell identity.

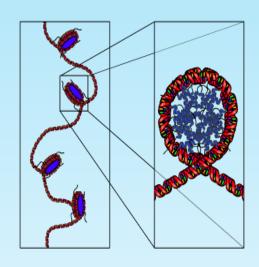
Topics:

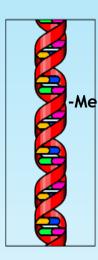
- 1. Tree of life
- 2. Building blocks of life
- 3. Structure (and differences) of DNA and RNA
- 4. DNA makes RNA makes protein
- 5. Genomes and genomic features
- 6. Genetics
- 7. Epigenetics

7. Epigenetics: Heritable traits not encoded in the DNA sequence.









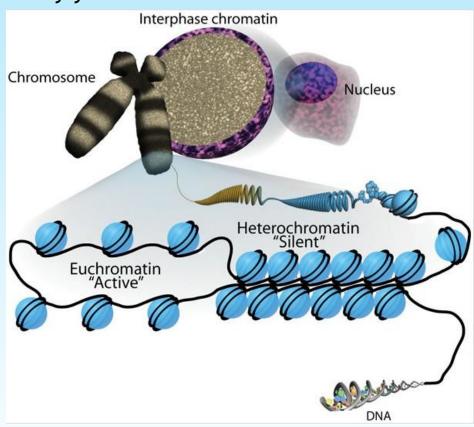
wikipedia

Mediated by:

- -DNA methylation
- -Histone variants
- -Histone covalent modification
- -Non-coding RNAs (miRNAs, long non-coding RNAs)
- -Nucleosome remodelling (positioning) and chromatin compaction
- -Nuclear position

DNA packaging (DNA + packaging proteins = chromatin)

- -Each cell in your body contains about 2m DNA (in 46 pieces i.e. chromosomes)
- All that has to be packed into a nucleus that of approx 5 µm diameter which is why you need chromatin!



All the DNA in your body (10¹³-10¹⁴ cells), if aligned end to end could stretch 5-50 times from the earth to the sun...

Histones + nucleosomes

Histone proteins:

- Small (ca. 100-140 aa)
- Highly conserved
- Package DNA

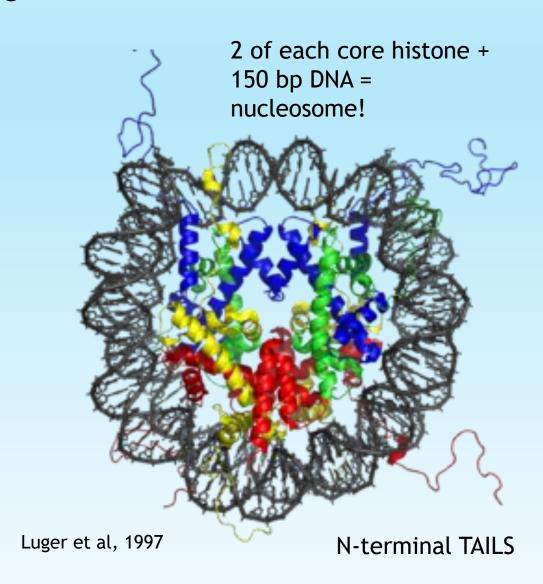
4 core histones:

H2A

H₂B

H3

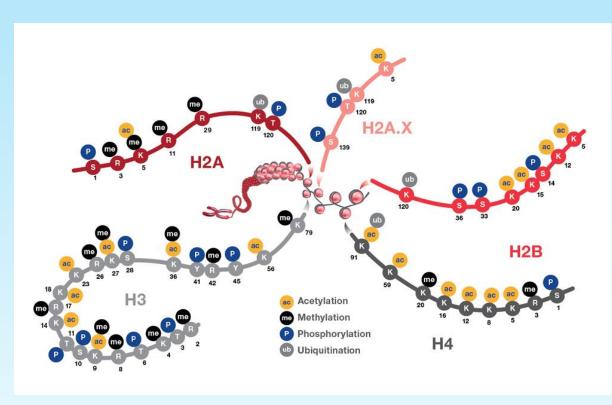
H4



Histone tails are targets for modifications

Post-translational modifications

- -acetylation
- -methylation
- -phosphorylation
- -Ubiquitylation
- -SUMOylation
- -Citrullination / deimination
- -Proline isomerisation
- -ADP-ribosylation
- -Palmitoylation
- -GlcNac



These modifications control how tightly the chromatin is packed (eg condenses to heterochromatin), attract or repulse transcription factors, signal damage etc.

DNA methylation

Cytosine

methylated Cytosine

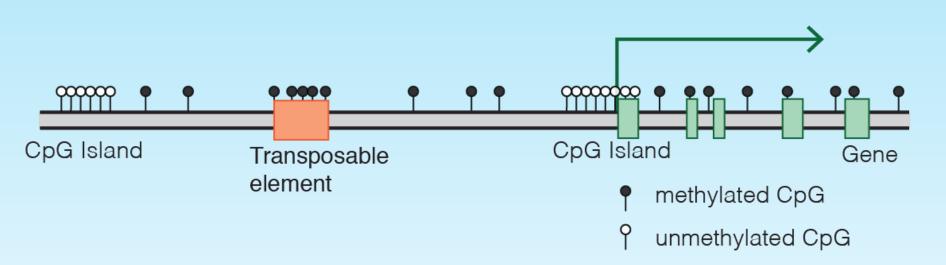
(aka 5mC)

Also get adenine methylation

- Occurs most in CG (also known as CpG) context in mammals.
- Also common in CHH and CHG in plants (H= A, C or T)
- Adenine and cytosine
 methylation common in selected
 sequence motifs in bacteria as
 an antiviral defense. Express an
 enzyme that cuts a DNA
 sequence (restriction enzyme):
 non-methylated viral DNA gets
 cut, methylated bacterial DNA
 protected.

DNA methylation in vertebrates

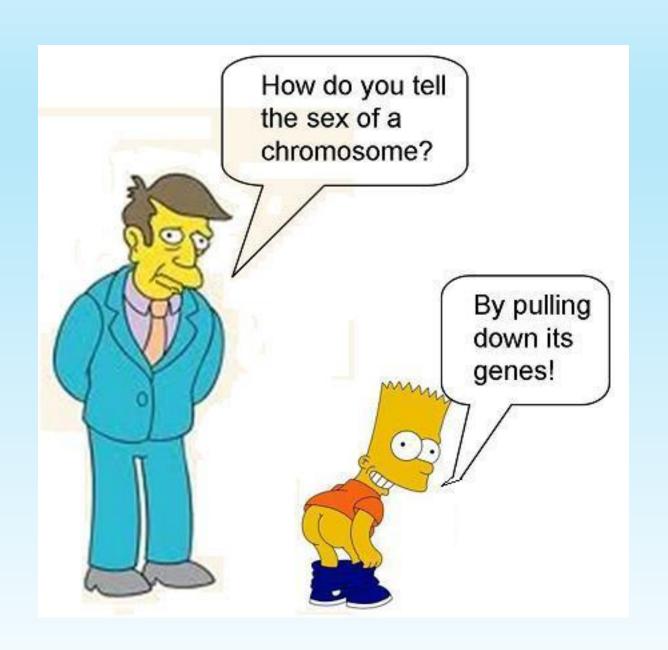
Typical mammalian DNA methylation landscape



- 5mC promotes transcriptional repression (can block TF binding directly, and can recruit proteins that cause chromatin condensation)
- Clusters of non-methylated C found at ca. 65 % of vertebrate promoters
- 5mC mutates more frequently to T, therefore gradual conversion of CG to TG in genome, except in promoters that are actively demethylated = gives rise to "CpG islands".

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Thanks for your attention

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