

CSI5126. Algorithms in bioinformatics

Essential Cellular Biology

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Version September 11, 2018

Summary

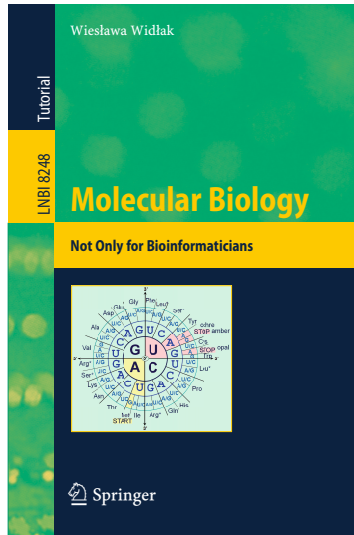
This lecture presents the **cell**, the **kinds of cells**, their **organization** and **composition**. Concepts from **molecular evolution** are introduced. It presents the **macromolecules** of the cell, with their basic organization. Throughout the presentation, we will highlight the importance of the notions for bioinformatics.

General objective

- **Describe** the organization of the cell and macromolecules

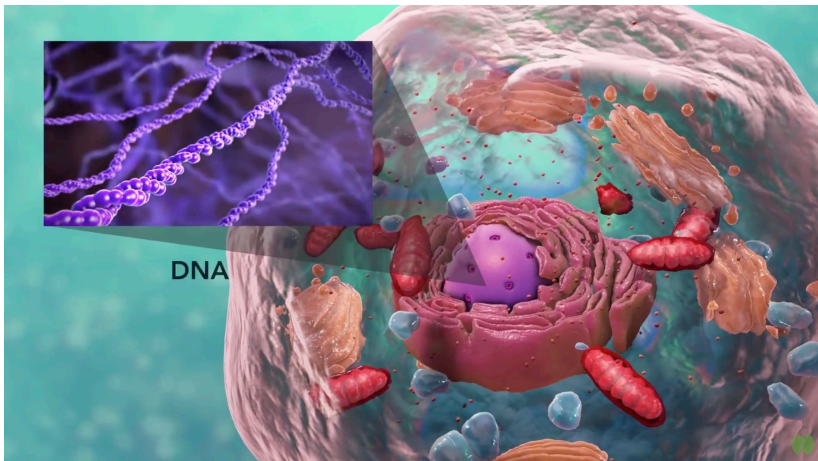
Reading

- Wiesława Widłak (2013). *Molecular Biology: Not Only for Bioinformaticians* (Vol. 8248). Springer. Chapters 1, 2, and 3.



link.springer.com/book/10.1007/978-3-642-45361-8

Cell Structure



<https://www.youtube.com/watch?v=URUJD5NEXC8>

Cells: building blocks of living organisms

Two **kinds** of cells (with and without nucleus)

Prokaryote (procaryote, prokaryotic cell, procaryotic organism):

Cell or organism **lacking** a membrane-bound, structurally **discrete nucleus** and other sub-cellular compartments. Bacteria are prokaryotes.

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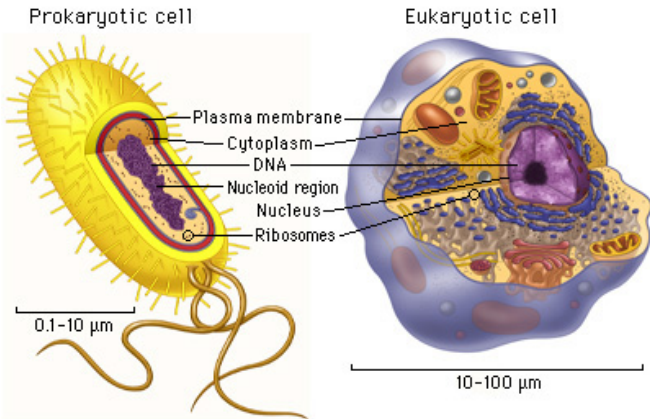
Eukaryote (eucaryote, eukaryotic cell, eucaryotic cell): Cell or organism **with** a membrane-bound, structurally **discrete nucleus** and other well-developed sub-cellular compartments. Eukaryotes include all organisms except viruses, bacteria, and cyanobacteria (blue-green algae).

Cells: building blocks of living organisms

- ❖ Eukaryotic cells are generally larger than prokaryotic cells.
- ❖ The packaging of the genetic information (DNA) is much more **structured** and compact in **Eukaryotes** compared to **Prokaryotes**.

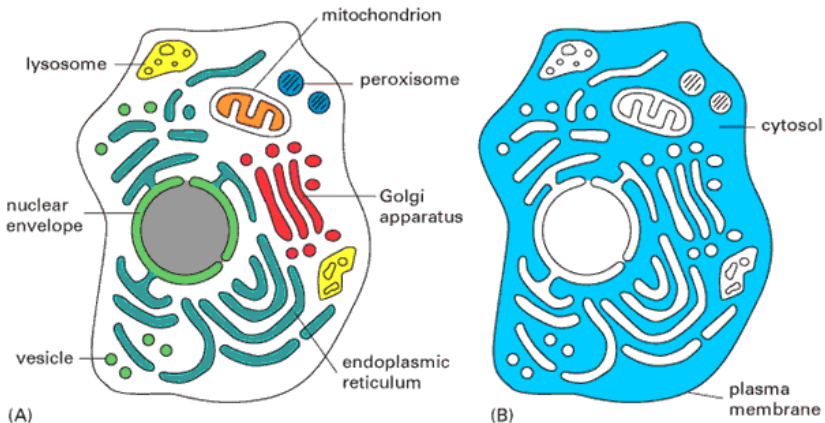
Cell theory: 1939 by Matthias Schleiden and Theodor Schwann.

Prokaryotic vs eukaryotic cell



www.phschool.com/science/biology_place/biocoach/cells/common.html

Organisation of an eukaryotic cell



Organelle genomes

- Organelles are discrete structures having **specialized functions**.

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- ❖ Several organelles are believed to be engulfed prokaryotes (**endosymbiotic theory** made popular by Lynn Margulis)
- ❖ Mitochondrial genes are **inherited from the mother only**.

Bioinformaticist's point of view

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- ❖ Eukaryotic cells being more complex provide a richer set of problems:
e.g. **protein sub-cellular localisation problem**.
- ❖ During the sequence assembly, one has to consider the possibility of contamination, mtDNA/nuclear DNA, bacterial DNA.

Resources

- ❖ Texas Education Agency
Advanced Biotechnology Collection on iTunes U
 - ❖ <https://itunes.apple.com/ca/itunes-u/tea-advanced-biotechnology/id876525204?mt=10>
 - ❖ Specifically the **Cell Structure and Function** segment
- ❖ Help Me Understand **Genetics**
 - ❖ <https://ghr.nlm.nih.gov/primer>
- ❖ **BBC** The Cell The Hidden Kingdom
 - ❖ <https://www.youtube.com/watch?v=aDuwkdQzb2g>
- ❖ <http://learn.genetics.utah.edu>

(3) kingdoms of life

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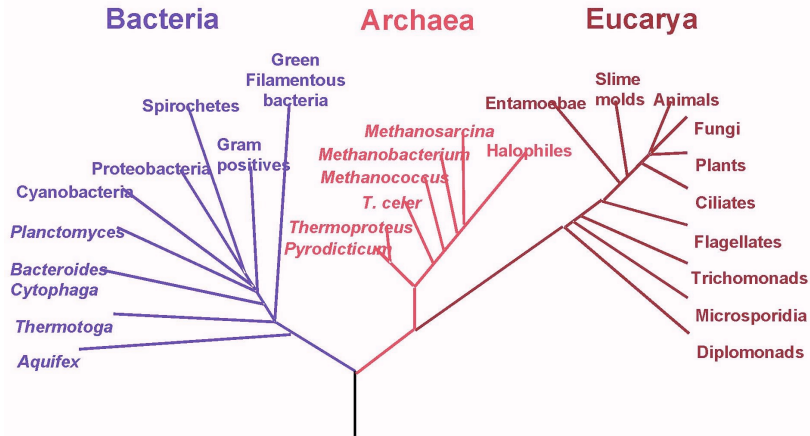
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- Archaea:** (archaebacteria) like the prokaryotes **they lack the nuclear membrane** but have **transcription and translation mechanisms close to those of the eukaryotes**.

(3) kingdoms of life: Archaea

Methanococcus jannaschii is an **methane producing archaeobacterium** which had its complete genome sequenced in 1996. This organism was discovered in 1982 in white smoker of a hot spot at the bottom of the Pacific ocean: depth **2600 meters**, **temperature 48-94° C (thermophilic)**, optimum at 85° C, 1.66 Mega bases, 1738 genes. 56% of its genes are unlike any known eukaryote or prokaryote, one kind of DNA polymerase (other genomes have several).

Phylogenetic Tree of Life



Phylogenetic tree

- ❖ “The objectives of phylogenetic studies are (1) to reconstruct the correct **genealogical ties between organisms** and (2) to estimate the **time of divergence** between organisms since they last shared a common ancestor.”

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- ❖ “The **nodes** represents the **taxonomic units**, and the **branches** define the **relationships** among the units in terms of descent and ancestry.”
- ❖ “The **branch length** usually represents the **number of changes** that have occurred in that branch.” (or some amount of time)

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Bioinformaticist's point of view

- ❖ **Bench-marking** (cross-validation) and **molecular evolution**
- ❖ **Molecular sequence alignment** : are the sequences evolutionary related?
- ❖ **Large phylogeny problem**: Reconstructing phylogenetic trees from molecular sequence data
- ❖ **Small phylogeny problem**: Reconstructing ancestral molecular sequences

Nothing in Biology Makes Sense Except in the Light of Evolution

Theodosius Dobzhansky

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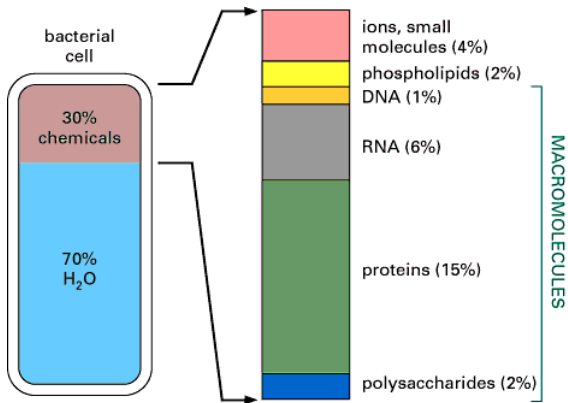
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- ❖ Virus that infect bacteria are called phages or bacteriophages.
- ❖ Viroids don't even have a capsid – consists of a single-stranded RNA.

Composition of the Cell



⇒ **DNA**, **RNA** and **proteins** will be the main focus of the course.

Macromolecules: DNA (deoxyribonucleic acid), RNA (ribonucleic acid) and Protein

Bioinformatics is mainly concerned with three classes of molecules:

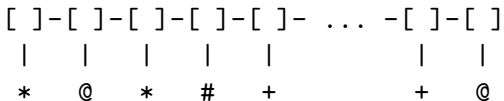
- **DNA, RNA and proteins** — collectively called **macromolecules** or **biomolecules**.

Macromolecules: DNA (deoxyribonucleic acid), RNA (ribonucleic acid) and Protein

All three classes of macromolecules are **polymers**, that is they are composed of smaller units (molecules), called **monomers**, that are **linked sequentially** one to another forming **unbranched linear structures**.

Macromolecules: DNA (deoxyribonucleic acid), RNA (ribonucleic acid) and Protein

Generally speaking, the units (monomers) consists of two distinct parts, one that is **common** to all the monomers and defines the **backbone** of the molecule, and another part that confers the **identity** of the unit, and therefore its **properties**.



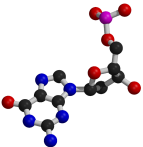
DNA's building blocks: ACGT



Adenine (A)



Cytosine (C)



Guanine (G)

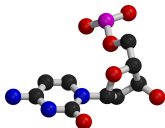


Thymine (T)

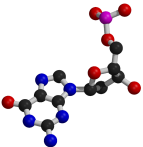
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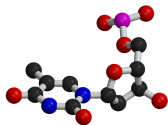
Adenine (A)



Cytosine (C)



Guanine (G)



Thymine (T)

⇒ **Identify** the common and unique parts of each monomer.

(20) Amino Acids (Naturally Occuring)



A (Ala)



D (Asp)



E (Glu)



K (Lys)



P (Pro)



W (Trp)



V (Val)



R (Arg)



C (Cys)



G (Gly)



I (Ile)



M (Met)



S (Ser)



Y (Tyr)



N (Asn)



Q (Gln)



H (His)



L (Leu)



F (Phe)



T (Thr)

⇒ Stick (licorice) representation.

Structure

It's useful to distinguish between four **levels of abstraction** or **structure**: **primary**, **secondary**, **tertiary** and **quaternary** structure.

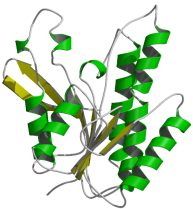
1, 2, 3, ...

EAR^RVLV^YGGRGALGSR^CVQ^NW ... (236) ...

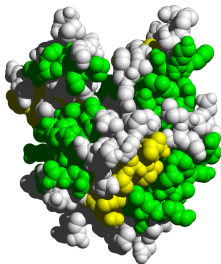
(a) primary structure



(b) secondary structure



(c) tertiary structure - ribbon



(d) tertiary structure - all atoms

Bioinformaticist's point of view

- ❖ A large number of computational problems are related to the **primary sequence**: sequence assembly, sequence alignment, phylogenetic tree inference, gene-finding, sequence motif discovery, etc.
- ❖ Predicting the **secondary, tertiary, and quaternary (docking) structure** are problems, on its own.
- ❖ These **abstractions** are allowing us to formulate efficient algorithms - understanding the implications is paramount.

Macromolecules: DNA (deoxyribonucleic acid), RNA (ribonucleic acid) and Protein

The primary structure or **sequence** is an ordered list of characters, from a given alphabet, written contiguously from left to right.

DNA : 4 letters alphabet,

$$\Sigma = \{A, C, G, T\}$$

RNA : 4 letters alphabet,

$$\Sigma = \{A, C, G, U\}$$

Proteins : 20 letters alphabet,

$$\Sigma = \{A, C, D, E, F, G, H, I, K, L, M, N, P, Q, R, S, T, V, W, Y\}$$

Examples

In the case of **nucleic acids** (DNA and RNA), the building blocks are called **nucleotides**, whilst in the case of **proteins** they are called **amino acids**.

Examples of DNA, RNA and protein sequences.

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> Chimpanzee Chromosome 1; A DNA sequence (size = 245,522,847 nt)
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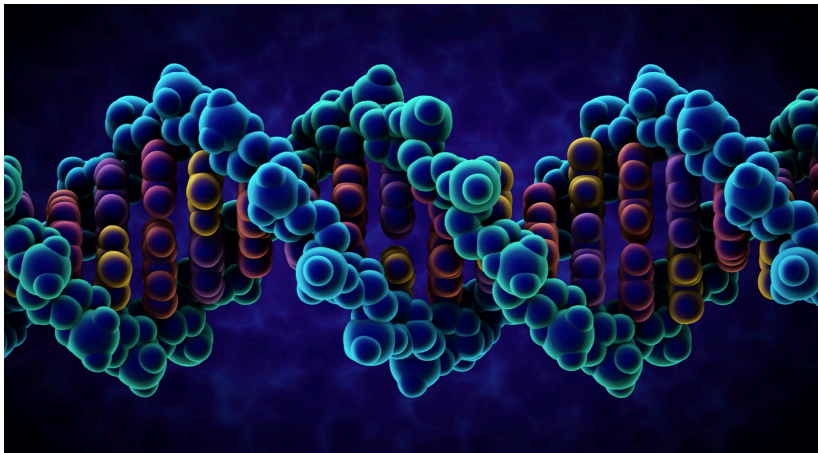
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> A01592; An RNA sequence (coding Beta Globin gene) (size = 441 nt)
AUGGUGCACCUGACUCCUGAGGAGAAGUCUGC ... GCAAGGUGAACGUGGAUGAAGUUGGUGGUG

Bioinformaticist's point of view

- ❖ **Exact string** (sequence) comparison, **approximate matching** (k -mismatches), comparison under the **edit-distance**, **significance** of match, **multi-way** sequence comparison
- ❖ Finding **repeats**, **approximate repeats**, finding interesting **patterns**
- ❖ Secondary, tertiary and quaternary structure inference

DNA



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

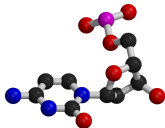
Deoxyribonucleic acids (**DNA**)

- ❖ **DNA** was discovered by **Johann Friedrich Miescher** in **1869**. Who discarded the possibility that DNA might be related to heredity!
- ❖ The **double-helical structure** of DNA was proposed in **1953** by James Watson and Francis Crick (who died on July 28, 2004).
- ❖ This discovery is often referred to as the **most important breakthrough in biology of the 20th century**.
- ❖ The proposed model finally explained Chargaff's rule (same amount of adenine and thymine, same amount of guanine and cytosine).
- ❖ More importantly, the model finally explains **how DNA and heredity are linked!** (replication)

DNA's building blocks: ACGT



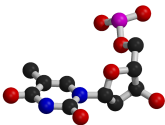
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DNA/RNA's building blocks

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- ❖ In the case of **RNA**, the bases are Adenine (**A**), Cytosine (**C**), Guanine (**G**) and Uracil (**U**).

DNA/RNA's building blocks

- ❖ The length of a DNA/RNA molecule is often expressed in **bases**, e.g. a 10 **mega base** long region.
- ❖ Or, since nucleic acids molecules **hybridize** (bind together) to form a duplex (double helical) structure, the length of a molecule is often expression is base pairs to avoid confusion, e.g. a 10 **mega base** pairs region.

DNA/RNA's building blocks

- ❖ **DNA** stands for deoxyribonucleic acid, and **deoxy** comes from the fact that the **C2' carbon of the sugar has no oxygen**; while RNA has one. **RNA's O2' oxygen is key to its functional versatility!**
- ❖ The other difference is the use of **T** (thymine) in the case of **DNA** vs **U** (uracil) in the case of **RNA**.
- ❖ Nucleotides are **always attached one to another in the same way** (well, almost always): the **C3' atom** of the nucleotide i is covalently linked to the **phosphate group** of the nucleotide $i + 1$.

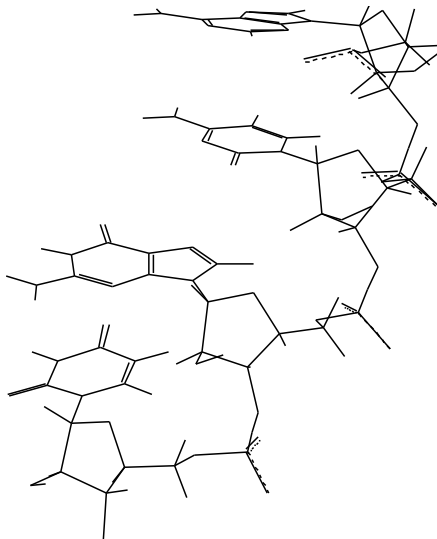
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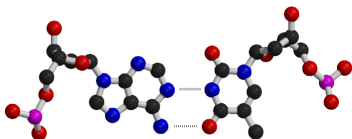
DNA/RNA's building blocks

- ❖ The **orientation of a DNA molecule** is important; just like the orientation of words are important in natural languages.
- ❖ The convention is to enumerate the string from its **5' end**; this correspond to the order into which information is process for certain key steps, to be described later. The features that are occurring **before the 5'** are said to be **upstream** while those occurring after the **3'** end are **downstream**, upstream and downstream signals.

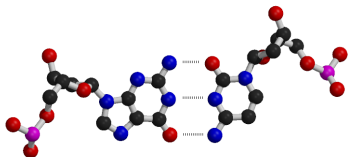
DNA strand



Watson-Crick (Canonical) base pairs



(Adenosine) A : T (Thymine)



(Guanine) G : C (Cytosine)

⇒ One of the two base pairs is stronger than the other, which one?

Watson-Crick (Canonical) base pairs

In the case of **DNA**, bases interact, i.e. form hydrogen bonds, primarily through the following set of rules:

- ❖ **A** interacts with **T** (and vice versa)
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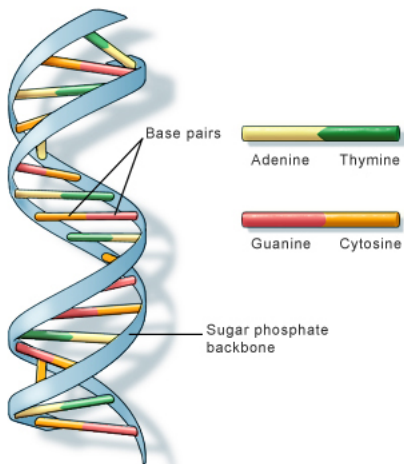
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Those rules are the consequence of the fact that A:T and G:C pairs position the backbone atoms roughly at the same three-dimensional location and therefore both produces the same double helical structure; isosteric base pairs.

- ❖ **DNA** molecules generally form **right-hand** side helices in **B form**, while **RNA** are **A form**, also right-hand side. A left-hand side helix exists that is called **Z DNA**.
- ❖ **DNA** molecules **cannot exist as a single strand**, they are degraded, i.e. cut into pieces.
- ❖ A **DNA** molecule is made of **two complementary strands running in opposite directions**.



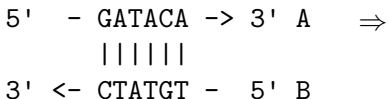
U.S. National Library of Medicine

DNA and Heredity

- + **DNA structure explains how information can be copied from one generation to the next**, or simply from one parent cell to its daughter cells during replication.

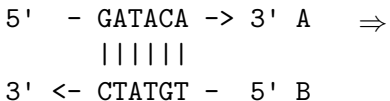
A is as a template to produce **B'**

Before replication

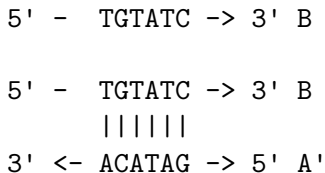


DNA and Heredity

Before replication

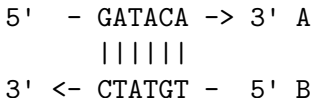


B is as a template to produce **A'**

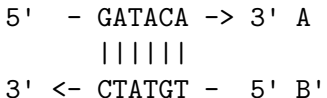


DNA and Heredity

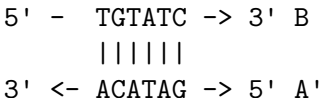
Parent cell (AB)



Daughter cell AB'



Daughter cell A'B

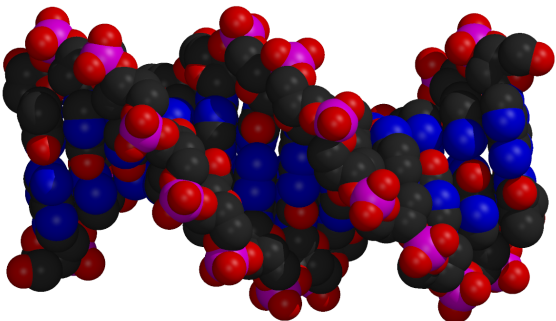


Two daughter cells, identical to their parent.
(semi-conservative process)

Remarks

- ❖ Complex organisms are growing from a single cell to billions of cells. Each cell contains an **exact copy** of the **DNA** of its **parent cell**.
- ❖ The information is redundant, the information on the second strand can be inferred from the information on the first strand. This is the basis of **DNA repair mechanisms**. A base that is deleted can be replaced. A mismatch can be detected.

CPK representation of a fragment of a DNA helix (B form)



TAAGTTATTA

|||||

ATTCAATAAT

... (580,074 bp) ...

AAAAAATAC

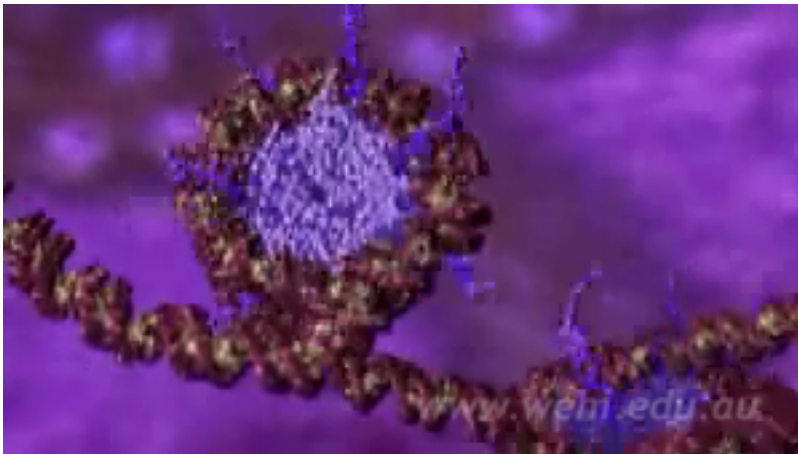
|||||

TTTTTTTATG

About CPK

CPK stands for **Corey-Pauling-Koltun** representation. **Every atom** is represented as a sphere, with **radius proportional to its van der Waals radius**. The usual color scheme is to represent carbon atoms in black, nitrogen in blue, oxygen in red and phosphorus atoms in pink.

Chromosome



<https://youtu.be/OjPcT1uUZiE?list=PLD0444BD542B4D7D9>

About the animation

- ❖ **Histone proteins** attach to the **DNA**.
- ❖ **Histones interact one with another** to form a complex called **nucleosome**, but also forcing the DNA to wrap around it.
- ❖ The histone, nucleosome and DNA models were derived from their PDB (<http://www.rcsb.org/pdb/>) structures and other published data.

About the animation

- ❖ **Histone proteins** attach to the **DNA**.
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- ❖ The histone, nucleosome and DNA models were derived from their PDB (<http://www.rcsb.org/pdb/>) structures and other published data.
- ❖ Macromolecular structures cannot be directly observed. A molecular bond is between 1 and 2 Å (angstrom – 10^{-10} m) long, wave length in the visible spectrum are 400 to 700 nm (10^{-9} m).

Bioinformaticist's point of view

- ❖ Given DNA sequence information alone, **predict the locations where the histones will be binding.**
- ❖ Knowing the location of the histones might help predicting the **location of genes** as well as the **location of regulatory elements.**
- ❖ The **three-dimensional organization of the genome** is a hot topic.

Summary

- ❖ Two kinds of cells: **prokaryotic** and **eukaryotic**.
- ❖ **Eukaryotic cells have organelles**, and some organelles, such as the mitochondria, contain DNA.
- ❖ Three Kingdom of life: **Prokarya**, **Eukarya**, and **Archea**
- ❖ A **phylogeny** specifies the relationships between organisms and time of divergence.
- ❖ Three kinds of macromolecules: **DNA**, **RNA**, and **proteins**.
- ❖ **Macromolecules are linear (unbranched) polymers**, such that all the monomers have a common and a specific part (remember the analogy with the linked nodes).

References



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Pensez-y!

L'impression de ces notes n'est probablement pas nécessaire!