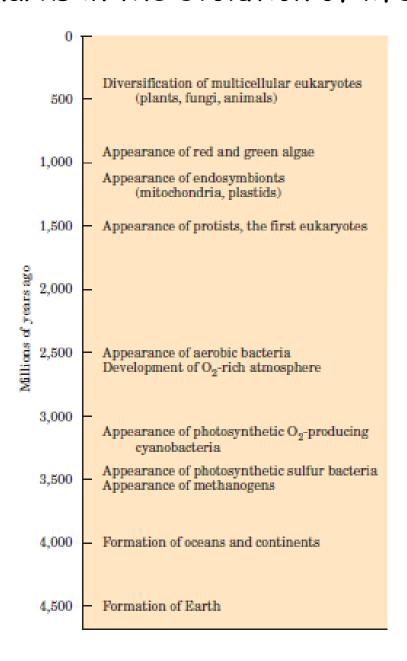


What is Life?

What are the principles of Biology?

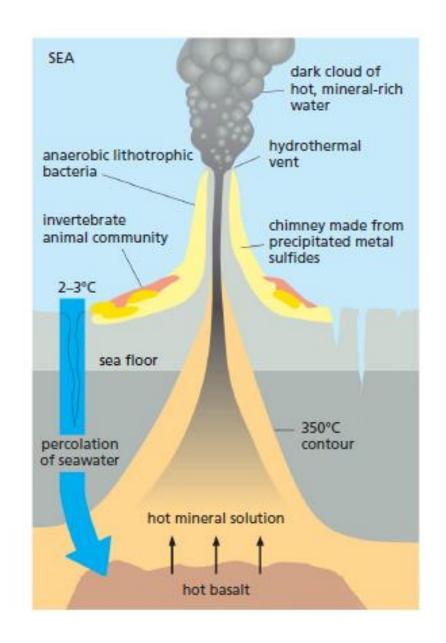
Landmarks in the evolution of life on Earth



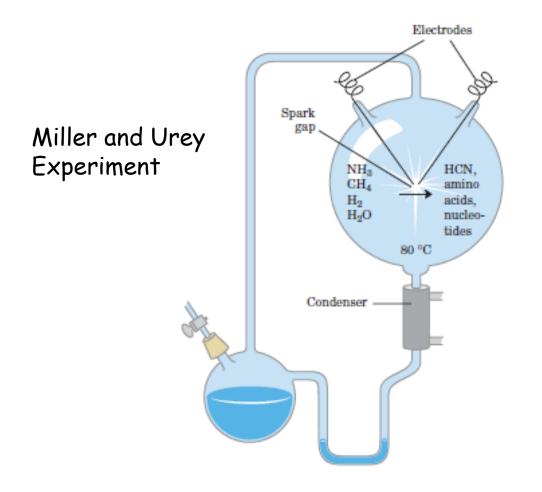
Life's Origins

Today, research into the origin of life is interdisciplinary with workers trying to answer four main questions:

- 1. What was the Earth's physical environment like when life first evolved?
- 2.What sorts of chemical reactions could produce the building blocks of life and could these occur naturally in the early Earth's environment?
- 3. How could the complex organic molecules be compartmentalized into a contained unit?
- 4. How did the genetic code evolve?



How did the first living organisms acquire their characteristic organic building blocks?



Hypothesis:

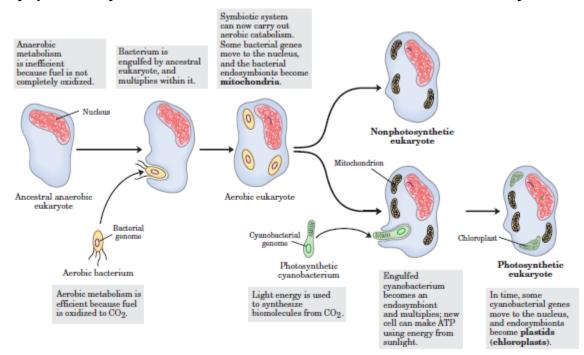
Effects of powerful atmospheric forces—ultraviolet irradiation, lightning, or volcanic eruptions—on the gases in the prebiotic Earth's atmosphere, and on inorganic solutes in superheated thermal vents deep in the ocean.

The appearance of the first living cell???

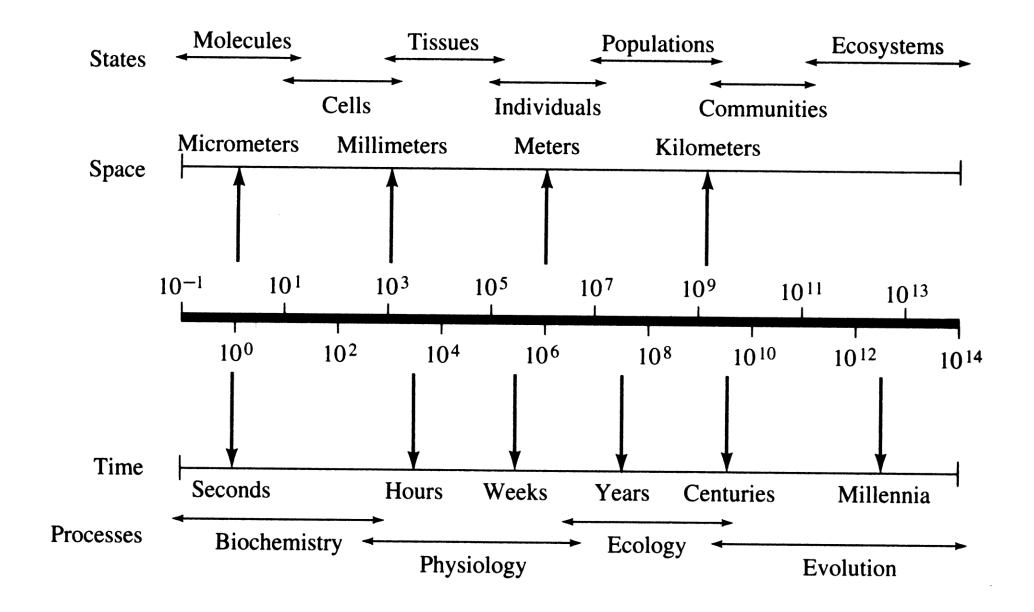
The First Cell Used Inorganic Fuels???

Lynn Margulis and the Theory of Endosymbiosis

- Eukaryotic cells originated from a series of endosymbiotic events involving multiple prokaryotes
- This idea was considered outrageous at the time (1967), but many of Margulis's ideas have since become widely accepted
- Mitochondria are the descendants of oxygen-respiring bacteria and chloroplasts were originally photosynthetic bacteria, are almost universally accepted



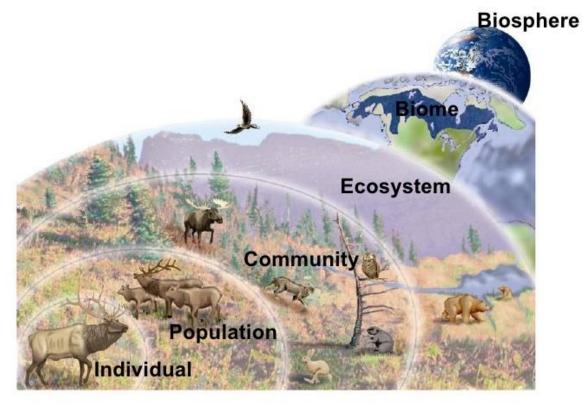
Biology is a vast discipline



The Hierarchical Structure of Life



Levels of Organization



Individual = 1 Species

Population = Many of the same species

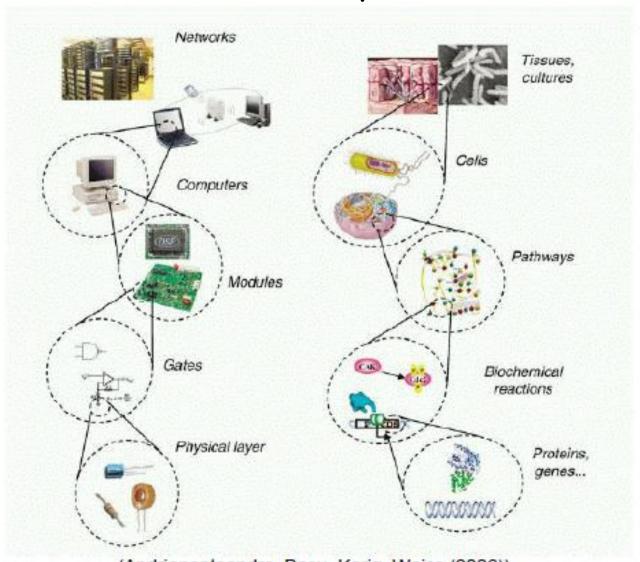
Community = Different populations (Biotic Factor = living)

Ecosystem = Various populations along with abiotic factors (non-living) coexisting.

Biome = Many ecosystems (Tundra, Tropical Rain Forest, Desert, etc...)

Biosphere = Many biomes

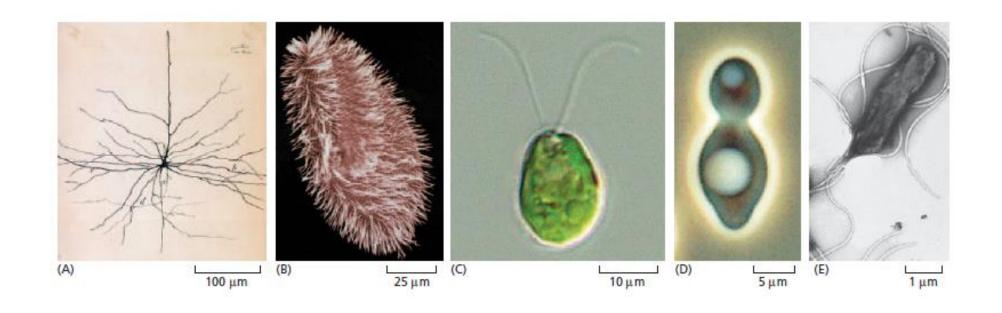
Cells compute?



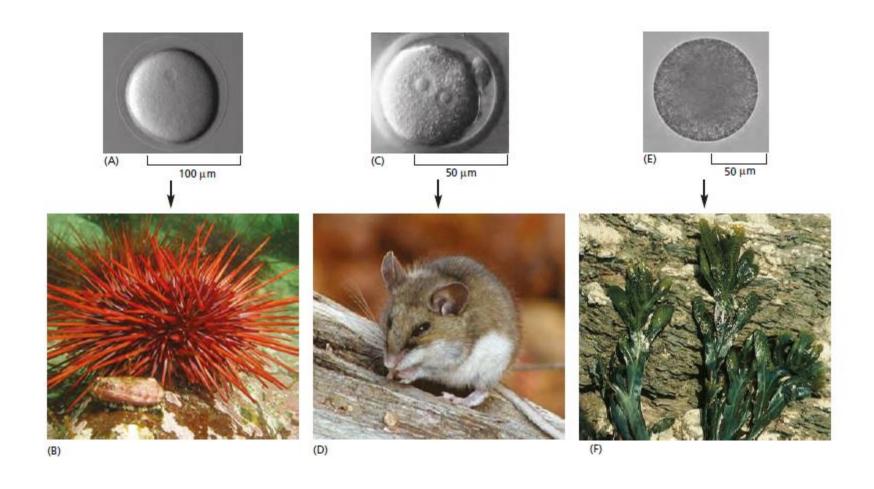
(Andrianantoandro, Basu, Karig, Weiss (2006))

Can we decipher the biological hardware and software?

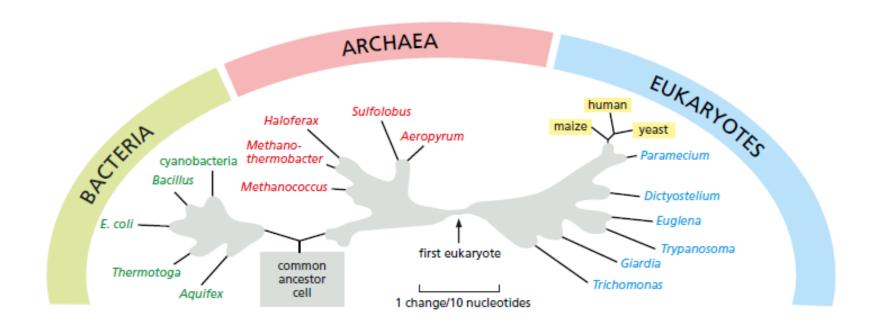
Cells come in variety of shapes and sizes



Origin from single cell

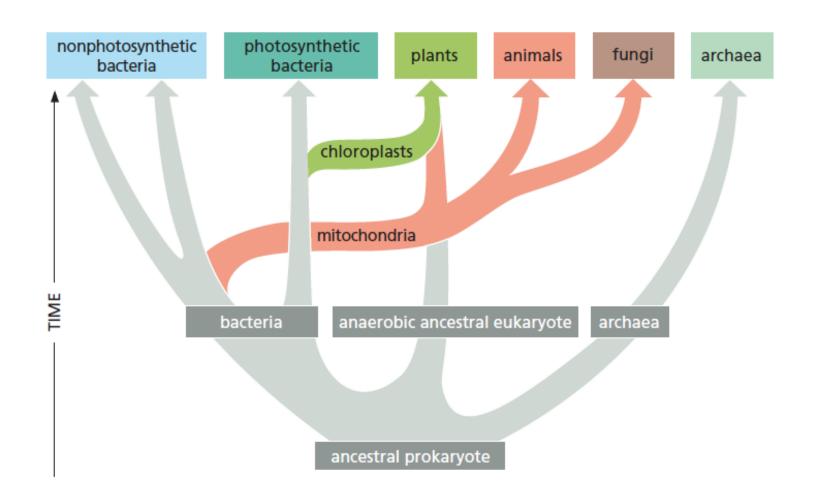


The Tree of Life

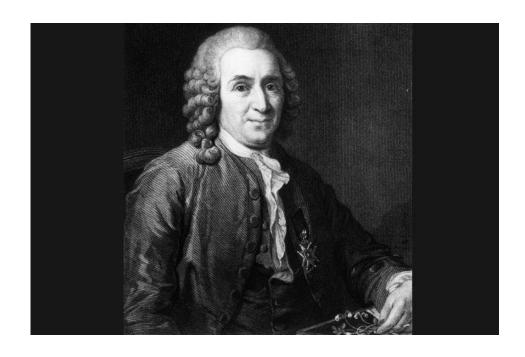


The domain system of classification was developed by Carl Woese and places organisms under three domains: **Archaea**, **Bacteria**, and **Eukarya**.

Under the domain system, organisms are further grouped into six Kingdoms. The Kingdoms include: **Archaebacteria** (ancient bacteria), **Eubacteria** (true bacteria), **Protista**, **Fungi**, **Plantae**, and **Animalia**.



Taxonomy and Organism Classification

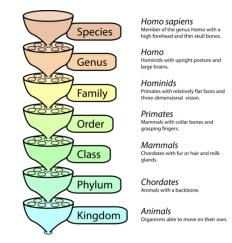


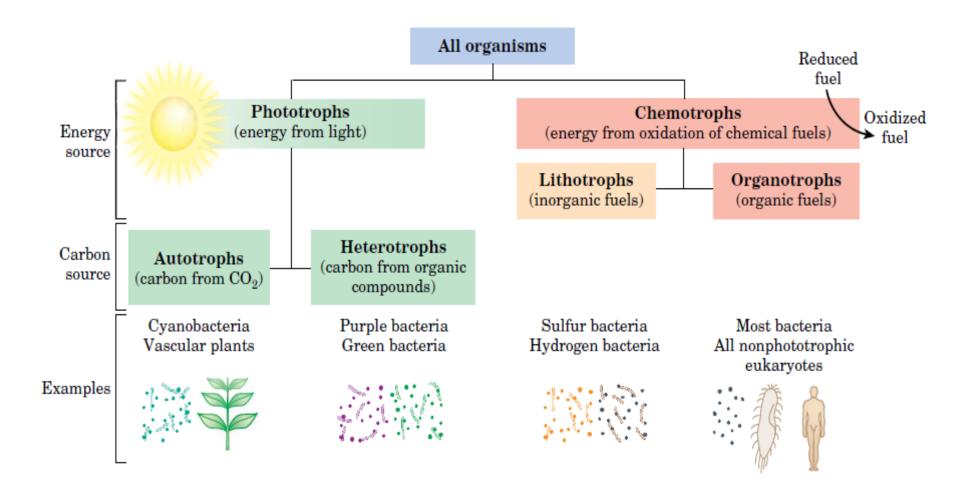
Botanist Carl von Linnaeus (1707-1778), founder of the modern system of binomial nomenclature for plants Kingdom, Phylum, Class, Order, Family, Genus, and Species.

A helpful aid for remembering the taxonomic categories:

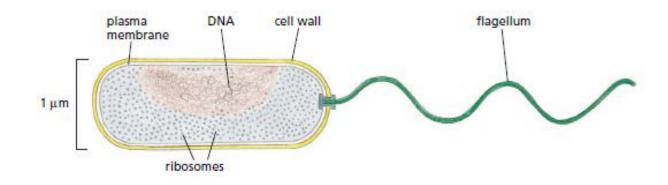
Domain, Kingdom, Phylum, Class, Order, Family, Genus, and Species is the mnemonic device:

Do Keep Plates Clean Or Family Gets Sick.

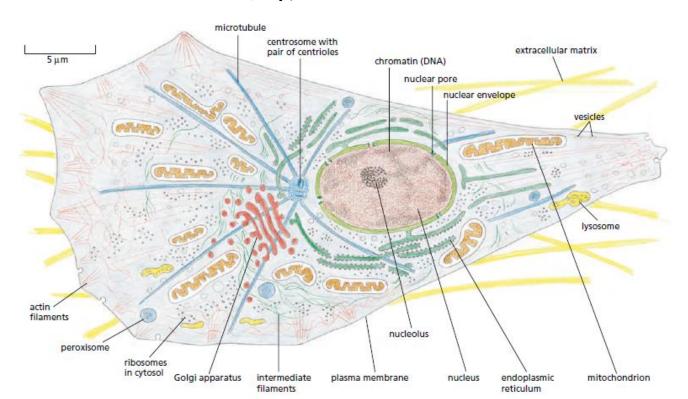




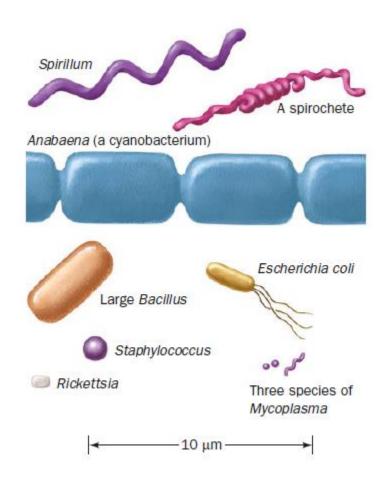
Bacterial cell



Animal cell



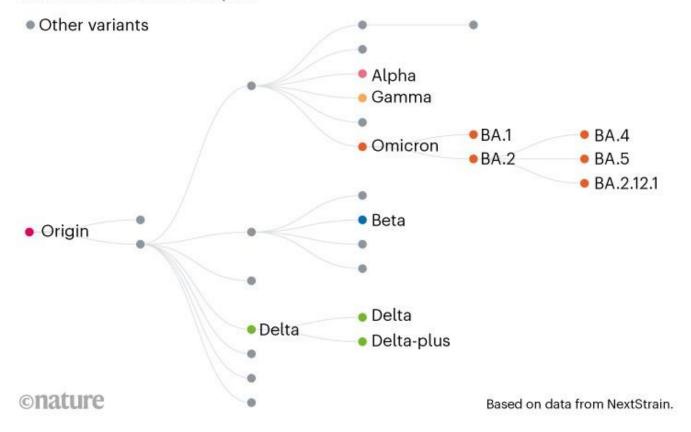
Scale drawings of some prokaryotic cells

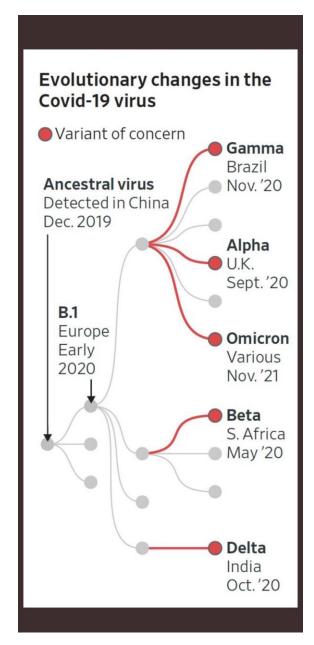


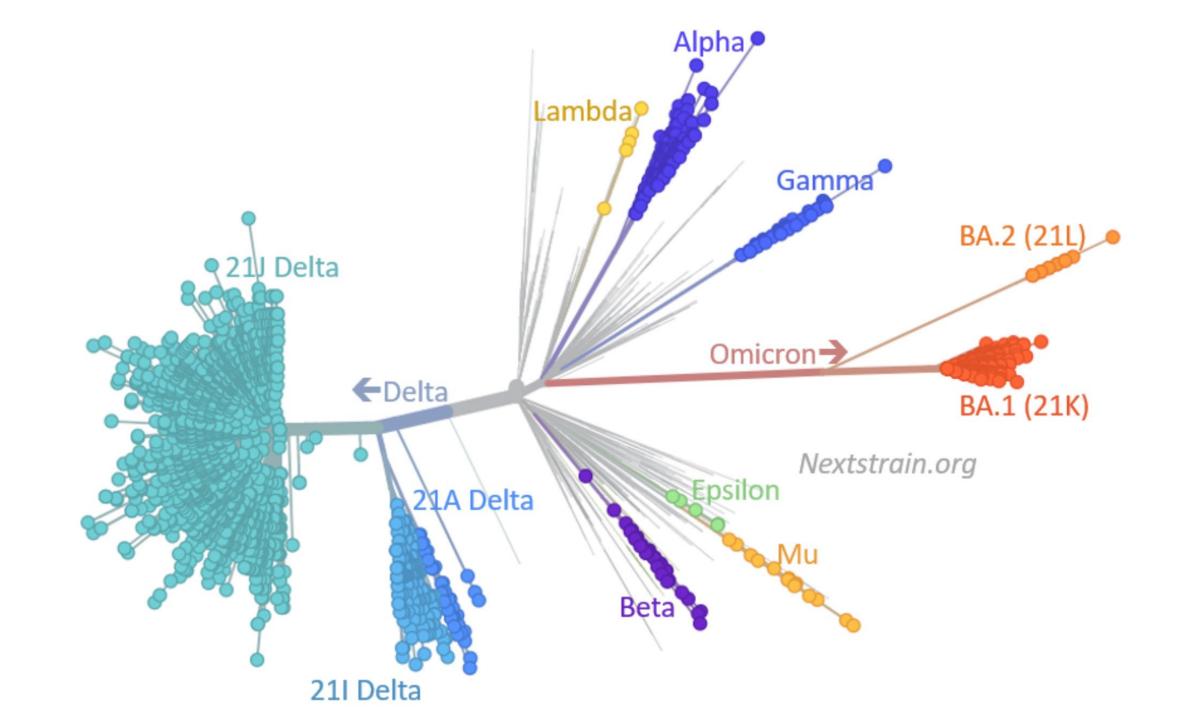
Three basic structures: spheroidal (cocci), rodlike (bacilli), and helically coiled (spirilla)

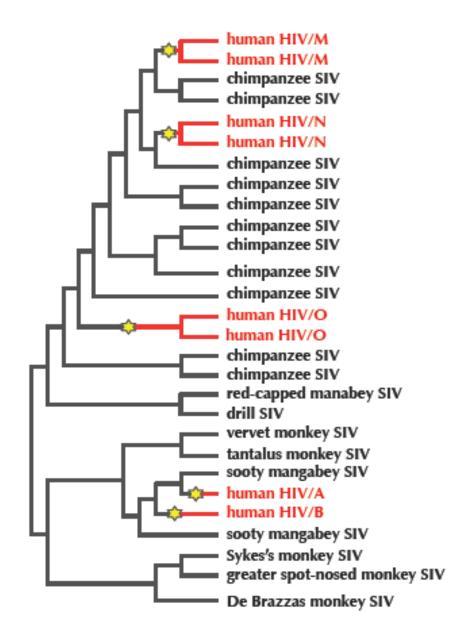
PATHOGEN PROGRESSION

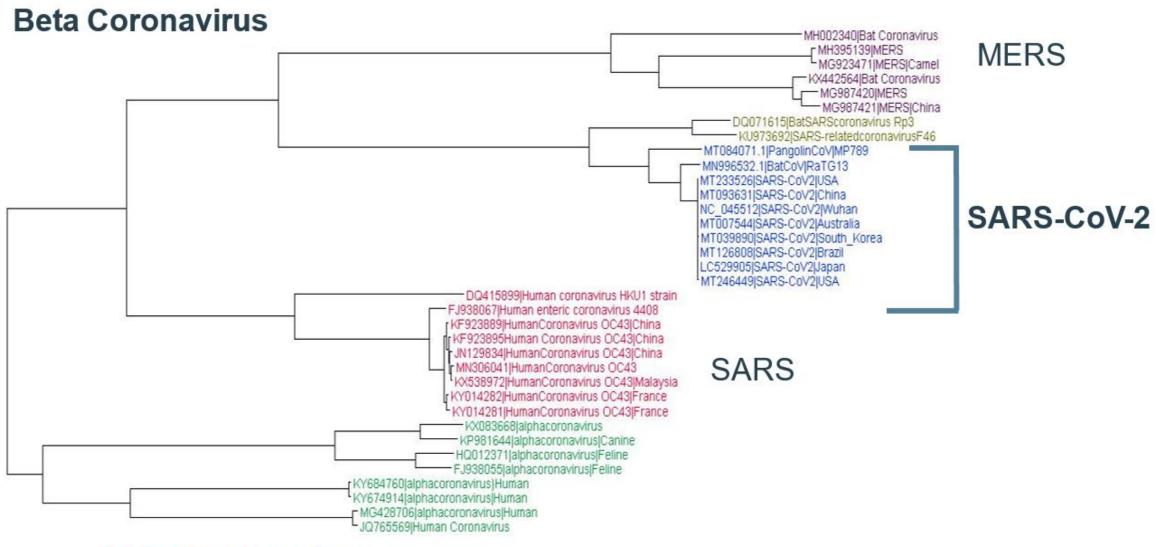
This diagram shows how the coronavirus SARS-CoV-2 has evolved to spawn several related variants. The latest are BA.4 and BA.5 along the Omicron lineage, which has dominated infections this year.











Alpha Coronavirus

Let us play the sequence alignment game!

Strings ATGCATGC and TGCATGCA!

ATGCATGC

TGCATGCA

ATGCATGC-

-TGCATGCA

We postulate a notion of a good alignment

as one that matches

as many symbols as possible.

Strings ATGCTTA and TGCATTAA

ATGC-TTA--TGCATTAA

Longest Common Subsequence Problem:

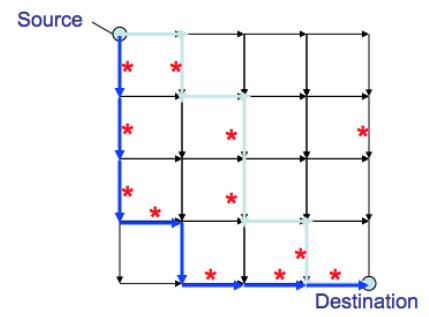
Find a longest common subsequence of two strings.

Input: Two strings.

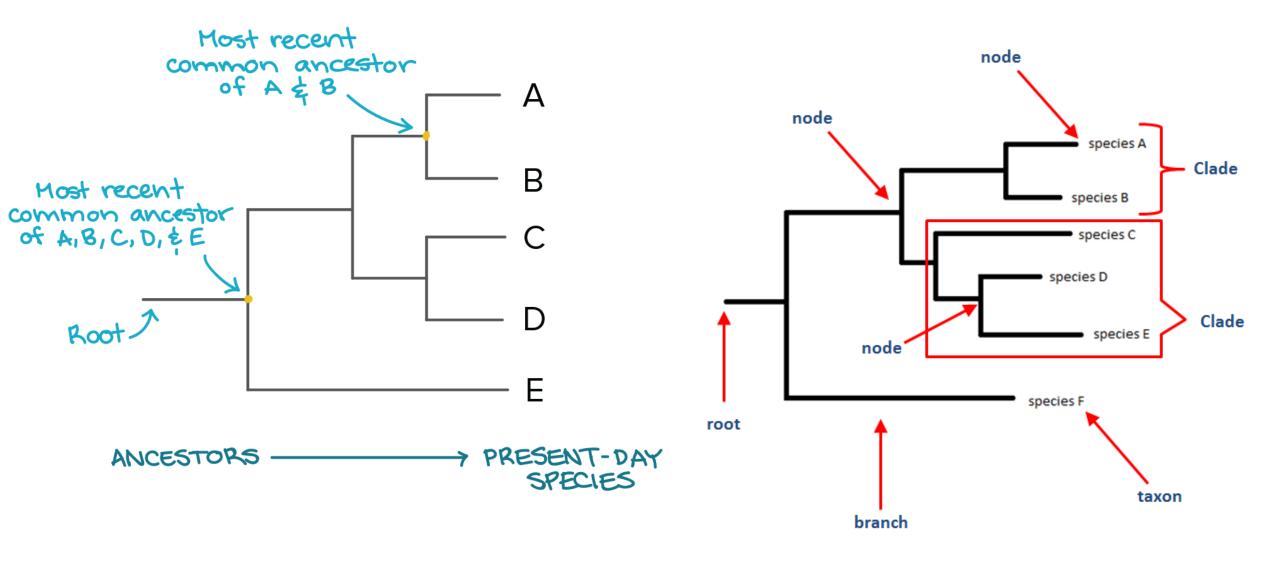
Output: A longest common subsequence of these strings.

The Manhattan tourist problem!



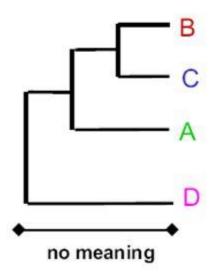


Parts of a phylogenetic tree



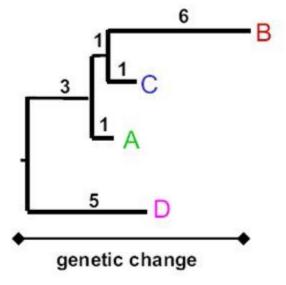
Types of trees

Cladogram



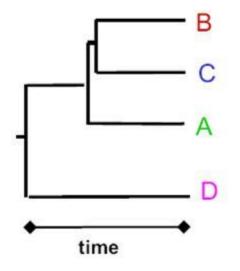
Simply shows relative recency of common ancestor

Phylogram

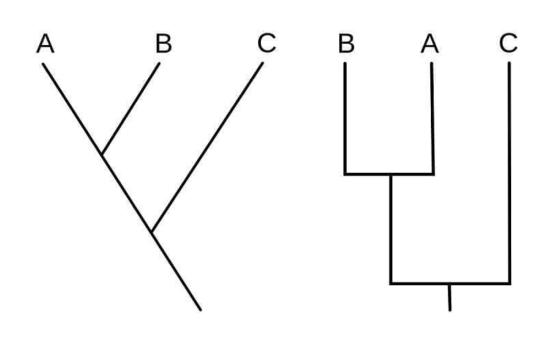


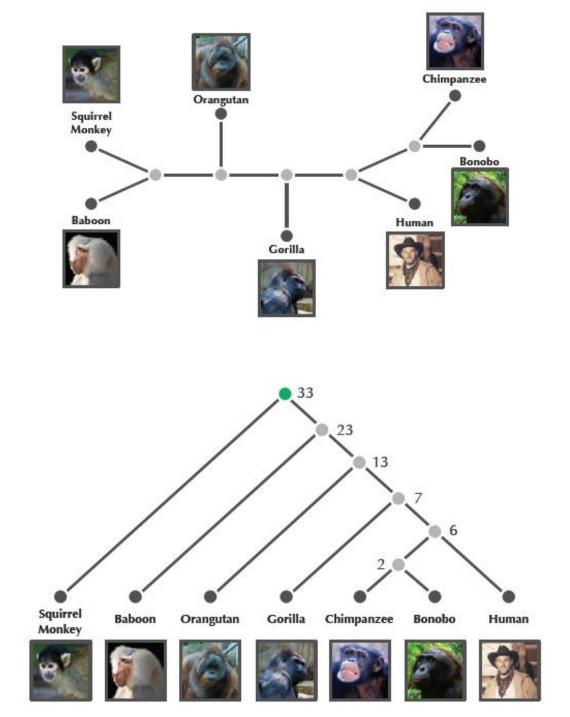
A cladogram with branch lengths

Ultrametric tree



A dendogram having all tips equidistant from root

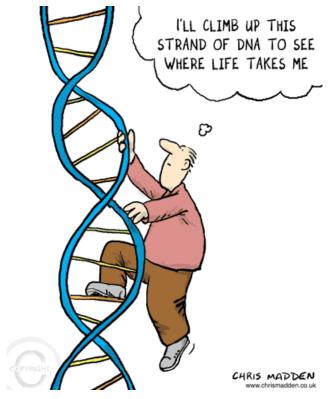




Some definitions

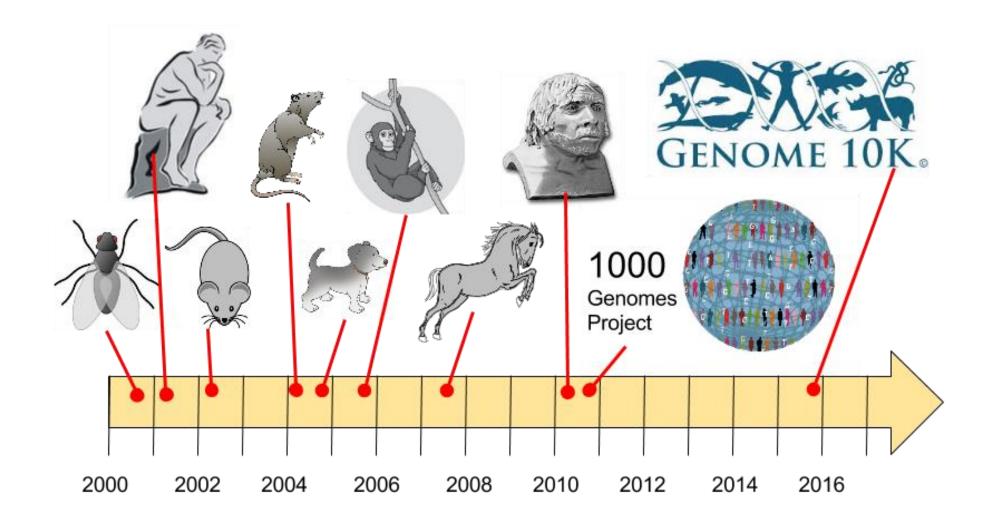
- Classification Grouping of things into classes.
 - e.g. Kingdom, phylum, class, order, family, genus, species.
- Taxonomy Giving names to things.
 - e.g. Taxons Mammalia, Homo sapiens, Felis
- Systematics- Understanding the relationships between things.
 - e.g. The daughter of my aunt is my cousin. The sister of my grandmother is my great-aunt.
- Phylogeny The evolutionary history of a species.
 - If you don't know your history, you don't know anything!
- Cladistics Phylogenetic systematics. Using the evolutionary history of species to understand their relationships, classify them, and give them names!

Universal feature-Chemical code



Organism	Genome size* (nucleotide pairs)	Approximate number of genes
Escherichia coli (bacterium)	4.6 × 10 ⁶	4300
Saccharomyces cerevisiae (yeast)	13 × 10 ⁶	6600
Caenorhabditis elegans (roundworm)	130 × 10 ⁶	21,000
Arabidopsis thaliana (plant)	220 × 10 ⁶	29,000
Drosophila melanogaster (fruit fly)	200 × 10 ⁶	15,000
Danio rerio (zebrafish)	1400 × 10 ⁶	32,000
Mus musculus (mouse)	2800 × 10 ⁶	30,000
Homo sapiens (human)	3200 × 10 ⁶	30,000

^{*}Genome size includes an estimate for the amount of highly repeated DNA sequence not in genome databases.



TGCCAAGCAGCAAAGTTTTGCTGCTGTTTATTTTTTGTAGCTCTTACTATATTCTACTTTTAC TAATATTACTATTACACATAATTATTTTTTATATATATGAAGTACCAATGACTTCCTTTTCCAG AGCAATAATGAAATTTCACAGTATGAAAATGGAAGAAATCAATAAAATTATACGTGACCT GTGGCGAAGTACCTATCGTGGACAAGGTGAGTACCATGGTGTATCACAAATGCTCTTTCC AAAGCCCTCTCCGCAGCTCTTCCCCTTATGACCTCTCATCATGCCAGCATTACCTCCCTGG ACCCCTTTCTAAGCATGTCTTTGAGATTTTCTAAGAATTCTTATCTTGGCAACATCTTGTAG CAAGAAATGTAAAGTTTTCTGTTCCAGAGCCTAACAGGACTTACATATTTGACTGCAGT AGGCATTATATTTAGCTGATGACATAATAGGTTCTGTCATAGTGTAGATAGGGATAAGCCA TTCCAGATGGAGTCTCGCACTTCTCTGTCACCCGGGCTGGAGCGCAGTGGTGCAATCTT GGCTCACTGCAACCTCCACCTCCTGGGTTCAGGTGATTCTCCCACCTCAGCCTCCCGAGT AGTAGCTGGAATTACAGGTGCGCGCTCCCACACCTGGCTAATTTTTTTGTATTCTTAGTAG AGATGGGGTTTCACCATGTTGGCCAGGCTGGTCTCAAACTCCTGCCCTCAGGTGATCTG CCCACCTTGGCCTCCCAGTGTTGGGTTTACAGGCGTGAGCCACCGCGCCTGGCCTGGA GGAAACTCTTAACAGGGAAACTAAGAAAGAGTTGAGGCTGAGGAACTGGGGCATCTG ACACCAGCCATTGTTTTCCTCTGGTAATGTCAGCCTCATCTGTTGTTCCTAGGCTTACTTG ATATGTTTGTAAATGACAAAAGGCTACAGAGCATAGGTTCCTCTAAAATATTCTTCCT GTGTCAGATATTGAATACATAGAAATACGGTCTGATGCCGATGAAAATGTATCAGCTTCTG ATAAAAGGCGGAATTATAACTACCGAGTGGTGATGCTGAAGGGAGACACAGCCTTGGA TATGCGAGGACGATGCAGTGCTGGACAAAAGGCAGGTATCTCAAAAGCCTGGGGAGCC AACTCACCCAAGTAACTGAAAGAGAGAAACAAACATCAGTGCAGTGGAAGCACCCAAG GCTACACCTGAATGGTGGGAAGCTCTTTGCTGCTATATAAAATGAATCAGGCTCAGCTAC TATTATT

The Human Genome



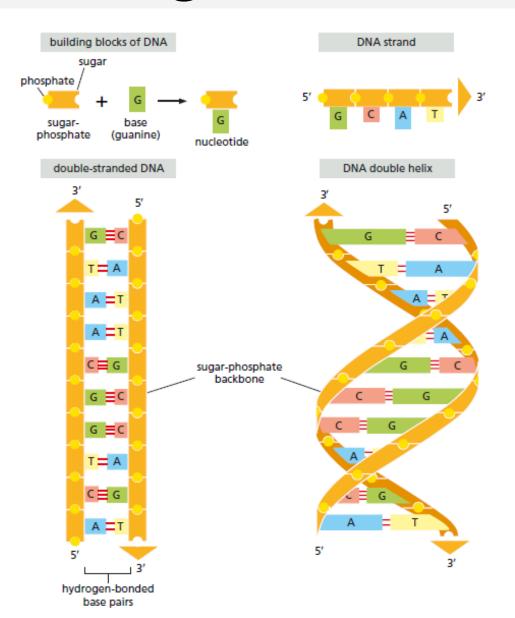
2003

"Fifteen years, six countries, twenty centers. Three billion dollars, three billion letters. One dollar per letter—such a deal!"

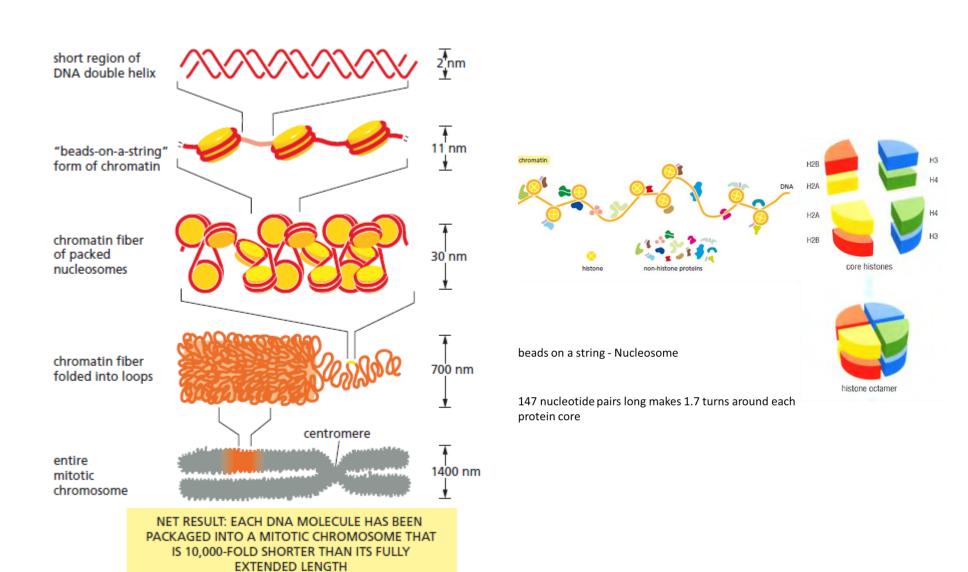
Eric Lander's seven-word "Nano Lecture"

"Genome: bought the book; hard to read"

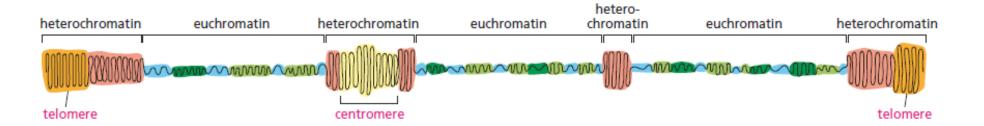
DNA and its building blocks



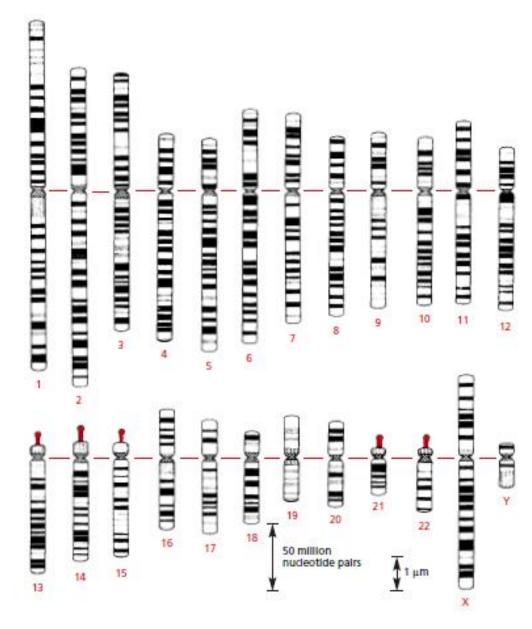
The organization of Chromosomes



The structure of chromatin varies



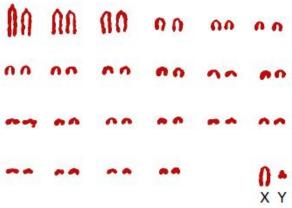
The banding patterns of human chromosomes



Giemsa Staining

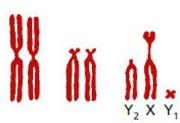


Chinese muntjac

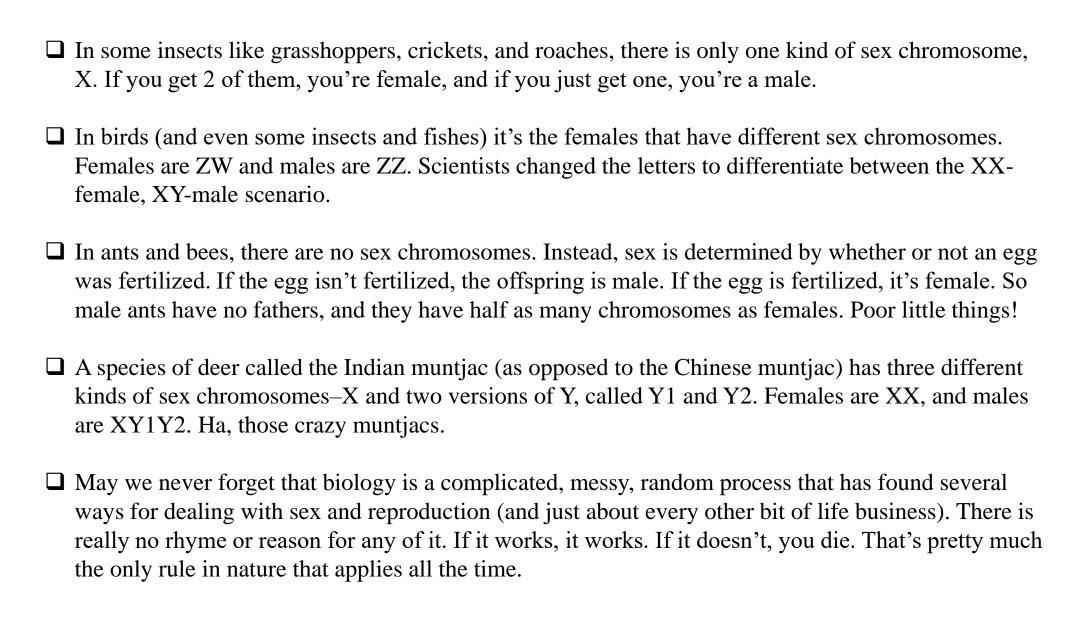




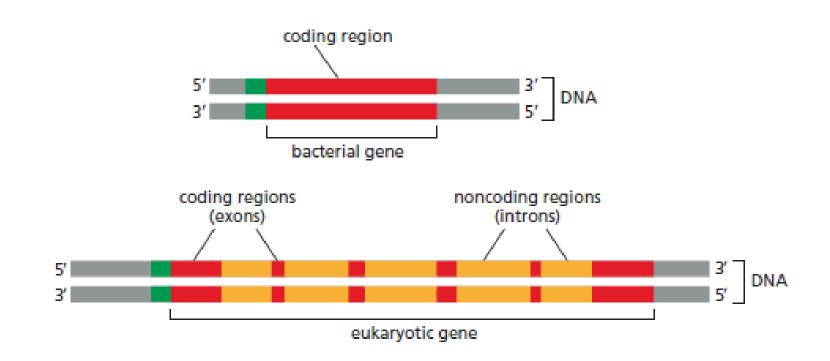
Indian muntjac



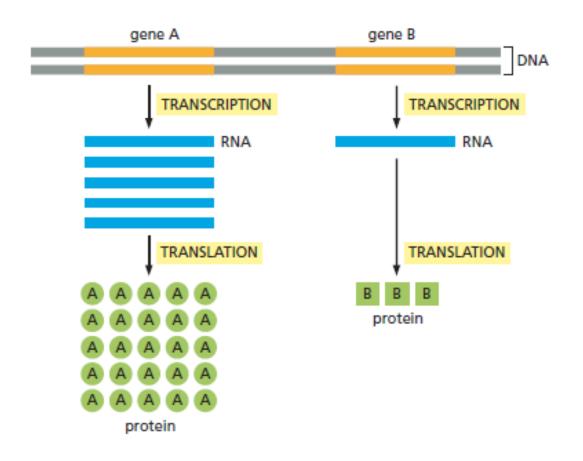
Interesting observations from Beatrice!



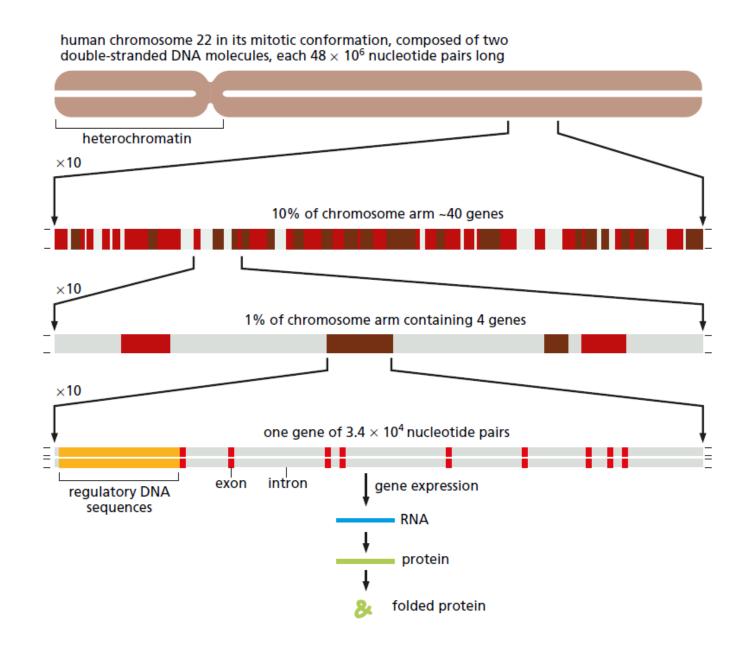
Eukaryotic and bacterial genes are organized differently



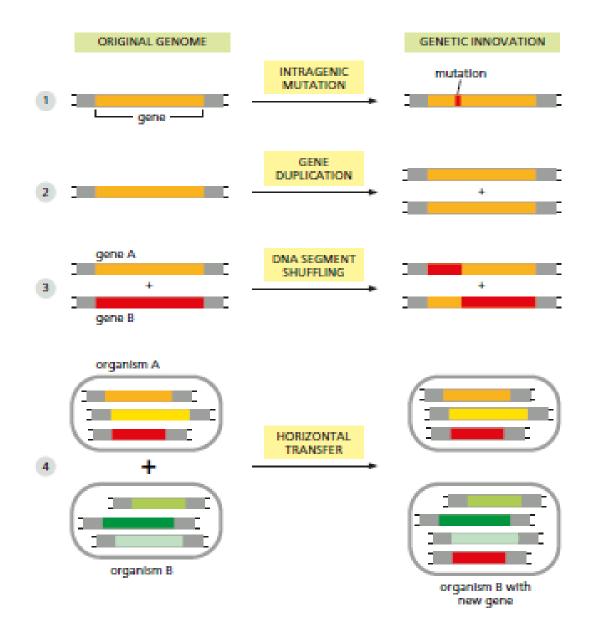
How Cells Read the Genome



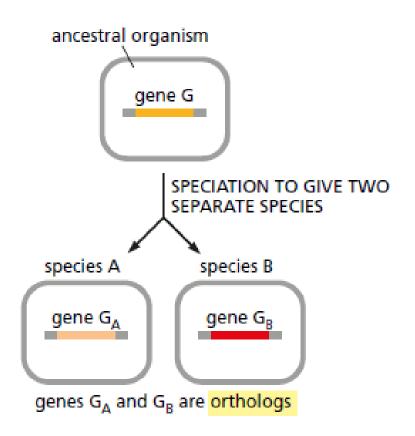
The organization of genes on a human chromosome

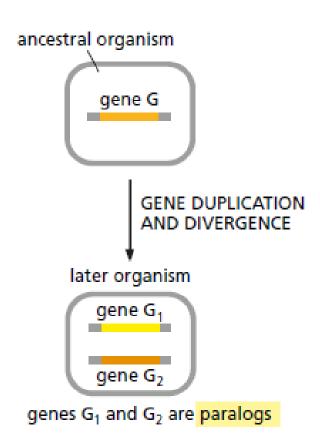


New genes are generated from existing genes



Gene duplications give rise to families of genes

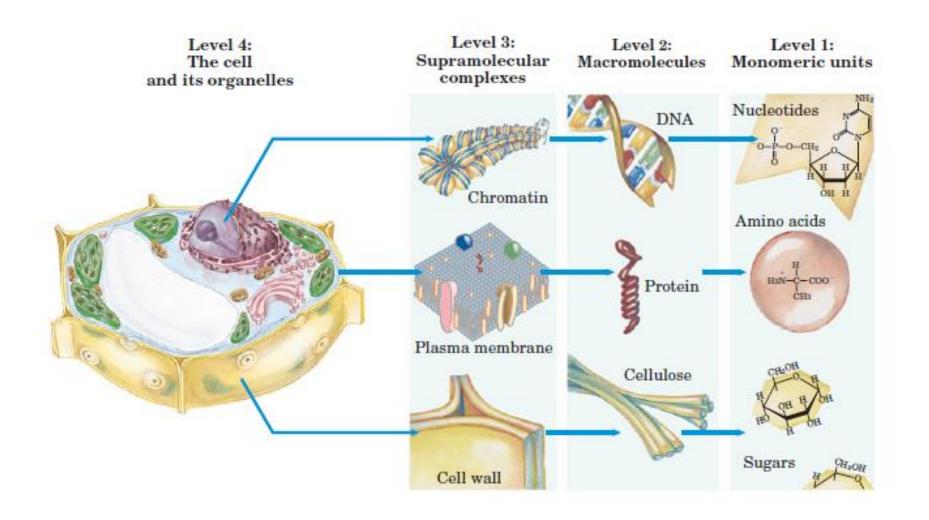




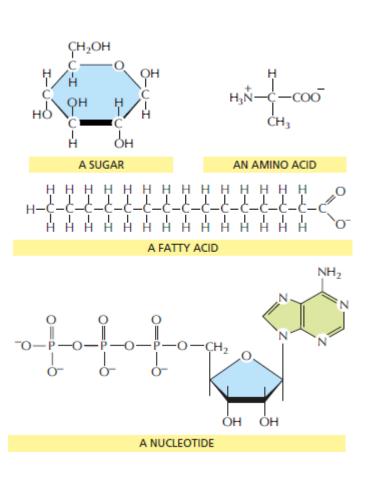


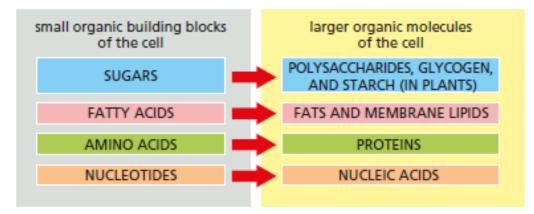


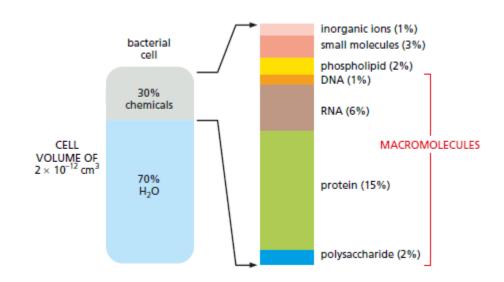
Structural hierarchy in the molecular organization of cells



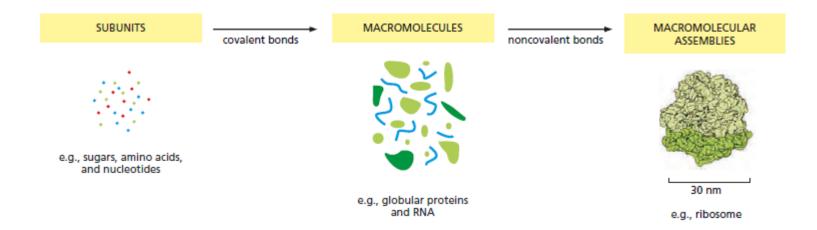
Cells function as biochemical factories

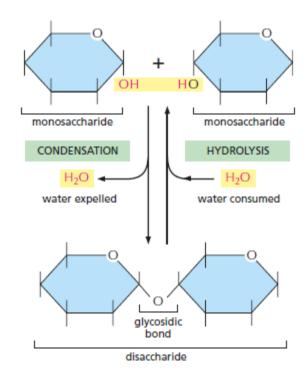




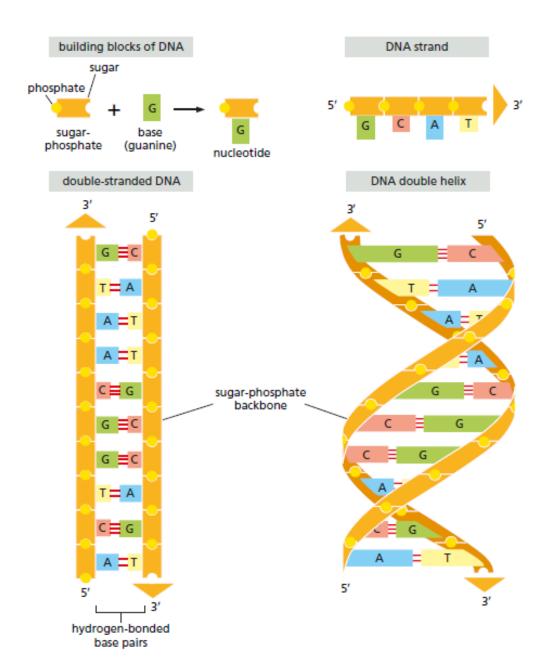


Biochemical bond formation



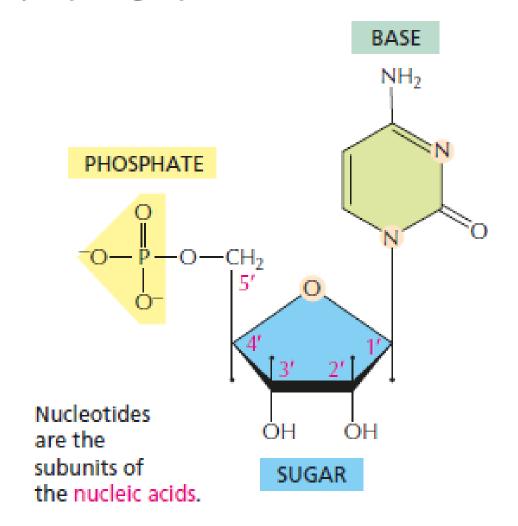


DNA and its building blocks

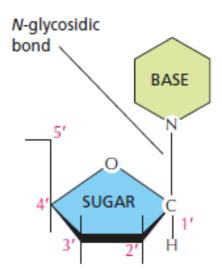


Nucleotides

A nucleotide consists of a nitrogen-containing base, a five-carbon sugar, and one or more phosphate groups.

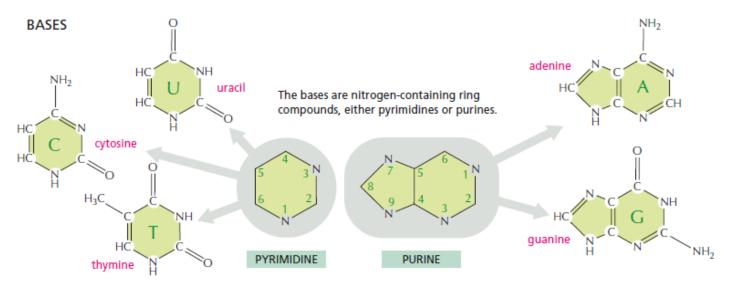


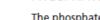
BASE-SUGAR LINKAGE



The base is linked to the same carbon (C1) used in sugar–sugar bonds.

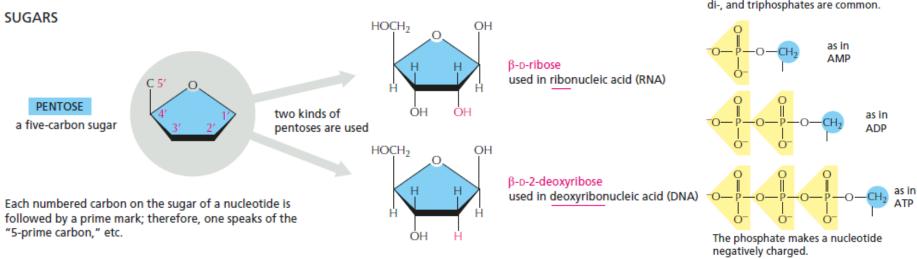
Nucleotides





PHOSPHATES

The phosphates are normally joined to the C5 hydroxyl of the ribose or deoxyribose sugar (designated 5'). Mono-, di-, and triphosphates are common.



Nucleotides vs Nucleoside

NOMENCLATURE

The names can be confusing, but the abbreviations are clear.

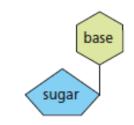
BASE	NUCLEOSIDE	ABBR.
adenine	adenosine	Α
guanine	guanosine	G
cytosine	cytidine	С
uracil	uridine	U
thymine	thymidine	Т

Nucleotides are abbreviated by three capital letters. Some examples follow:

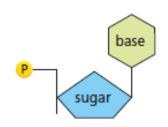
AMP = adenosine monophosphate

 $dAMP = deoxyadenosine\ monophosphate$

UDP = uridine diphosphate ATP = adenosine triphosphate

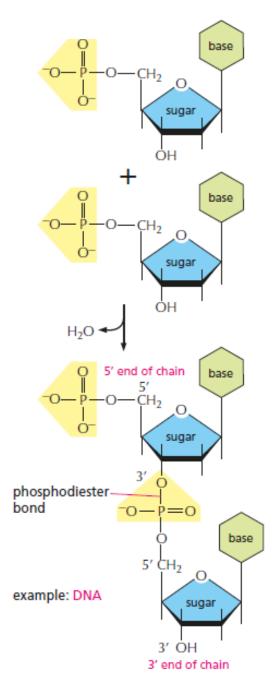


BASE + SUGAR = NUCLEOSIDE



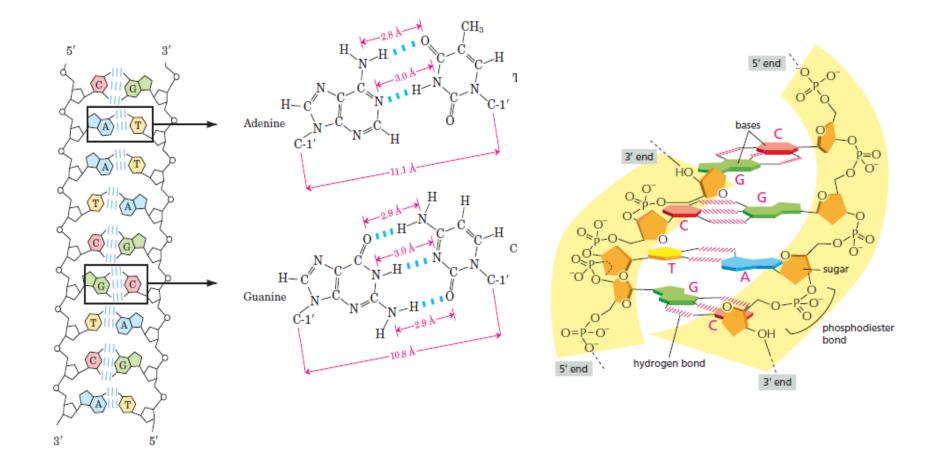
BASE + SUGAR + PHOSPHATE = NUCLEOTIDE

Nucleic Acids



Nucleotides are joined together by phosphodiester bonds between 5' and 3' carbon atoms of the sugar ring, via a phosphate group, to form nucleic acids. The linear sequence of nucleotides in a nucleic acid chain is commonly abbreviated by a one-letter code, such as AGCTTACA, with the 5' end of the chain at the left.

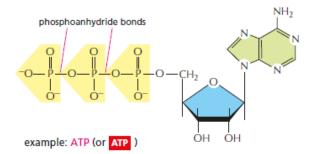
Hydrogen-Bonding Patterns

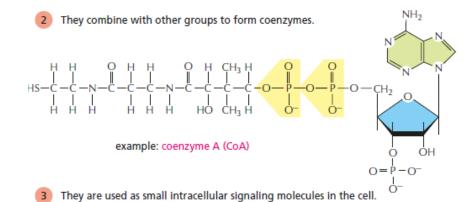


Nucleotides

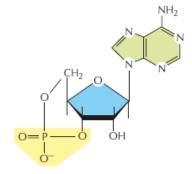
NUCLEOTIDES HAVE MANY OTHER FUNCTIONS

1 They carry chemical energy in their easily hydrolyzed phosphoanhydride bonds.

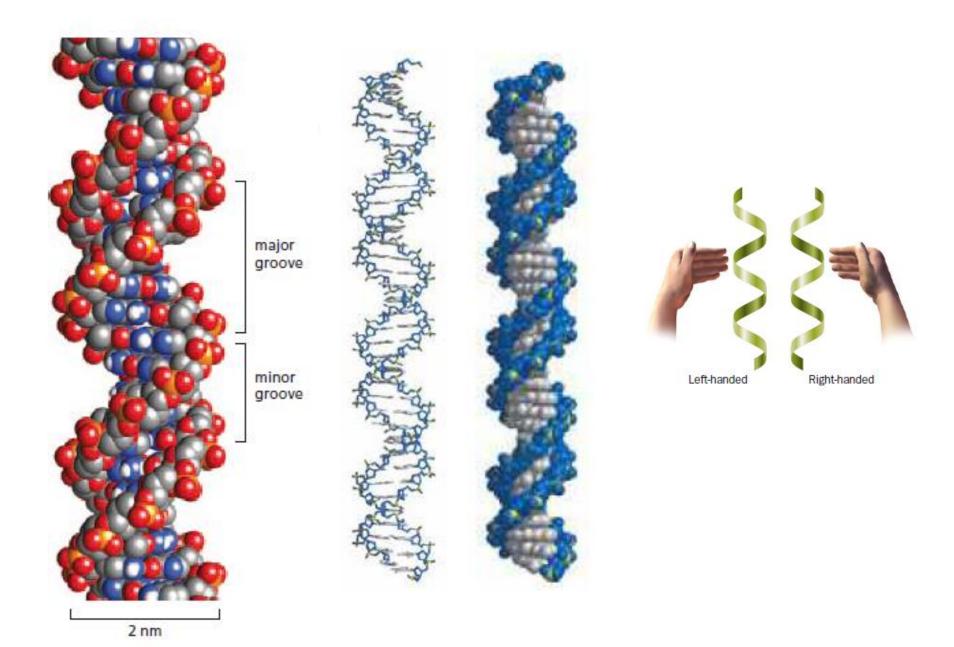




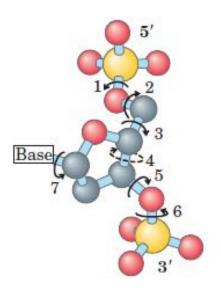
example: cyclic AMP

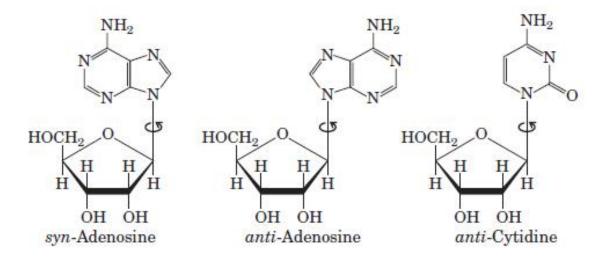


Structure of DNA

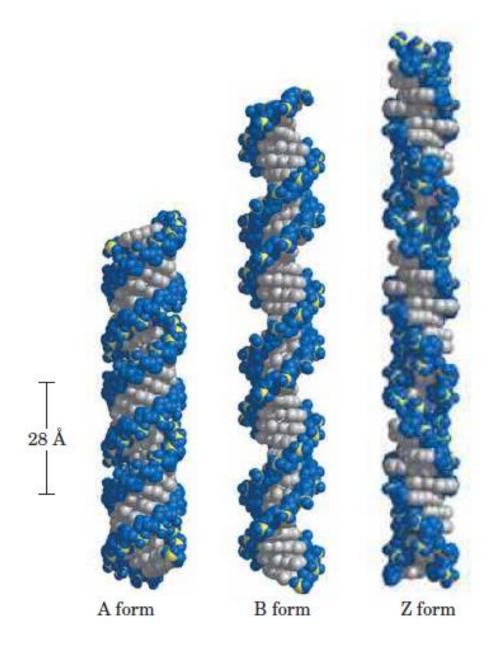


Structural variation in DNA



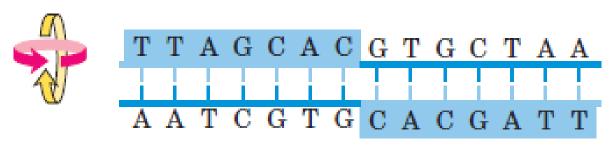


Structural variation in DNA

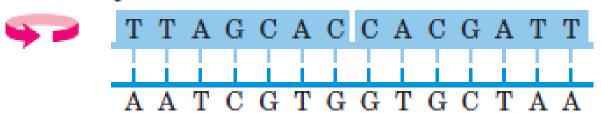


Certain DNA Sequences Adopt Unusual Structures

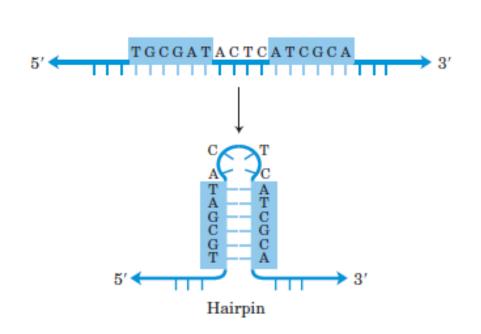
Palindrome

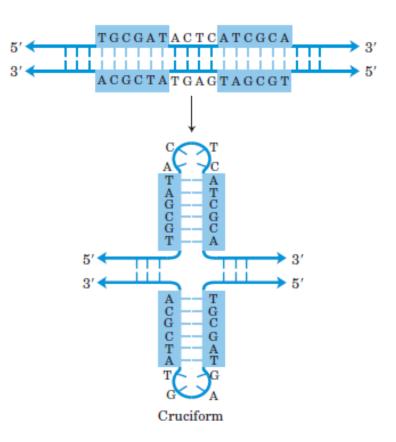


Mirror repeat



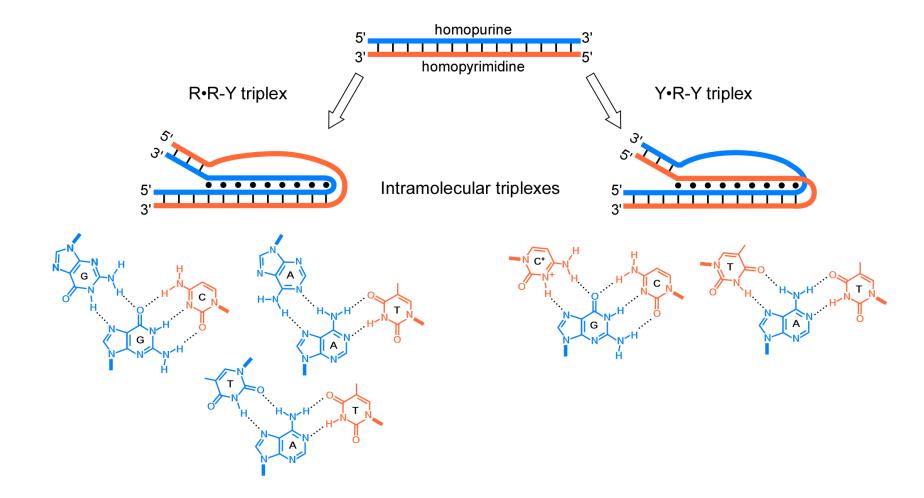
Hairpins and cruciforms

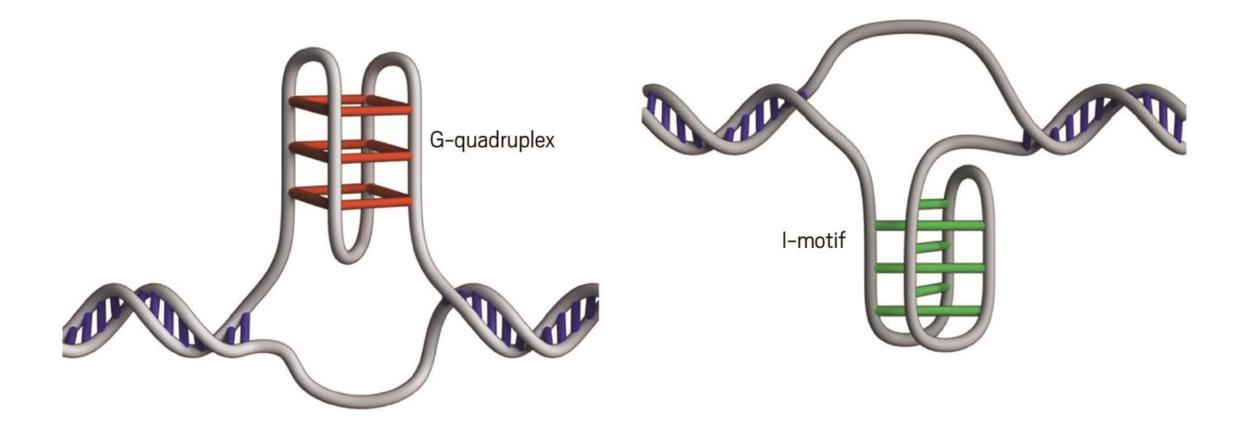




Several unusual DNA structures involve three or even four DNA strands

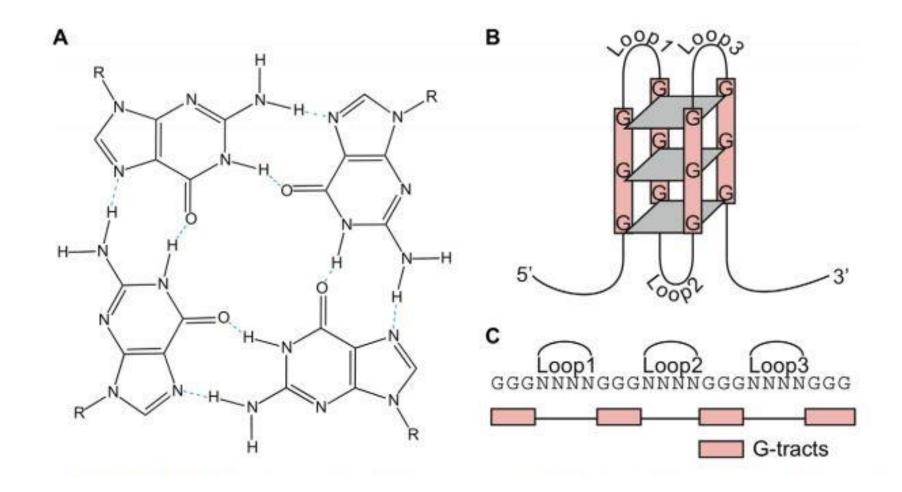
The atoms that participate in the hydrogen bonding of **triplex DNA**, are often referred to as **Hoogsteen positions**, and the non-Watson-Crick pairing is called **Hoogsteen pairing**, after Karst Hoogsteen, who in 1963 first recognized the potential for these unusual pairings



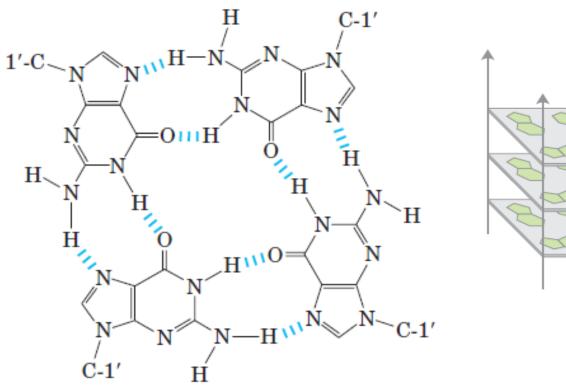


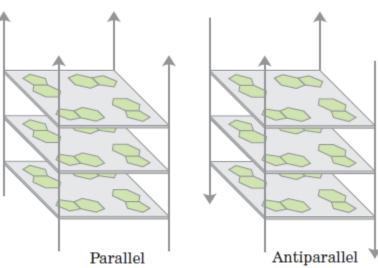
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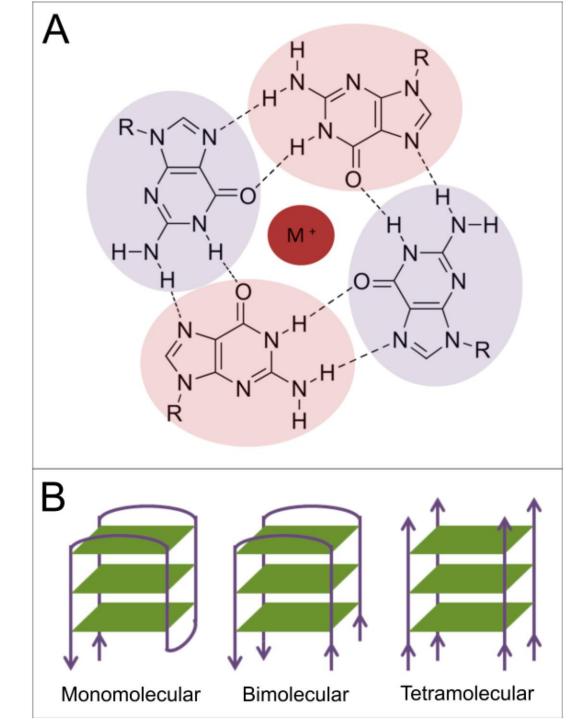
G-quadruplexes

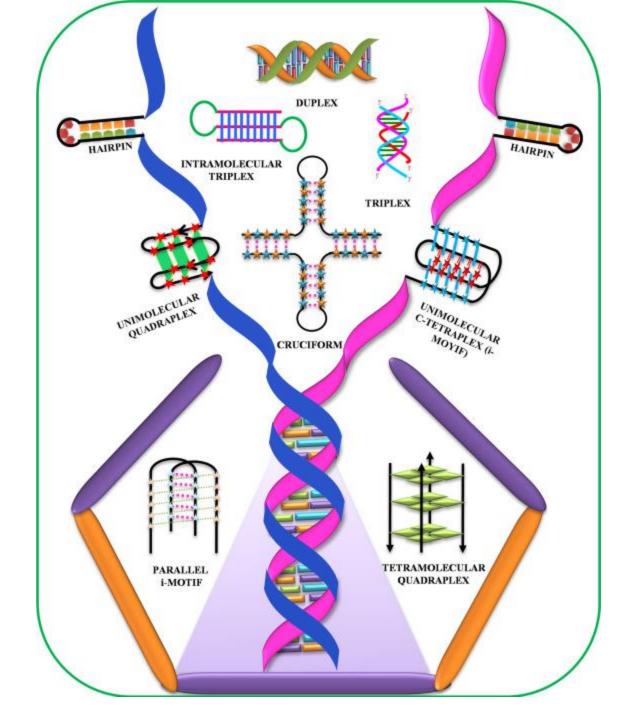


G-quadruplexes







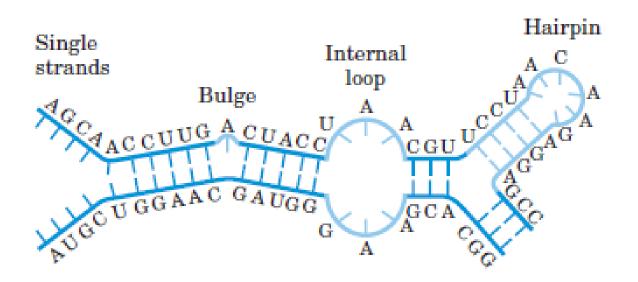


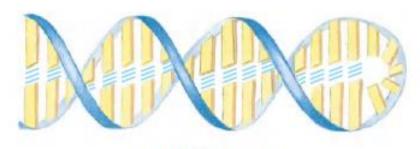
Intramolecular

Vs

Intermolecular

Structure of RNA





Hairpin double helix

The paired regions generally have an A-form right-handed helix, as shown for a hairpin