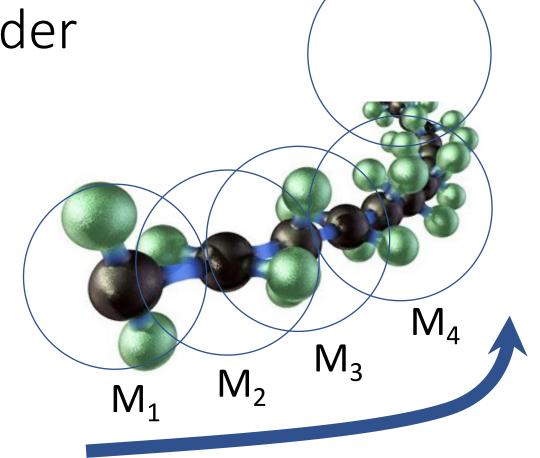
Prof. Sabrina Pricl A.Y. 2021-2022

Lesson 3 Nucleic acid polarity and structure



Macromolecules' Law & Order

- Macromolecules carry information as:
 - They have two ends (starting and terminal end)
 - They have a direction (from start to end)

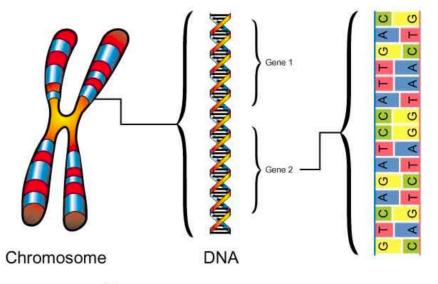


Macromolecules' Law & Order

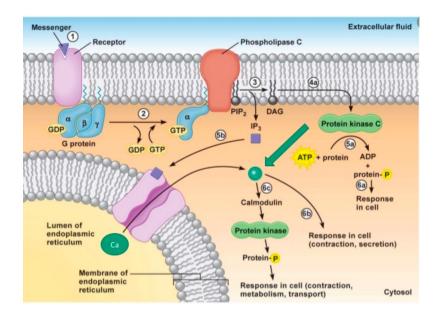
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- Macromolecules possess ORDER and POLARITY
 - These two features constitute the FUNDAMEMTAL INFORMATION cellules can read and understand

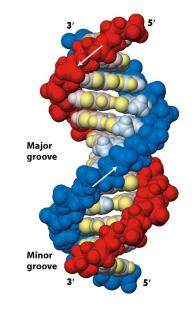
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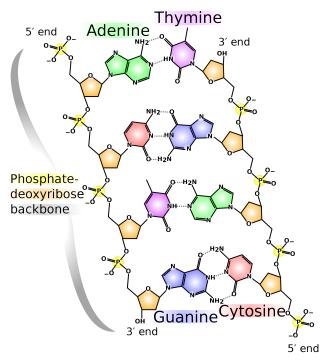
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- NUCLEIC ACIDS = carries of hereditary information
- PROTEINS = all other info/instructions



Genes



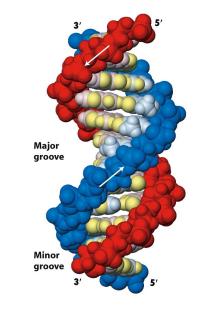


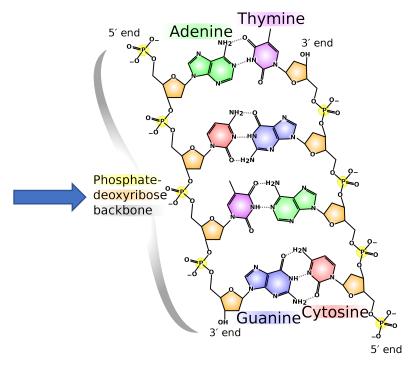


• Have two hands: 5' e 3'

5'P-S-P-S-P-S3'
B1 B2 B3

 P-S = sugar-phosphate backbone joined by phosphodiester bonds



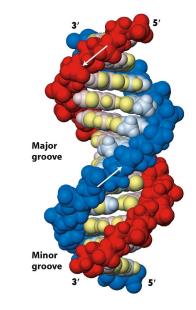


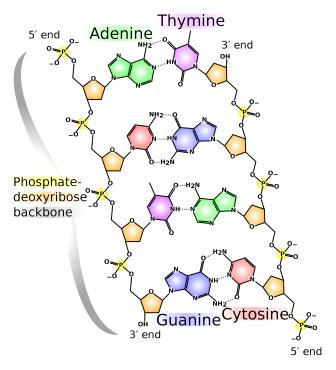
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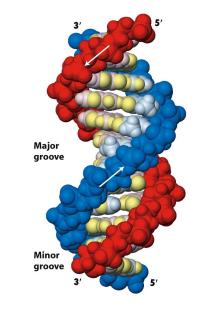
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- 5' and 3' ends are **CHEMICALLY DIFFERENT ENDS** and cells can distinguish them from one another

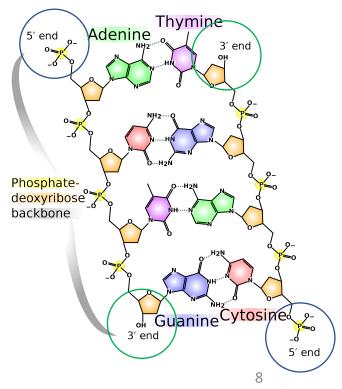




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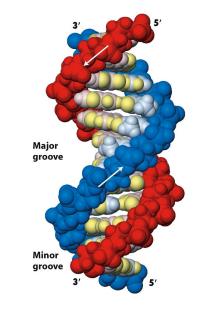
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 - 5'end = P-5'C on sugar
 - 3'end = 3'OH on sugar

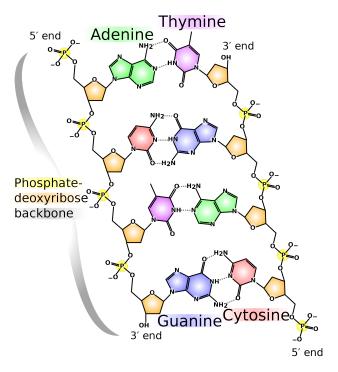




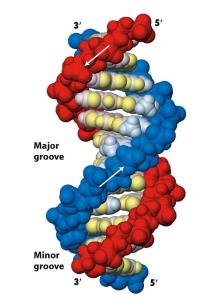
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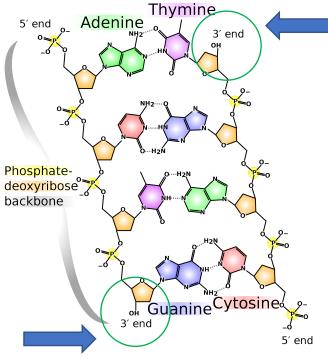
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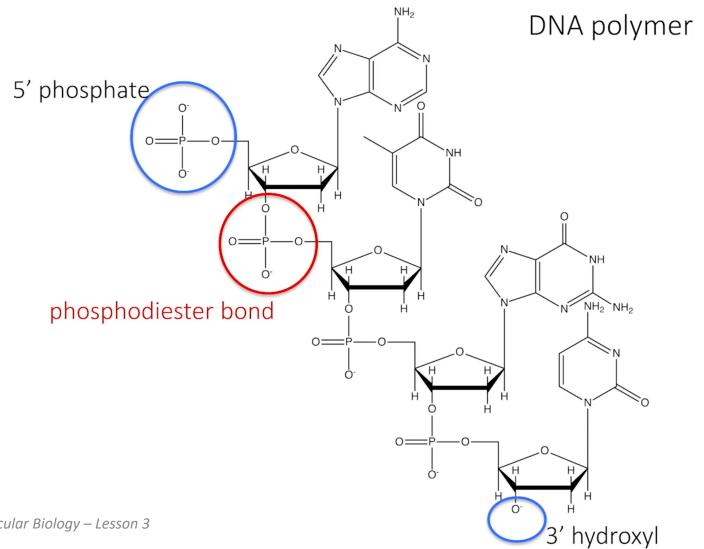


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 - 5'end = P-5'C on sugar
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- Base order along the polymer
- 3' is the last base added (any new incoming base will be added to 3'OH)





Nucleic acids



 S-P backbone is not written, just the bases + polarity

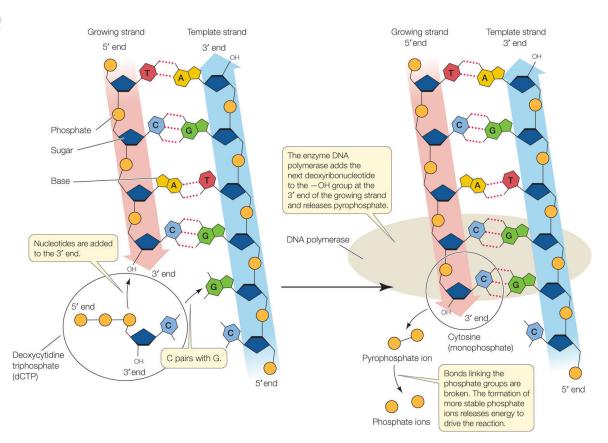
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- ALWAYS write 5' and 3' on each nucleic acid strand

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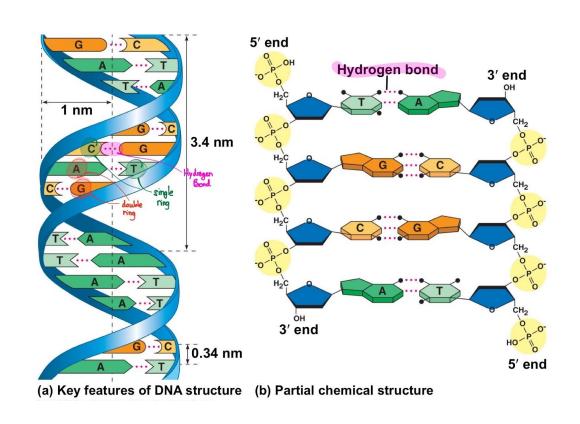
 $5'B_1B_2B_3B_4B_5B_63' = 5'GAATCC3'$

- Base order = INFORMATION
- Polarity = 5' and 3' ends: shows
 - First to last nucleotide added
 - Direction to read information

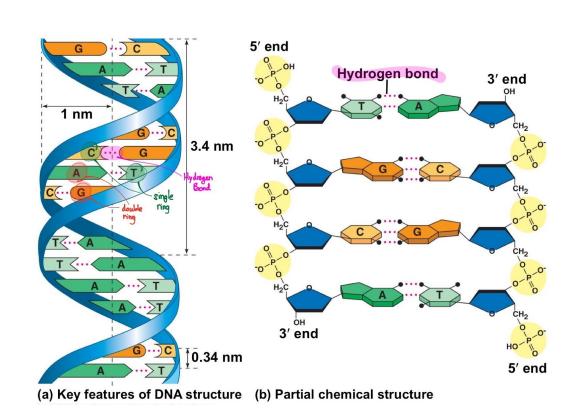


Get to know your DNA structure

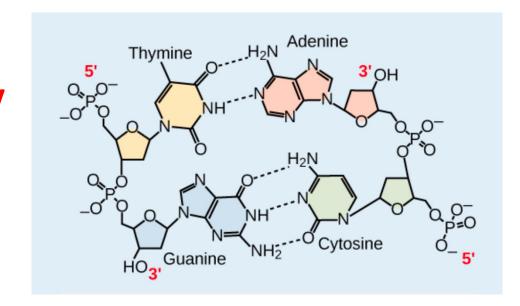
- DNA has a right-handed double-helix (DH) structure (2 nm diameter)
 - Ss and Ps lie on the outside of the helix (backbone)
 - Bs are stacked in the interior, in pairs
 - B pairs (BPs) are bound to each other by Hbonds
 - Every BP in the DH is separated from the next base pair by 0.34 nm (axial rise)
 - There are 10.4 BPs per helical trun
 - The distance to complete one helical turn (helical pitch) = 3.4 nm



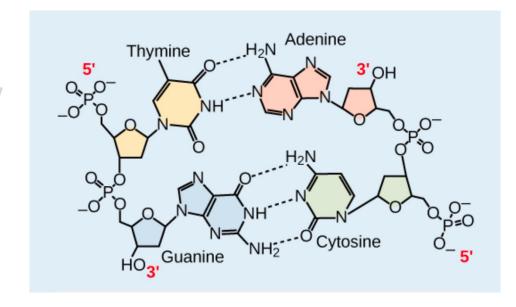
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 - There are 10.4 BPs per helical trun
 - The distance to complete one helical turn (helical pitch) = 3.4 nm
- The two strands of the helix run in opposite directions
 - the 5' carbon end of one strand will face the 3' carbon end of its matching strand
- This is referred to as antiparallel orientation
 - **Key DNA PROPERTY** (for DNA replication and in many nucleic acid interactions)



- Only certain types of base pairing are allowed:
- Specifically: A can only pair with T and G can only pair with C (base complementary rule)

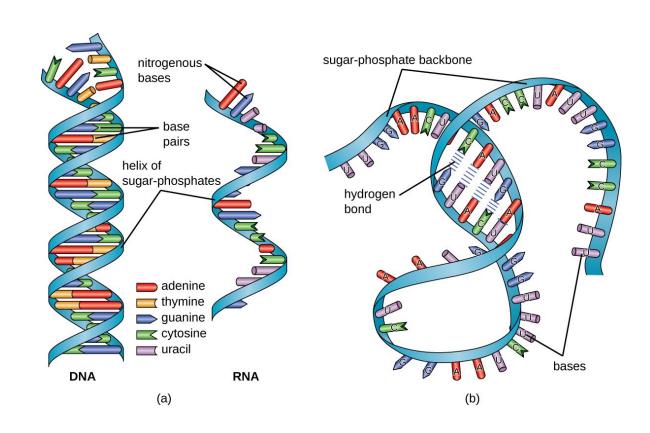


- Only certain types of base pairing are allowed:
- Specifically: A can only pair with T and G can only pair with C (base complementary rule)
- In other words, the DNA strands are complementary to each other
 - If the sequence of one DNA strand is **5'AATTGGCC3'**, the complementary strand would have the sequence **3'TTAACCGG5'**

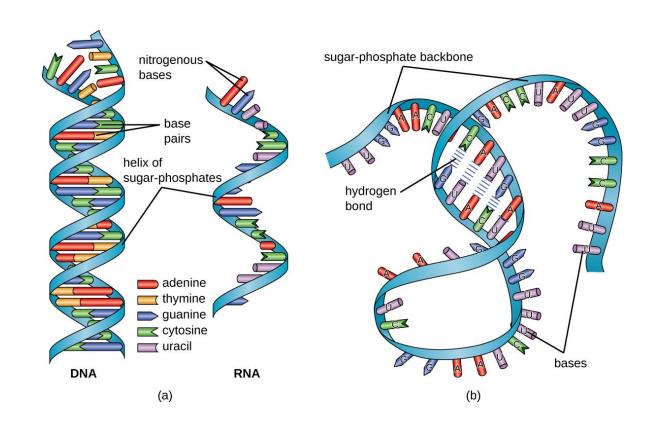


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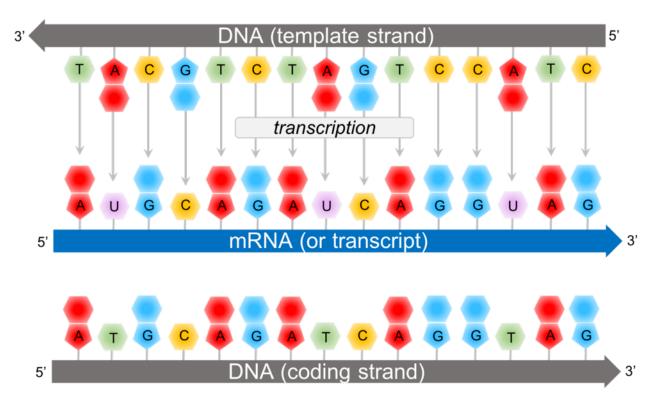
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- RNA is usually single-stranded
- The RNA four nitrogenous bases are A, U, G, and C
- There are four major types of RNA
 - messenger RNA (mRNA), ribosomal RNA (rRNA), transfer RNA (tRNA), and microRNA (miRNA) (more later)



- The mRNA carries the message from DNA
 - If a cell requires a certain protein to be synthesized, the gene for this product is "turned on" and the messenger RNA is synthesized in the cellular nucleus (more later in the course)

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- The RNA base sequence is complementary to the coding sequence of the DNA which it has been copied from but
- If the DNA strand to be copied (template strand) has a sequence 3'TACGTCTA...3', the sequence of the complementary mRNA strand (transcript) is 5'AUGCAGAU...3'



Note: template strand reading is $3' \rightarrow 5'$ mRNA synthesis is $5' \rightarrow 3'$ (more later)

Nucleic acid polarity and structure

• Take assignment 3: Nucleic acid polarity and structure