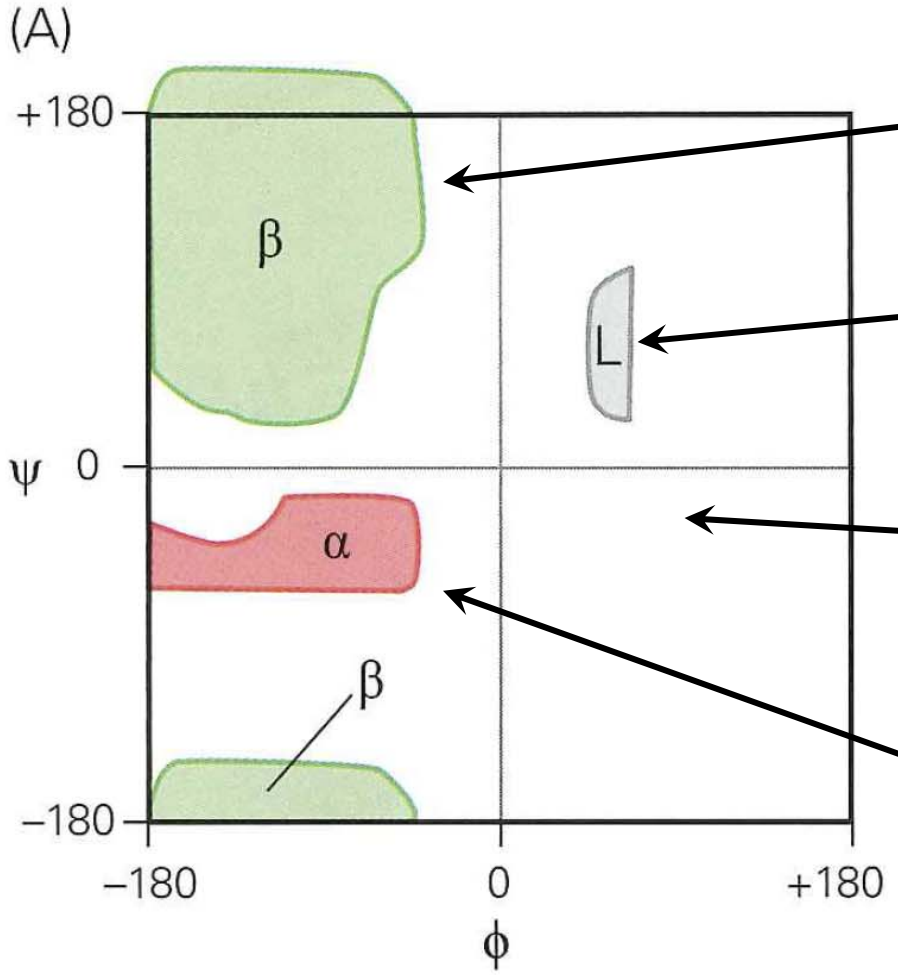
The Ramachandran Plot:

Sterically-allowed ϕ and ψ



Beta strand region

Left-handed alpha helix region
(Rare, but short helices do happen.)

White regions are “disallowed”
because of steric clash (atoms bump)

Alpha helix region

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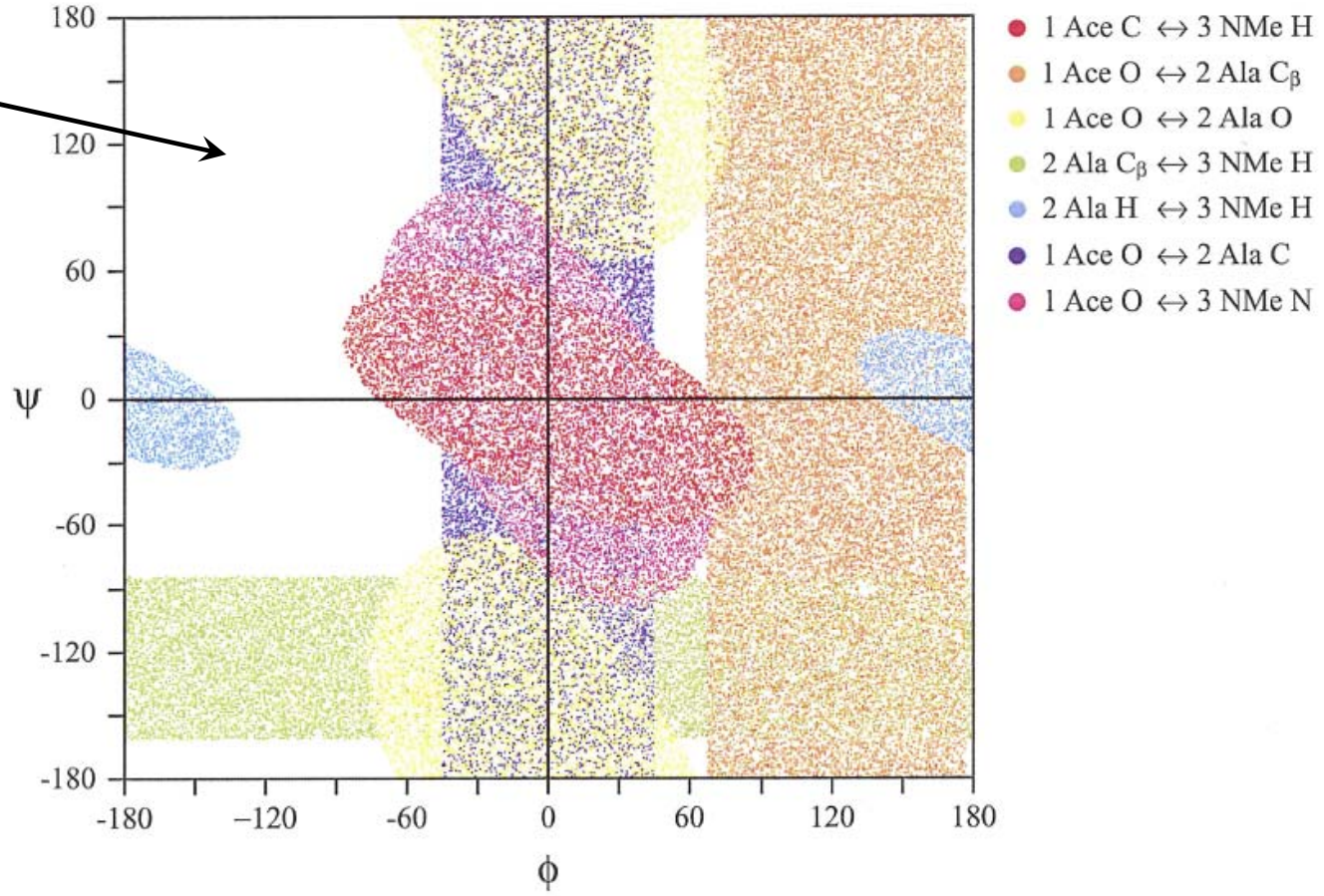
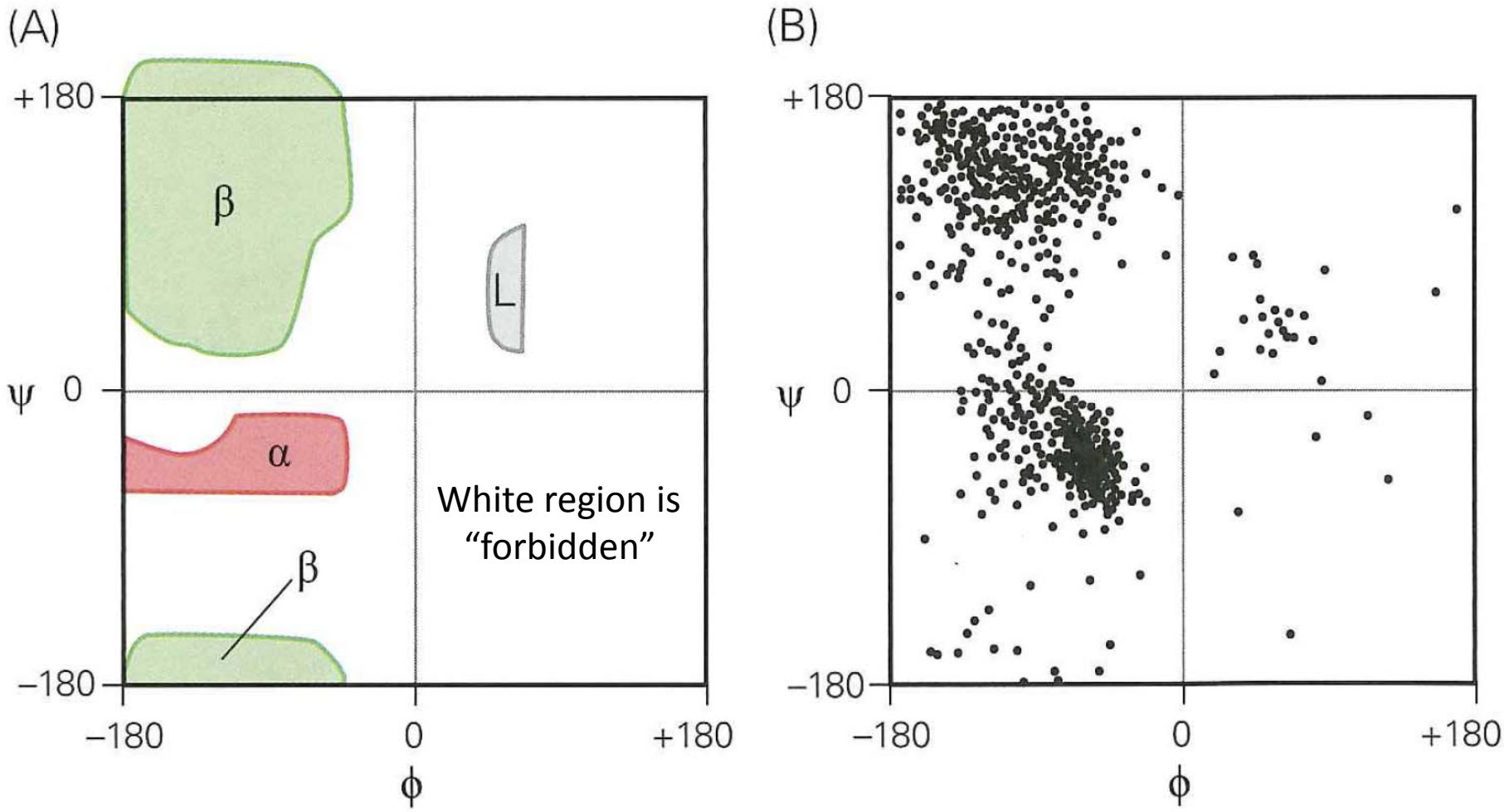
Ramachandran Reversed

White regions are allowed, colored dots represent steric clash.



Ramachandran Plot



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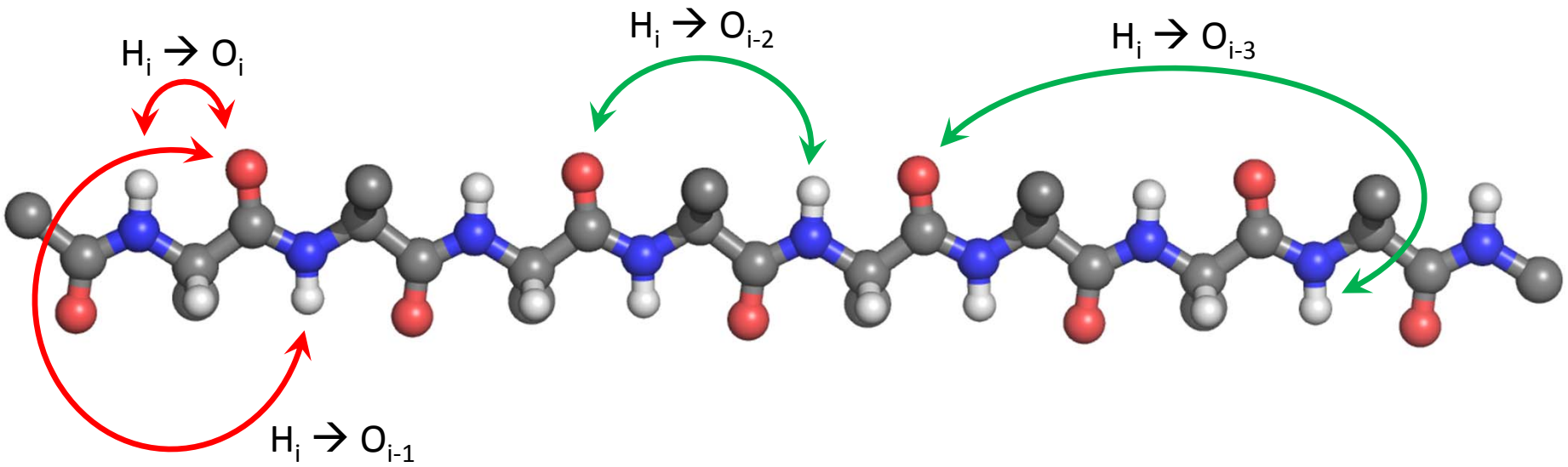
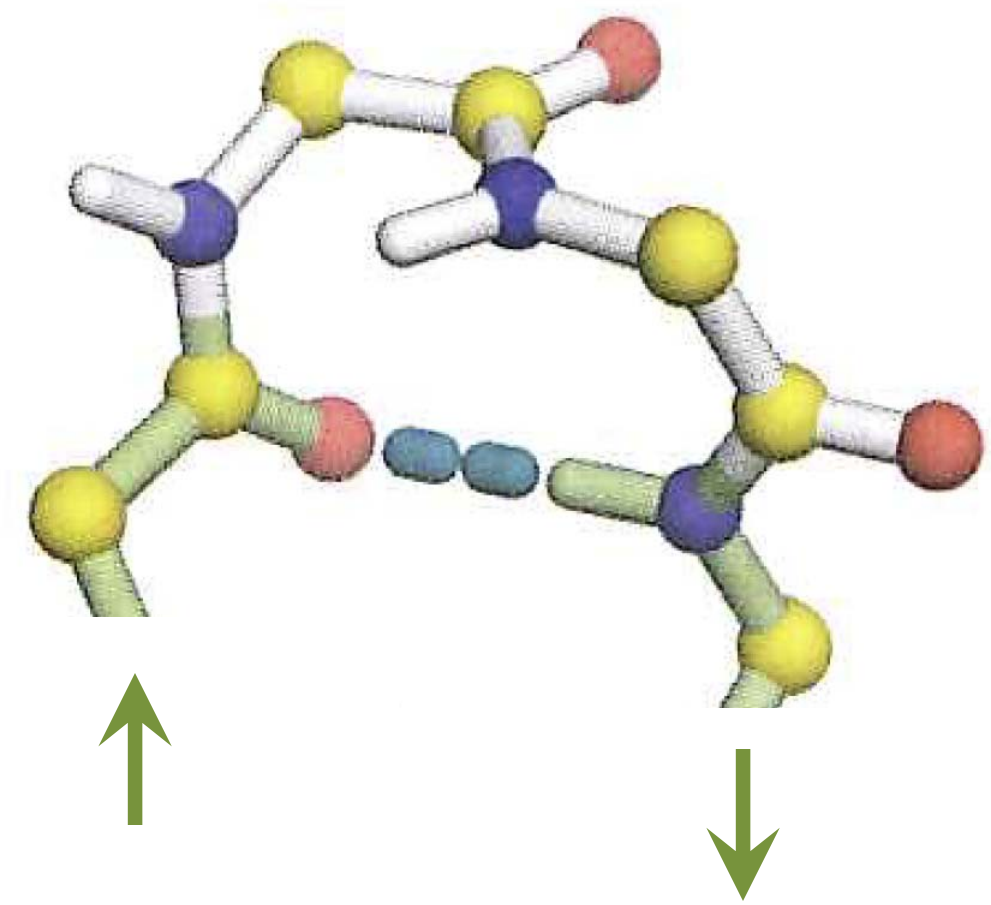
Backbone Hydrogen Bonding



- Non-repeating backbone patterns:
  - **Too much strain:** $H_i \rightarrow O_i$ and $H_i \rightarrow O_{i\pm1}$ hydrogen bonds
  - **Rare, but possible:** $H_i \rightarrow O_{i-2}$ (Gamma turns)
  - **Frequently observed:** $H_i \rightarrow O_{i-3}$ (Beta turns)

Beta Turns Reverse the Direction of the Backbone



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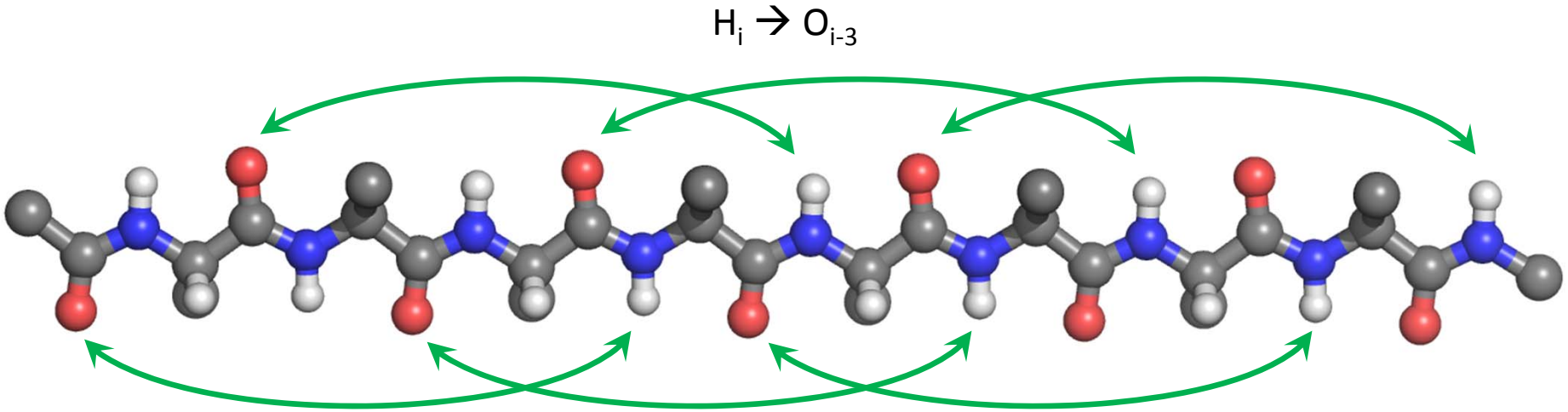
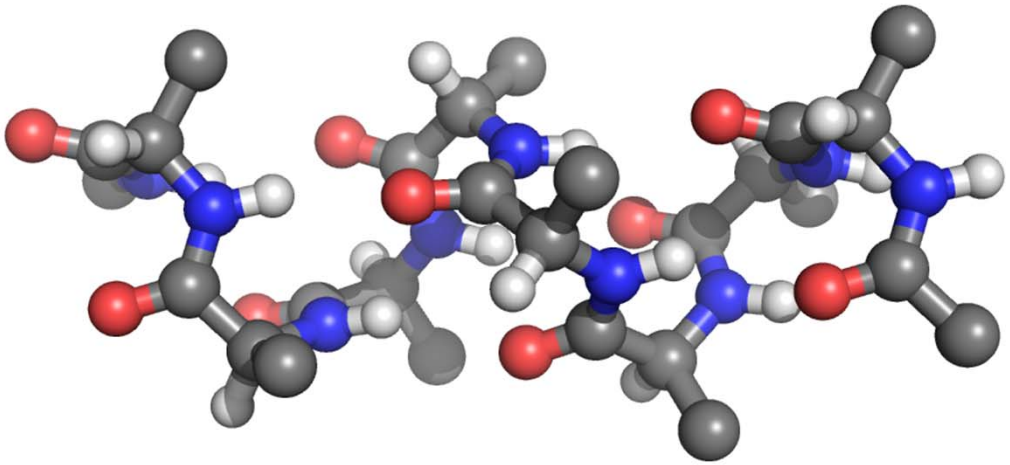
Backbone Hydrogen Bonding



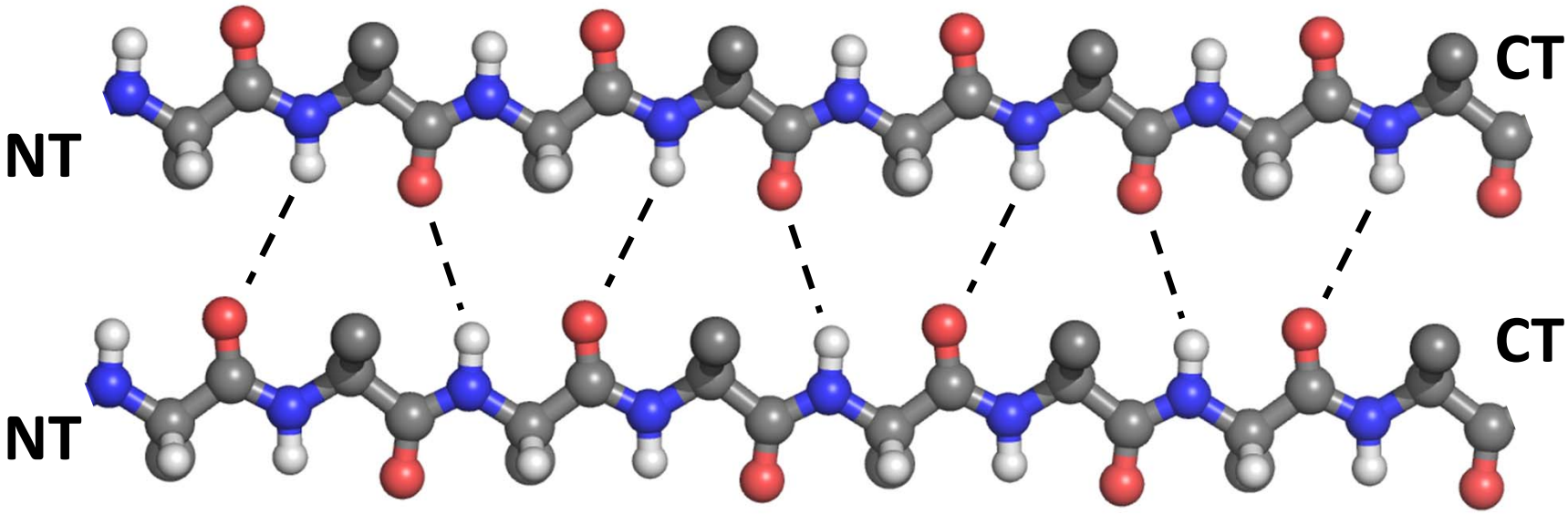
- The Alpha Helix Can be Repeated
  - $H_i \rightarrow O_{i-4}$ (Gamma turns)
  - Average ϕ is = -60^o, Average ψ is -40^o

A (Right-Handed) Alpha Helix



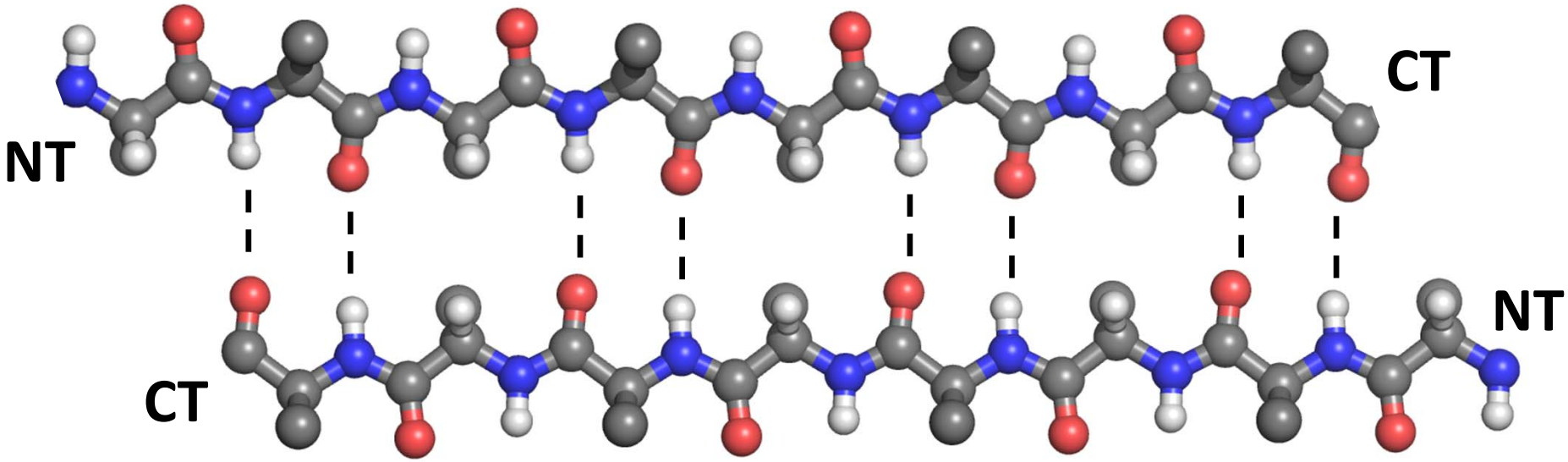
- Download helix.pdb from the course website for a model you can examine in PyMOL
- Notice that helix ends have unsatisfied H-bonds

Backbone Hydrogen Bonding



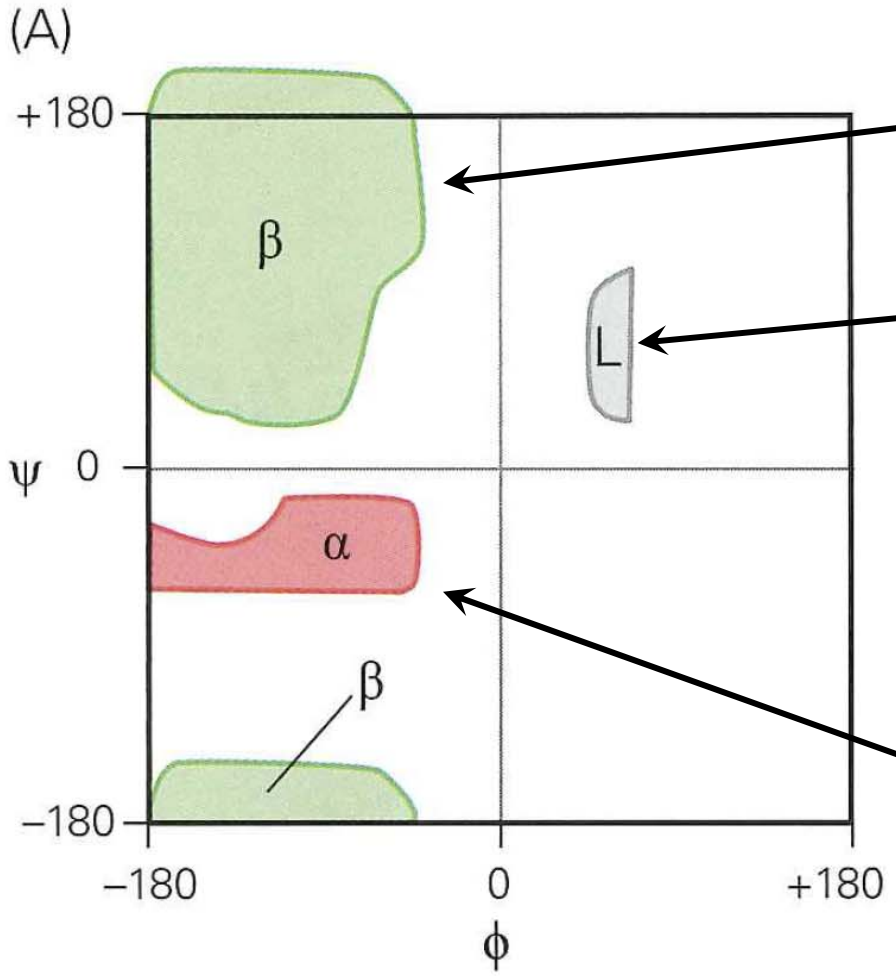
- Beta sheets are made of beta strands
  - No specified hydrogen bonding formula
  - Sheets can form between distant sets of residues
  - **Shown:** Parallel beta sheet

Backbone Hydrogen Bonding



- Beta sheets are made of beta strands
  - No specified hydrogen bonding formula
  - Average ϕ is = -120^{o}, Average ψ is 120^{o} (with large variation)
  - Sheets can form between distant sets of residues
  - **Shown:** Antiparallel beta sheet

Ramachandran Revisited



Beta strand region

Left-handed alpha helix region
(Rare, but short helices do happen.)

Alpha helix region

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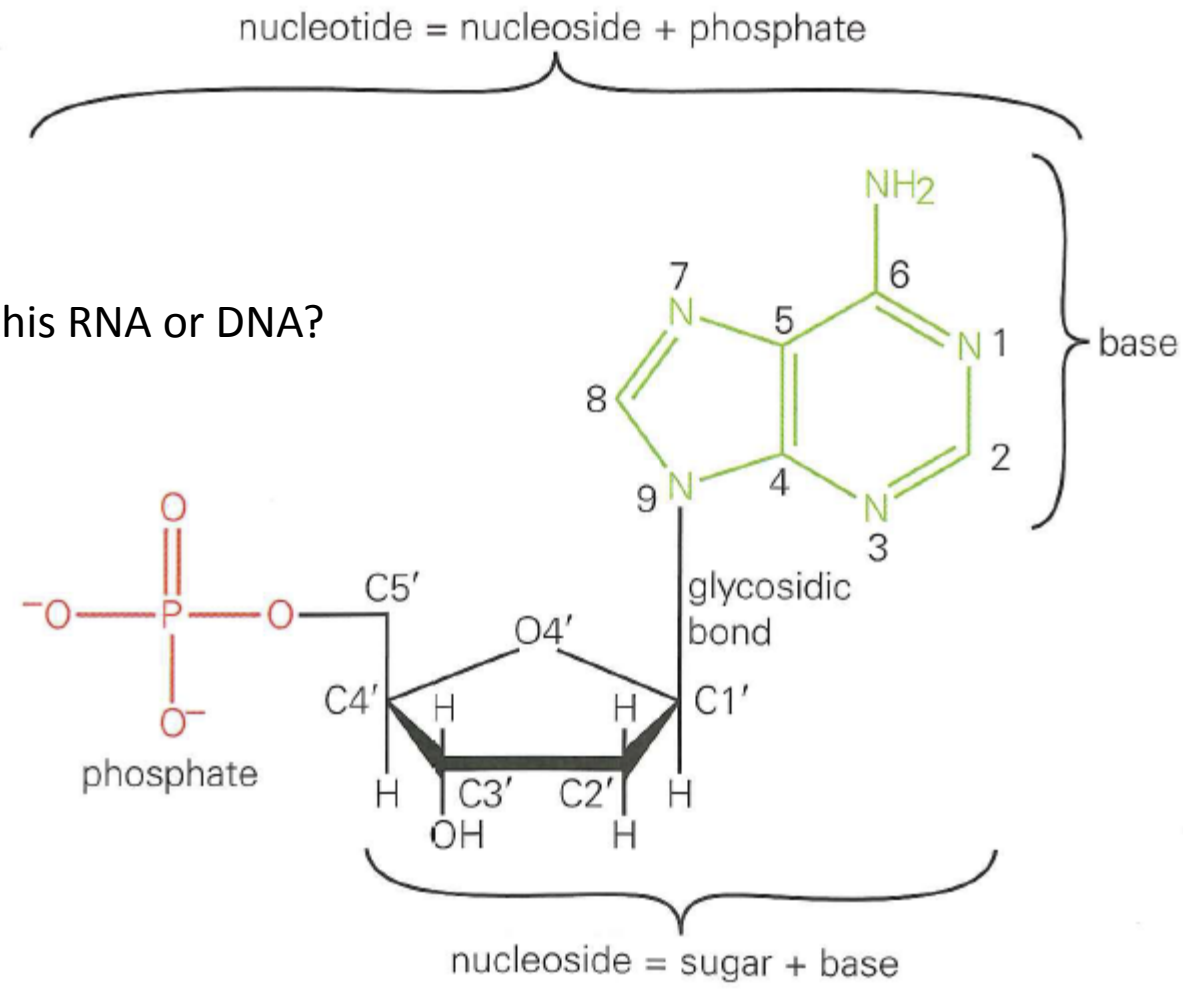
Nucleic Acid Structure

Question: Is this RNA or DNA?



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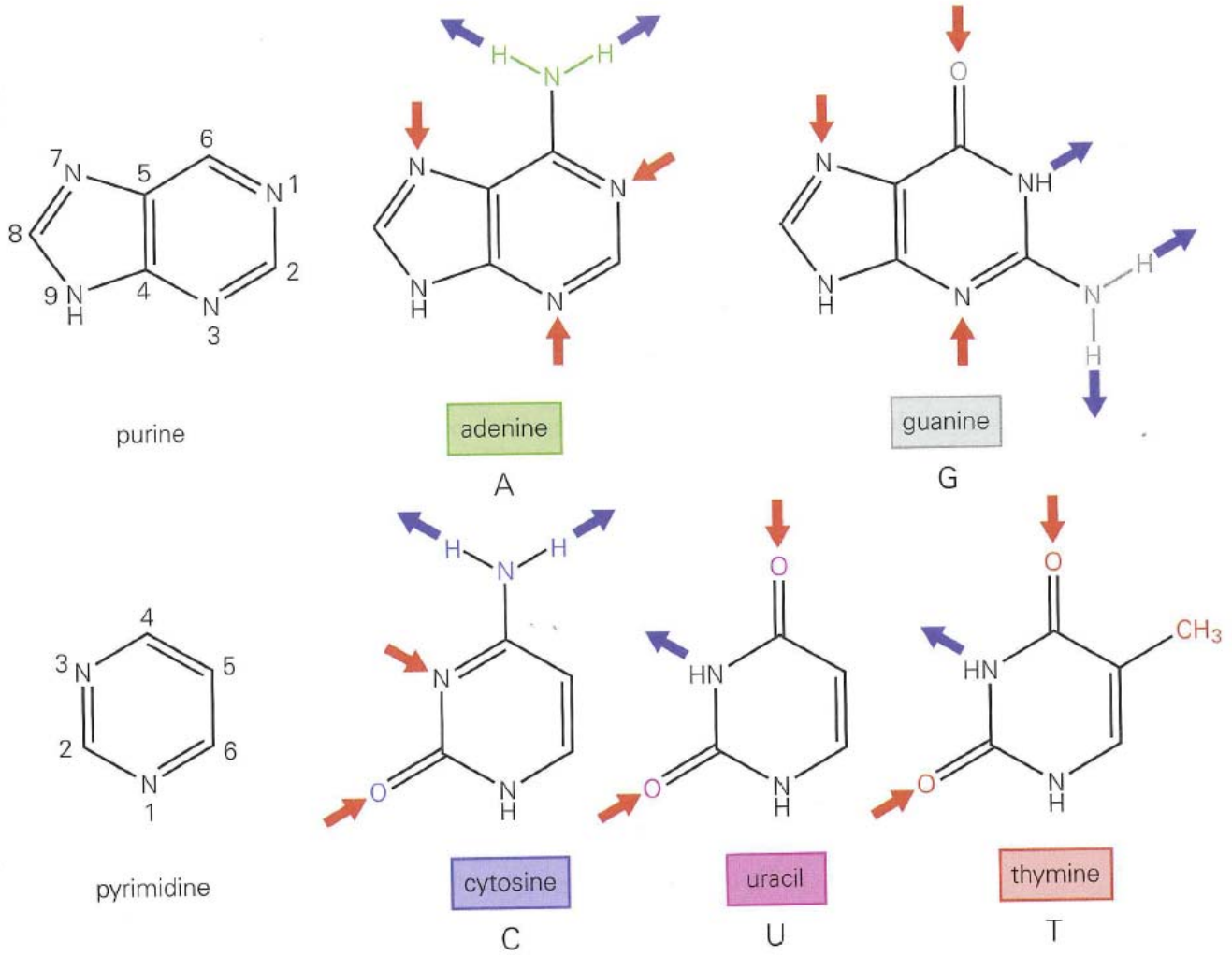
Nucleic Acid Bases

(You must memorize these.)



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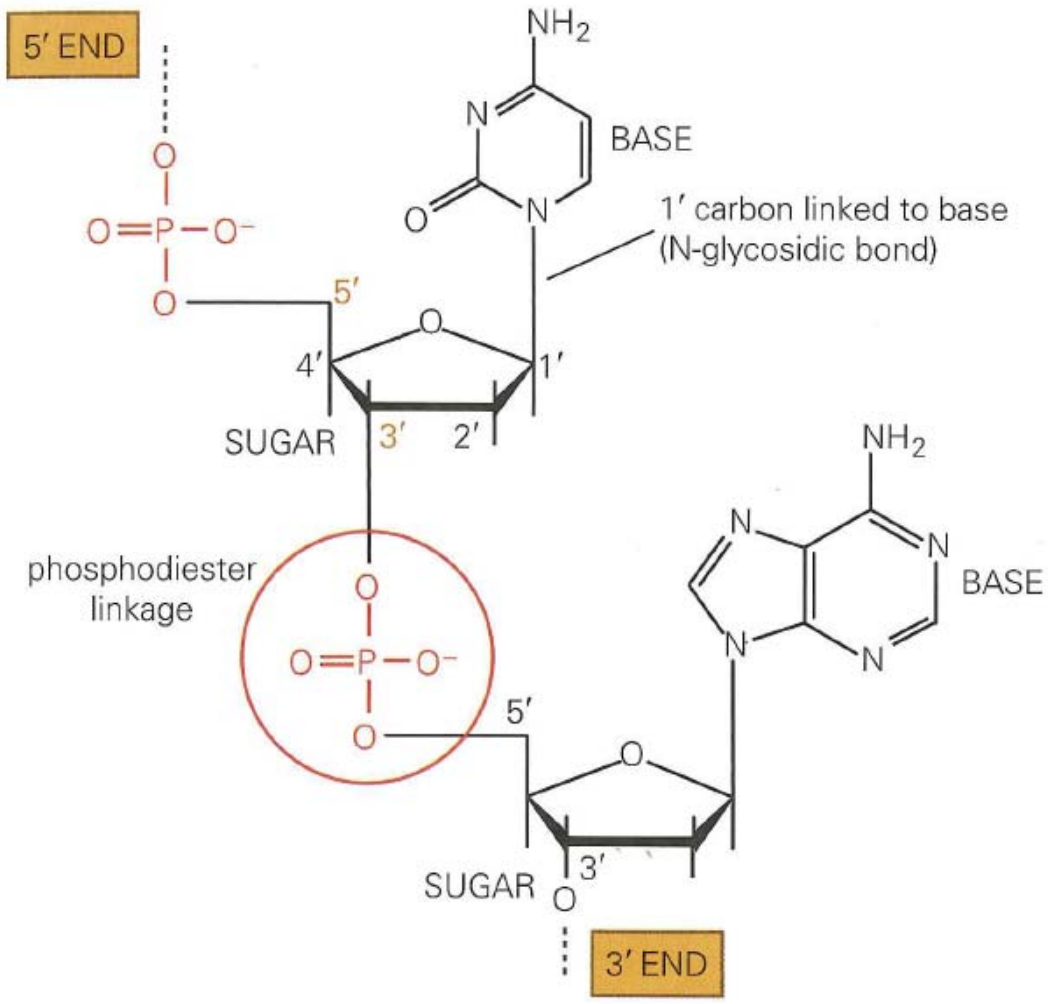
Nucleic Acids Are Also Polymers



DNA & RNA Polymerase: Build up DNA and RNA from nucleoside triphosphates (5' → 3' synthesis)

Convention: RNA/DNA typically is read from 5' to 3' direction (e.g. 5'-ATTGCAAC-3')

And yes, you must have the backbone structure of DNA memorized, too. (The level of detail in this slide is fine.)

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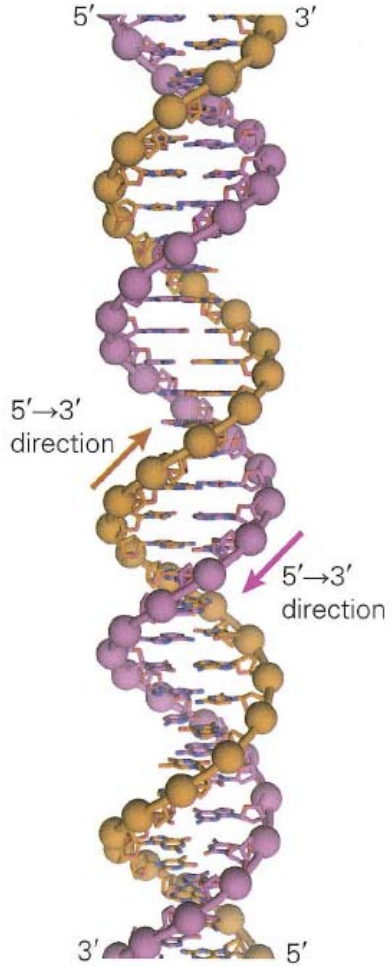
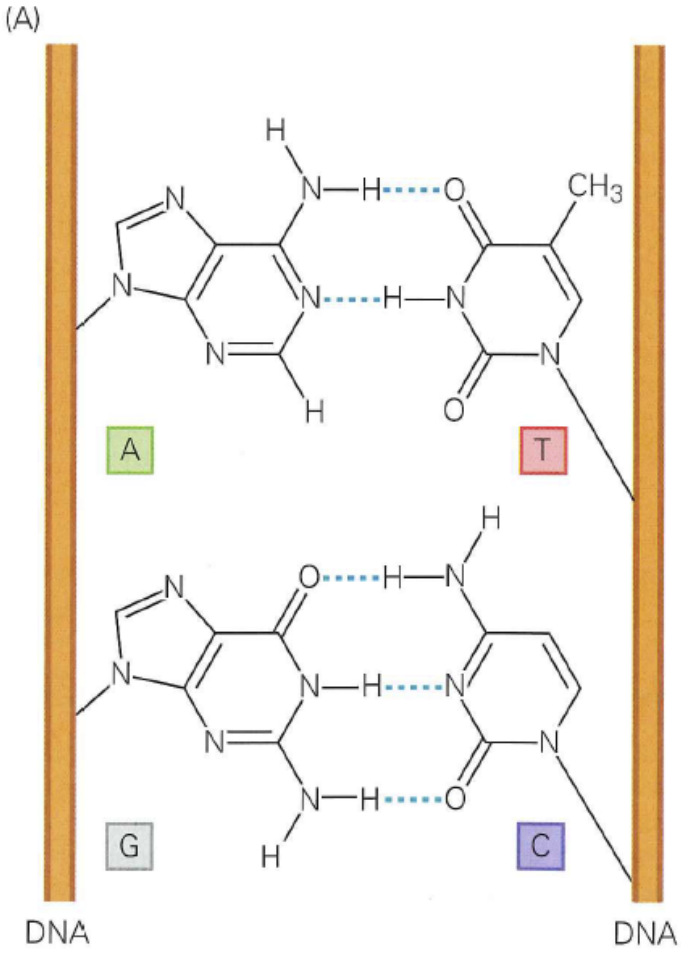
Watson-Crick Base Pairing in an (Antiparallel) Double Helix





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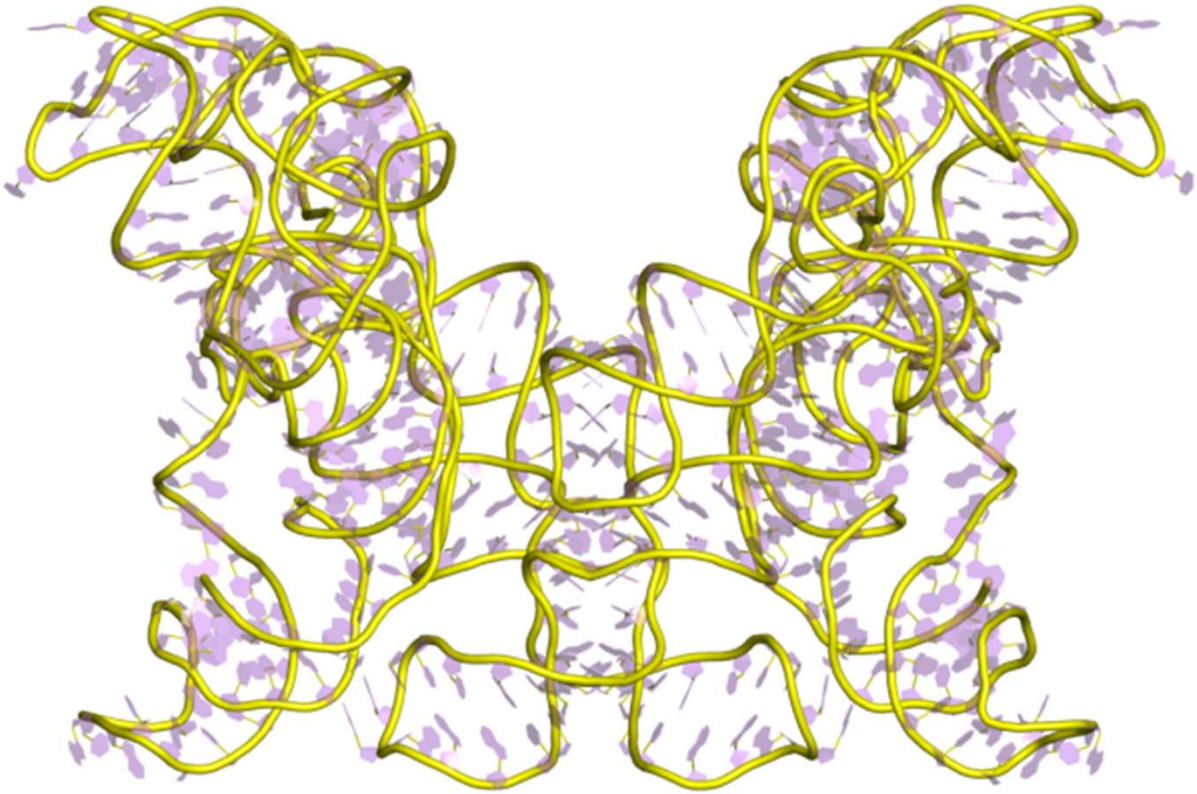
Diversity of Nucleic Acid Structure



Ribozyme: An RNA capable of catalyzing a chemical reaction

The ribosome contains a significant amount of RNA as well as proteins

Macromolecules can perform incredibly diverse structures!
(And we haven't even mentioned lipids and sugars.)

Wikipedia, "Group I Catalytic Intron." Accessed 8/23/2012.