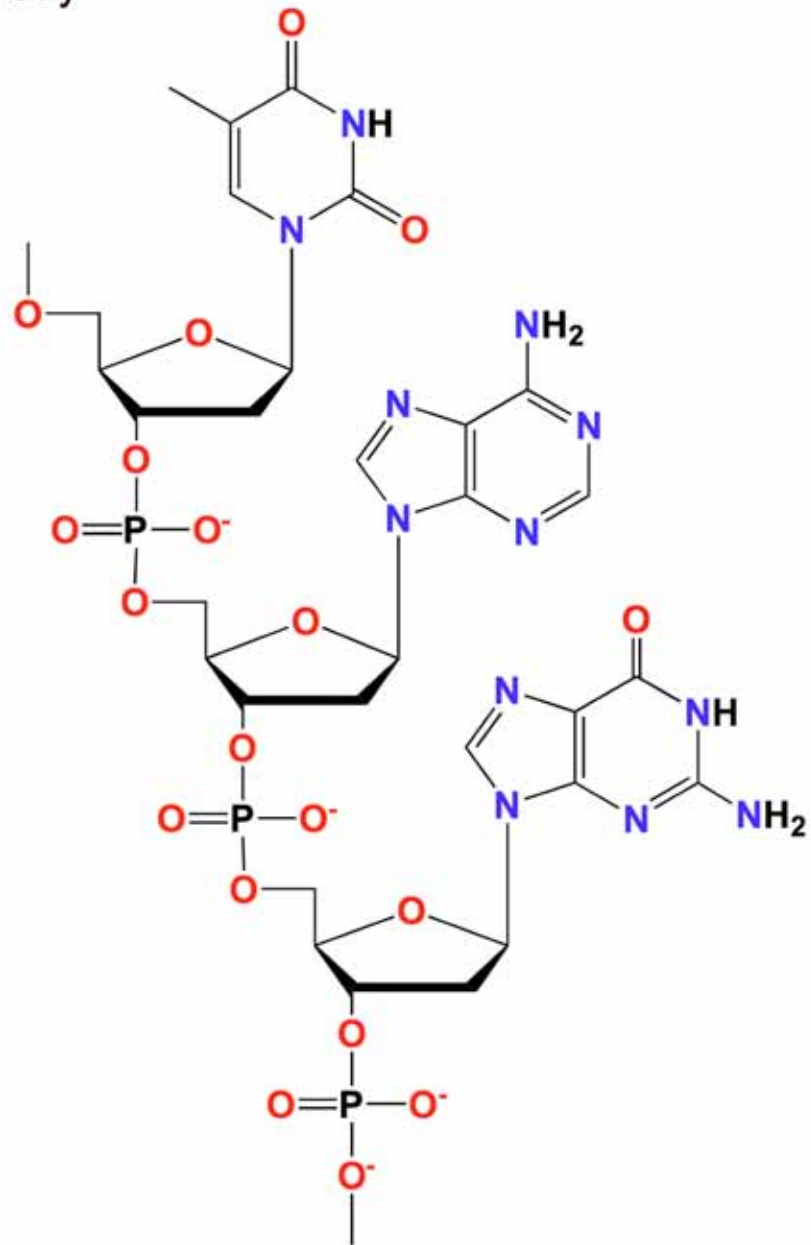
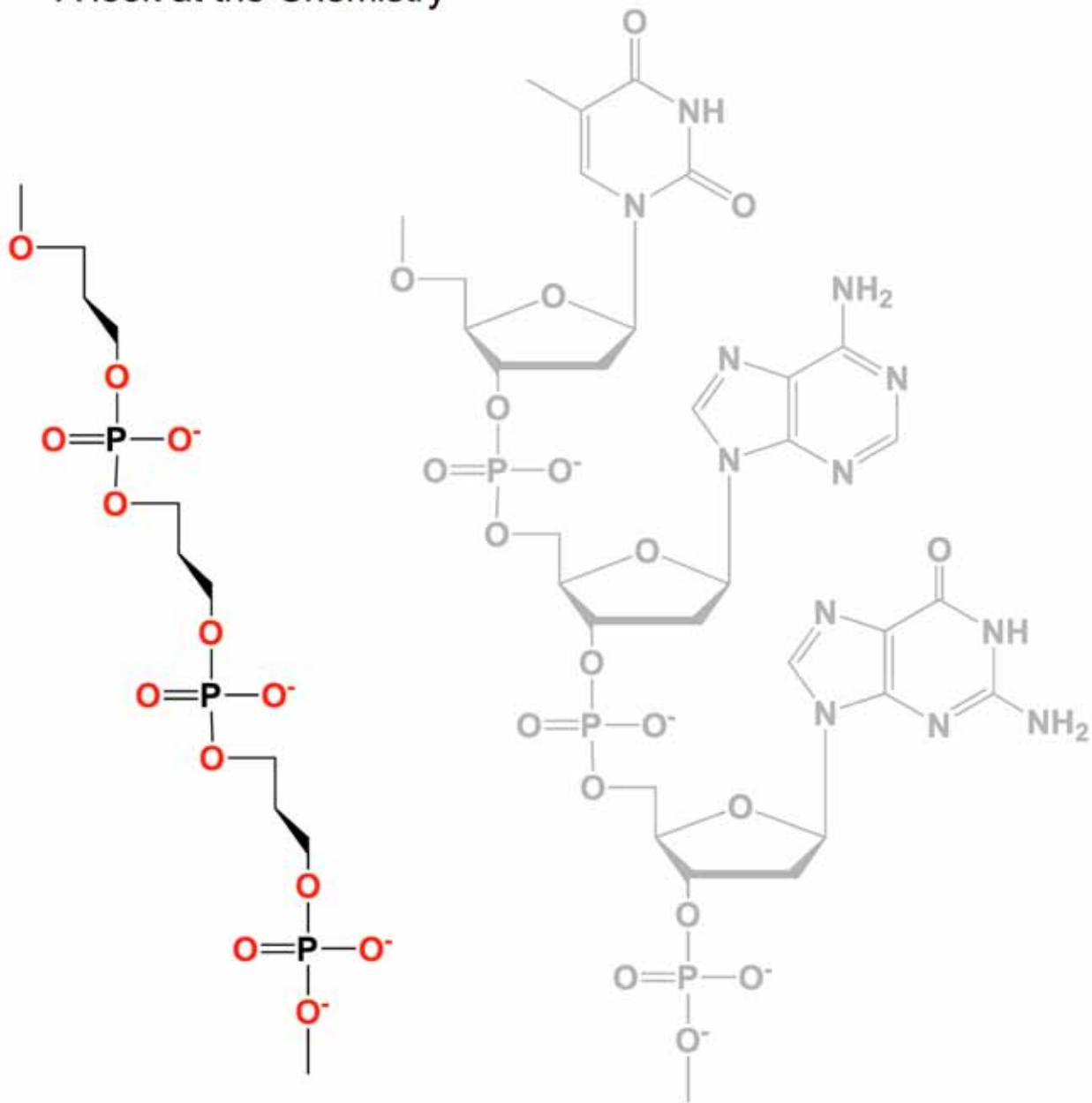


DNA

A look at the Chemistry

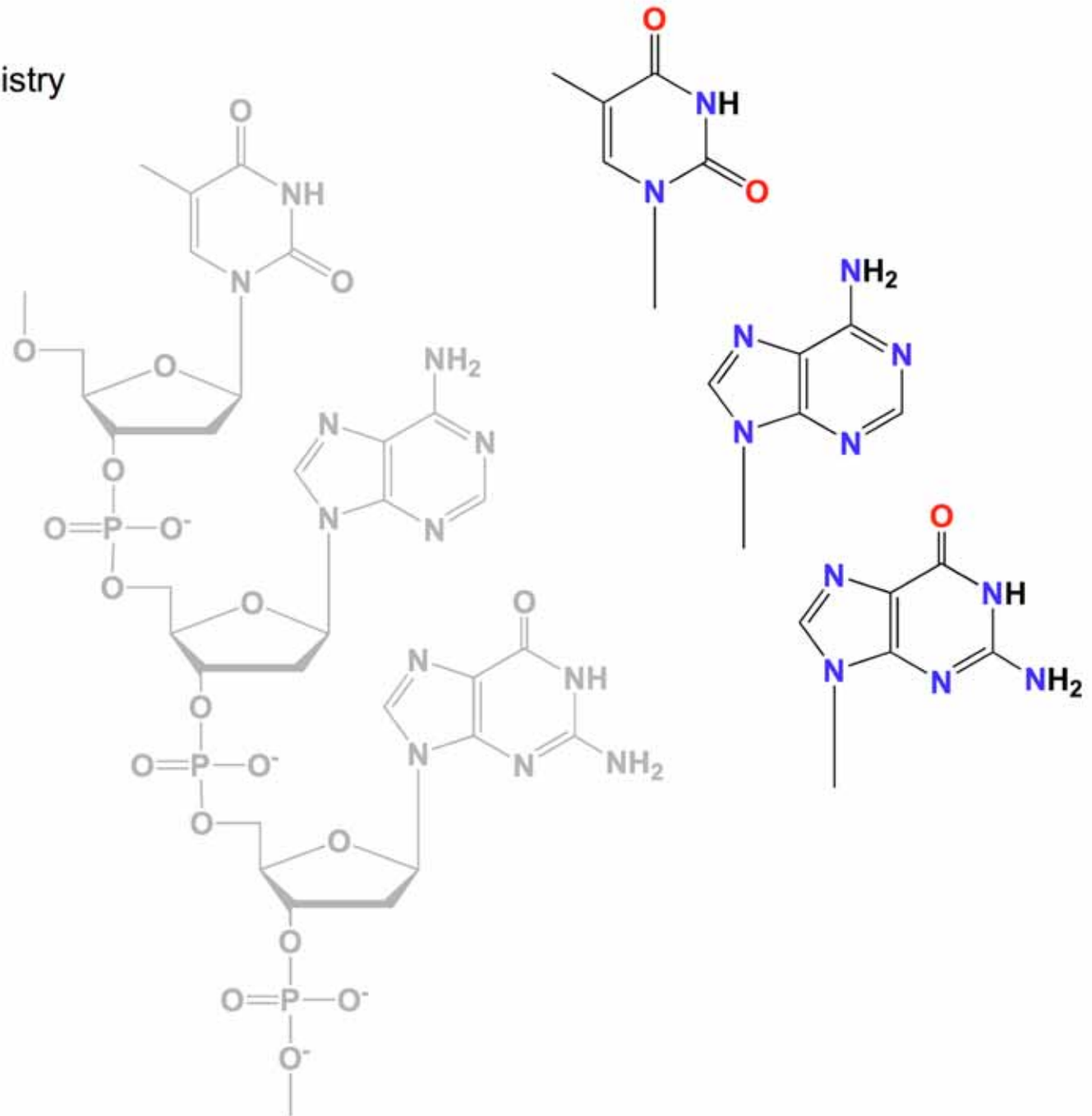


A look at the Chemistry



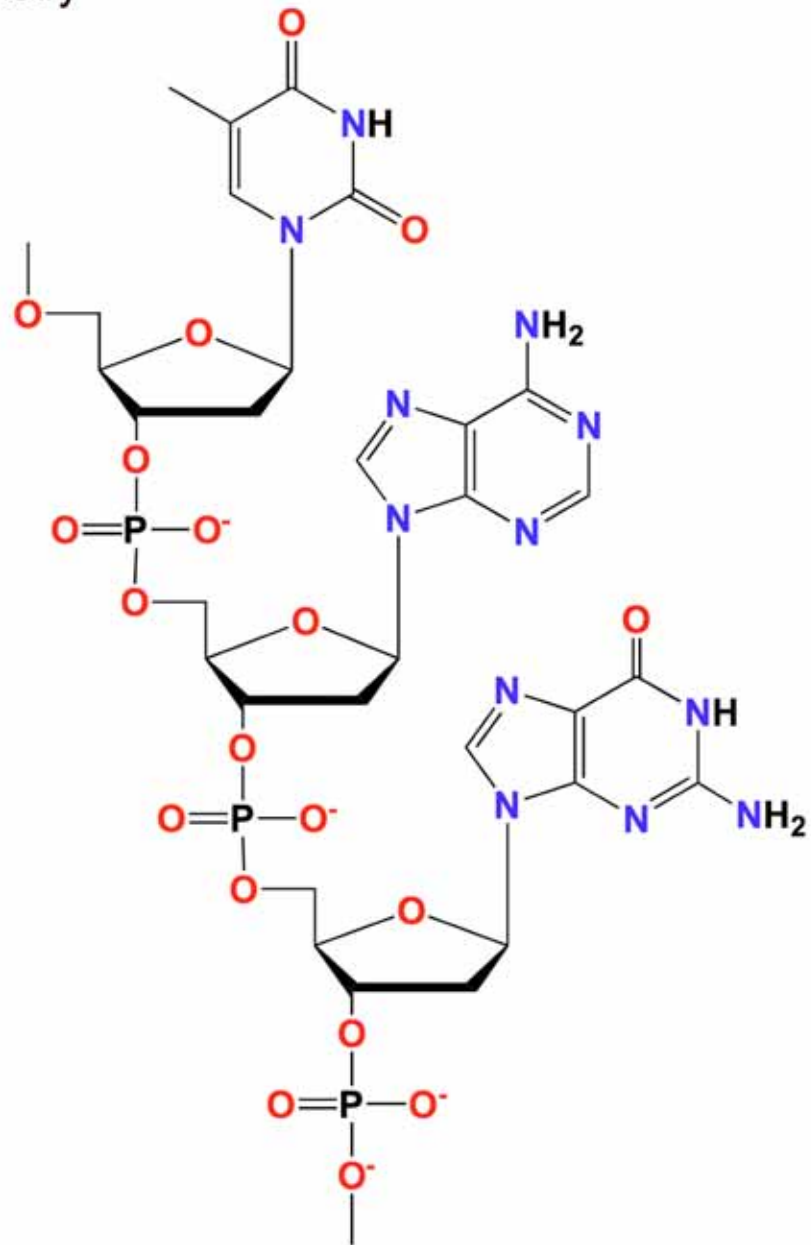
DNA

A look at the Chemistry



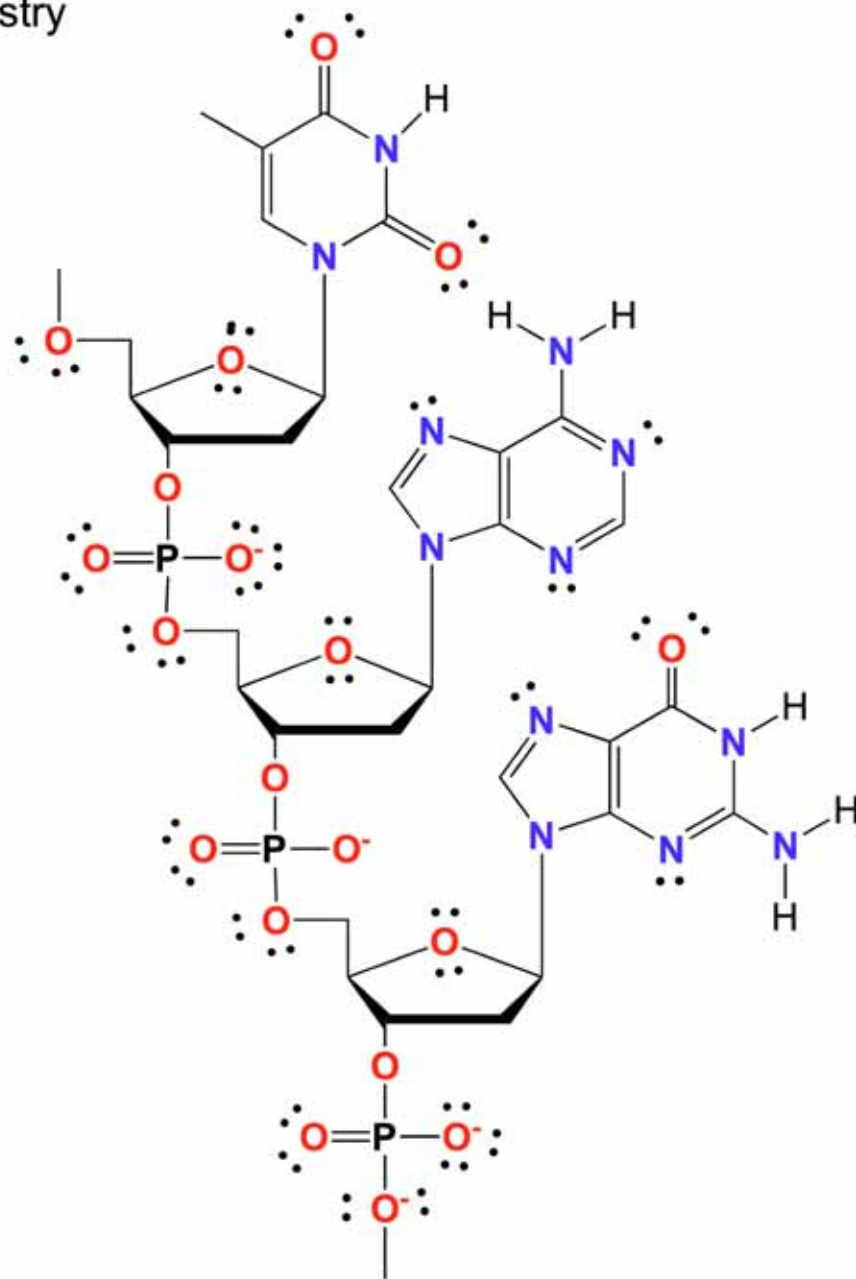
DNA

A look at the Chemistry



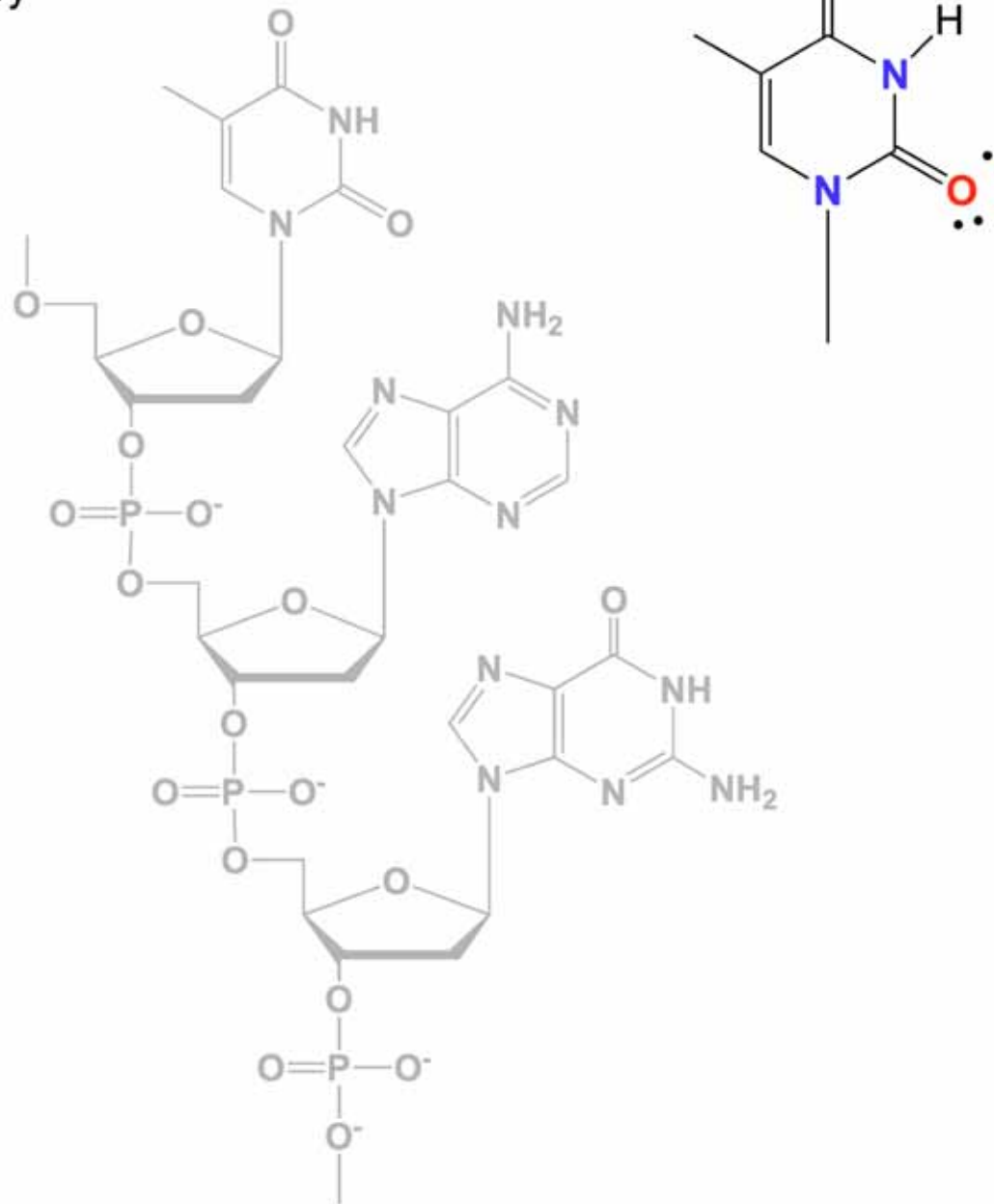
DNA

A look at the Chemistry



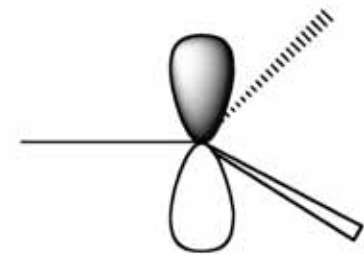
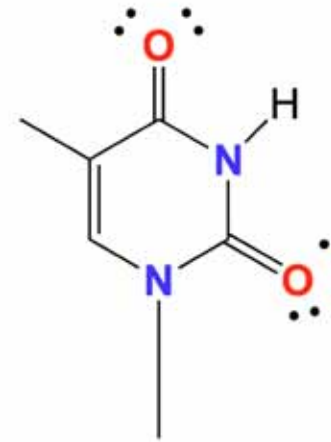
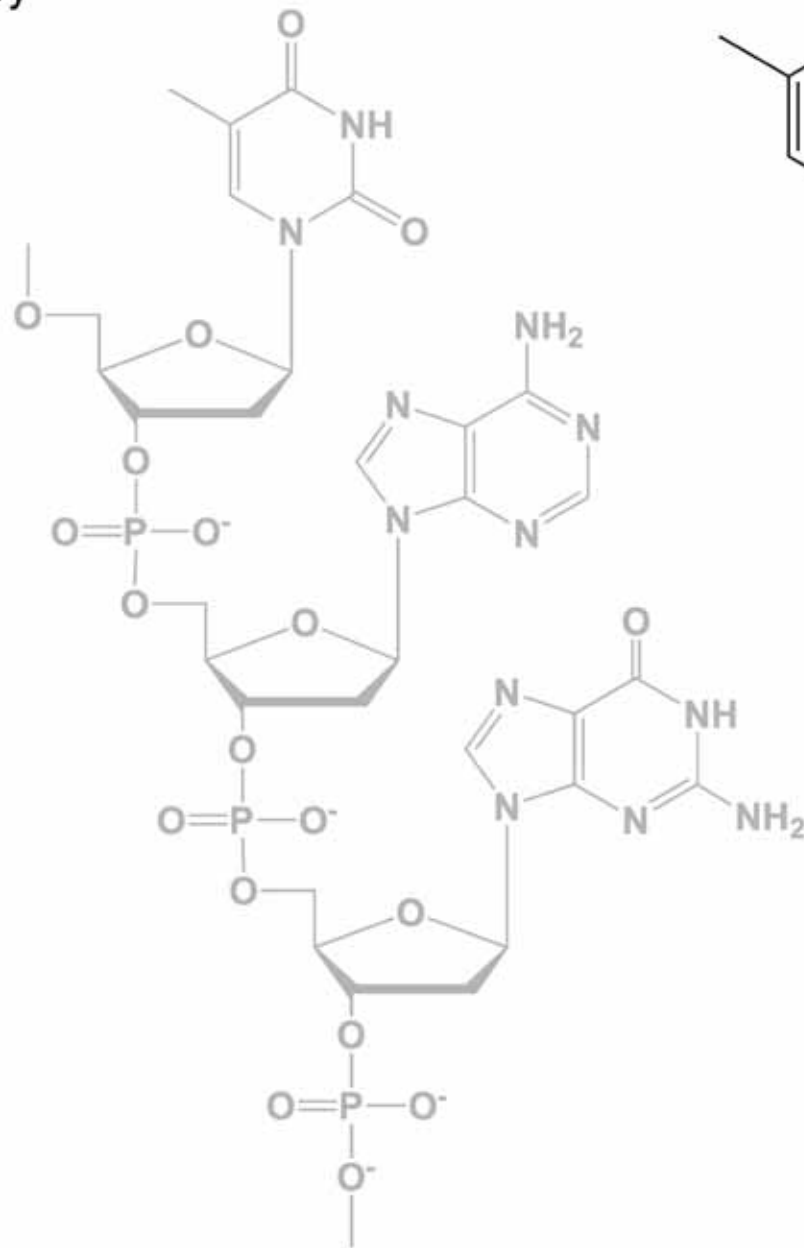
DNA

A look at the Chemistry



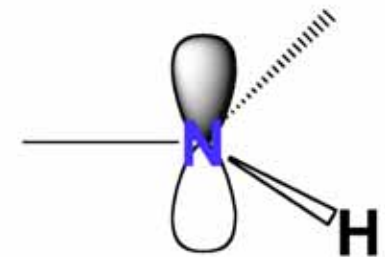
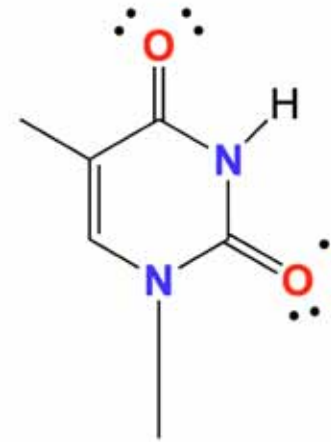
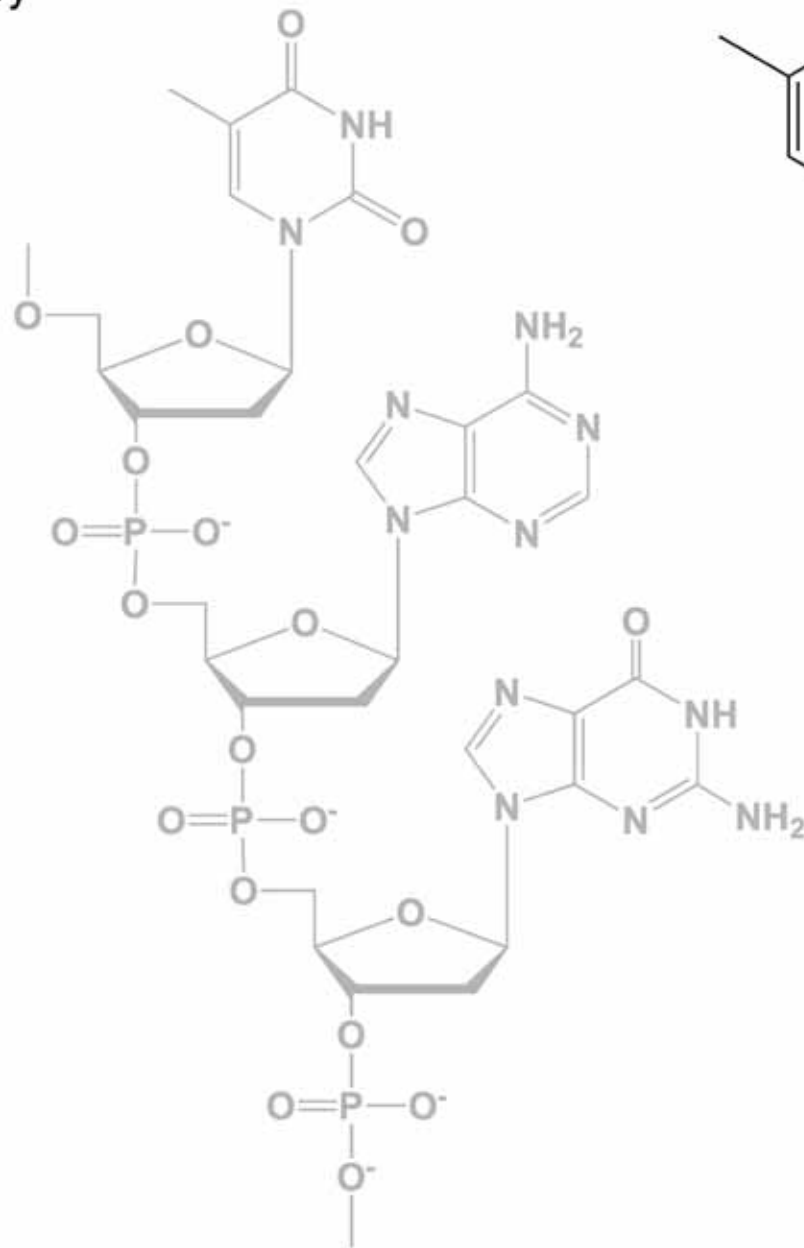
DNA

A look at the Chemistry



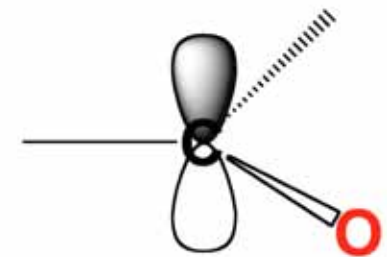
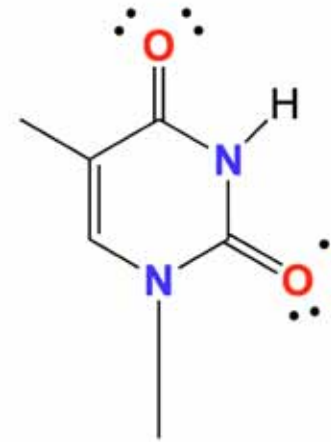
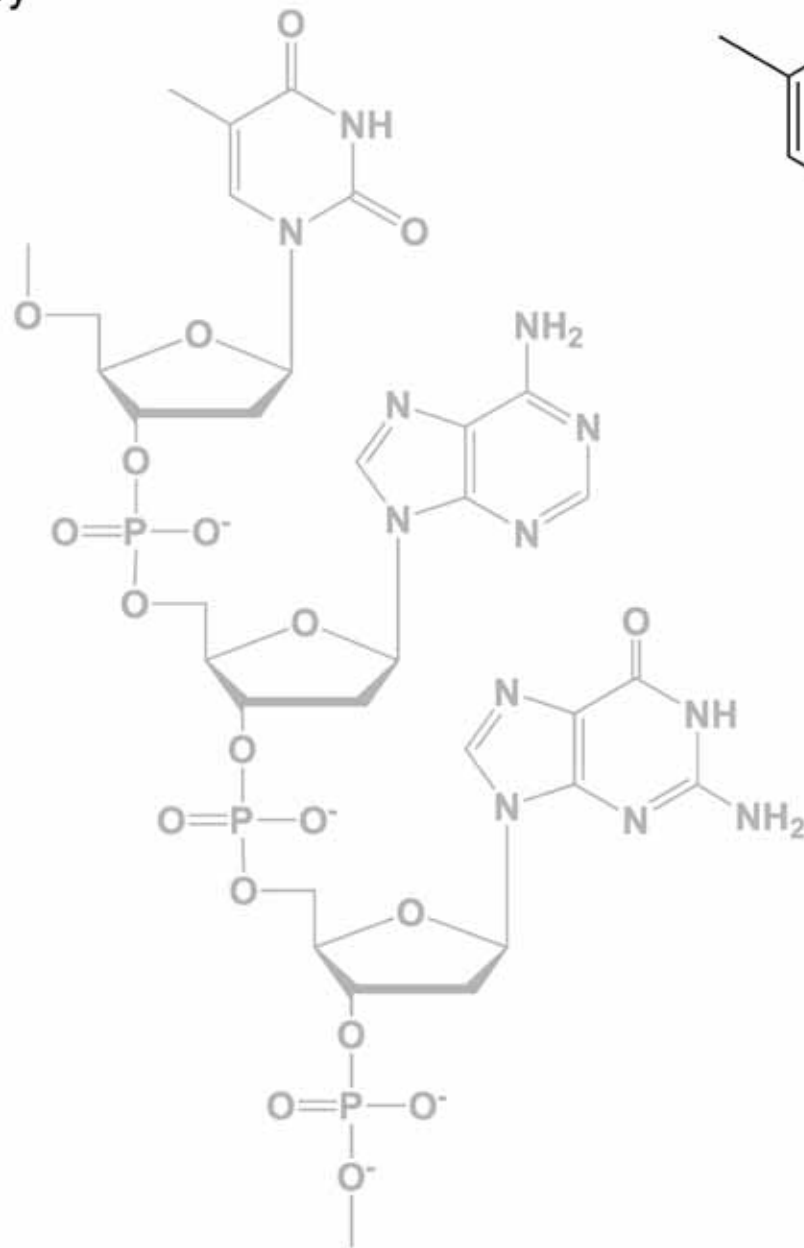
DNA

A look at the Chemistry



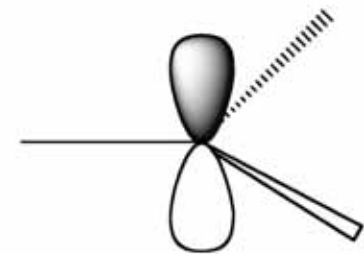
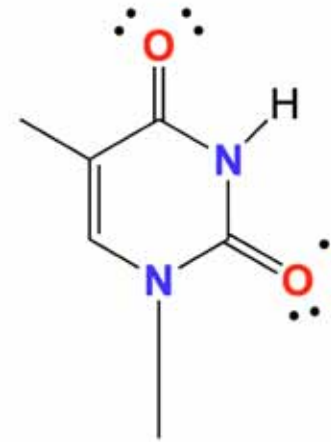
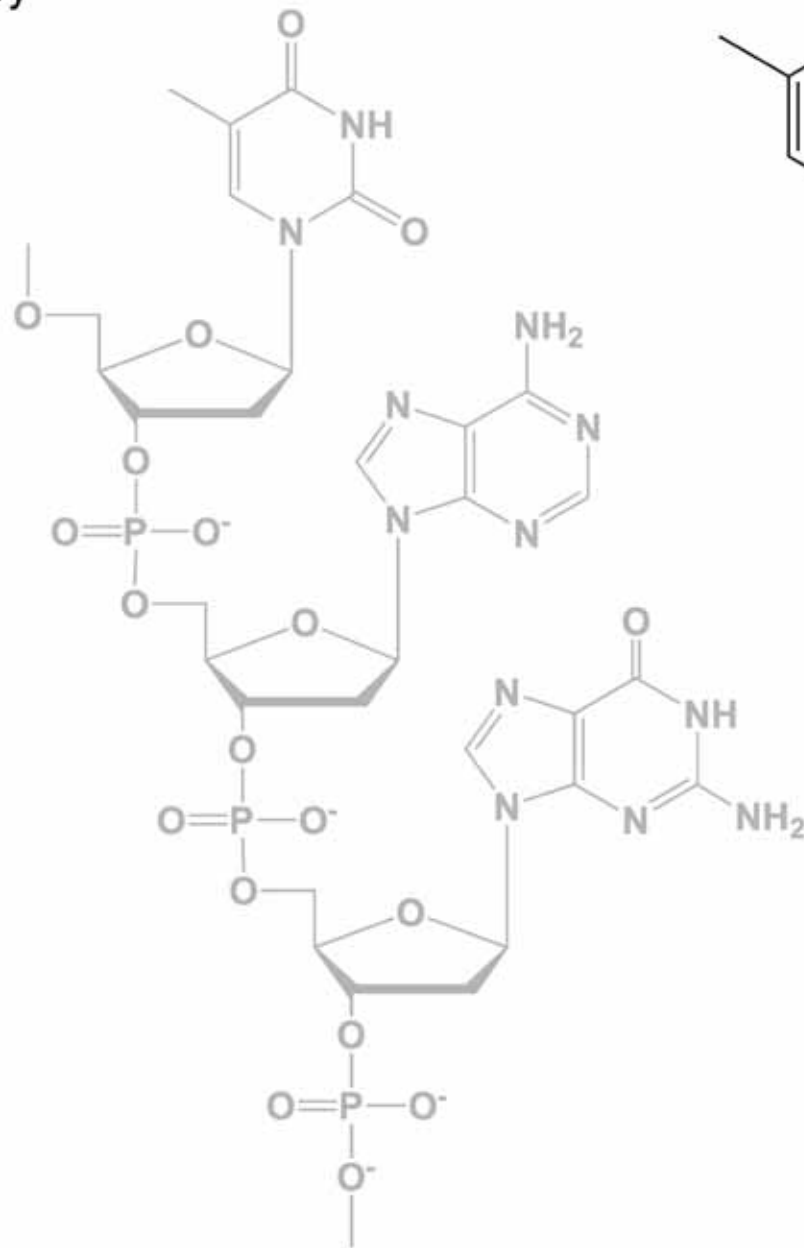
DNA

A look at the Chemistry



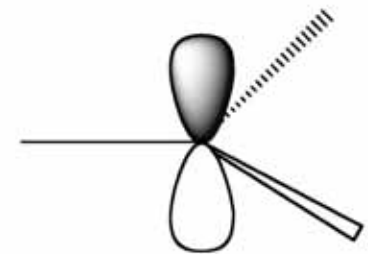
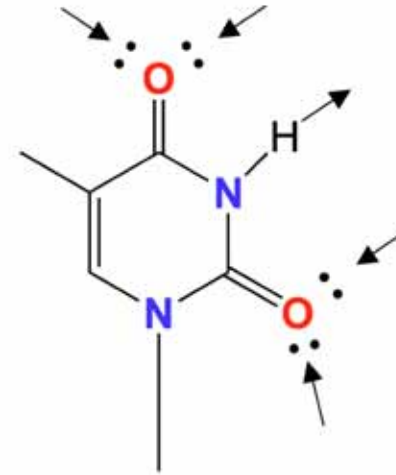
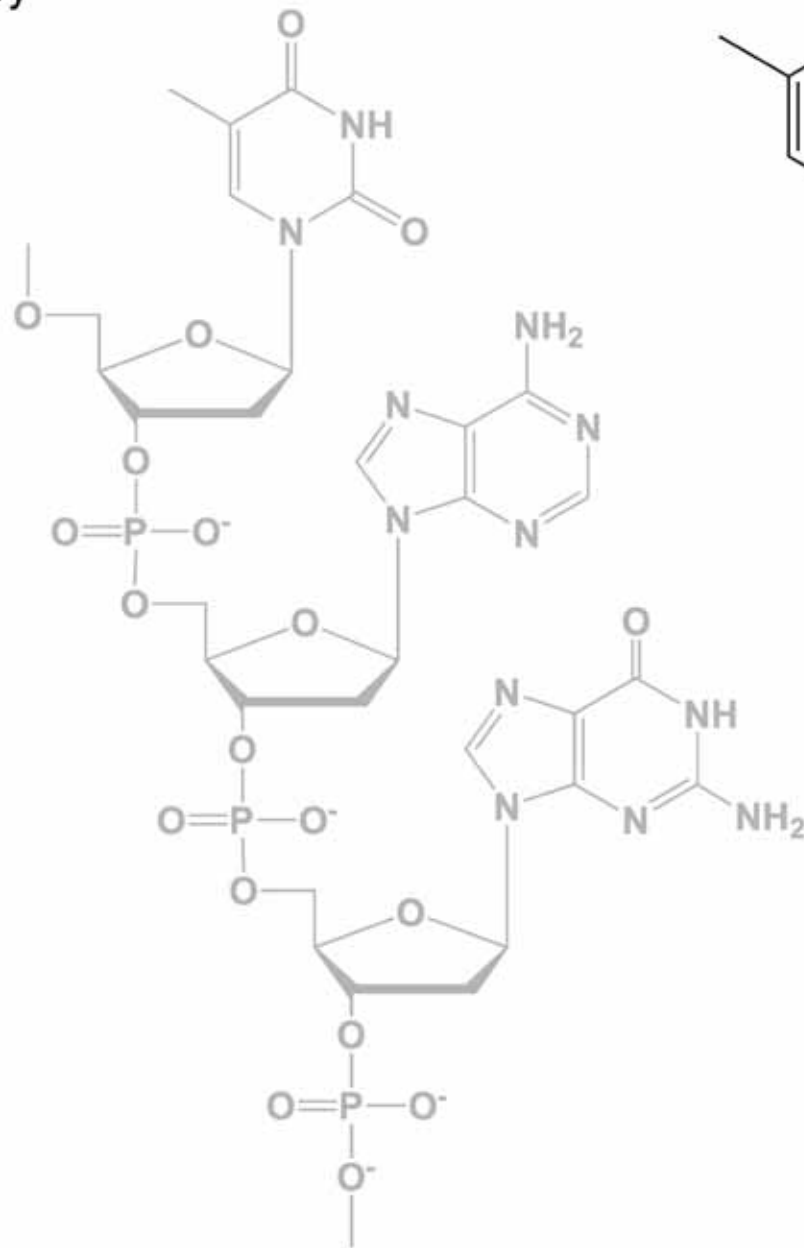
DNA

A look at the Chemistry



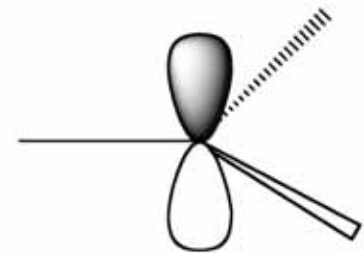
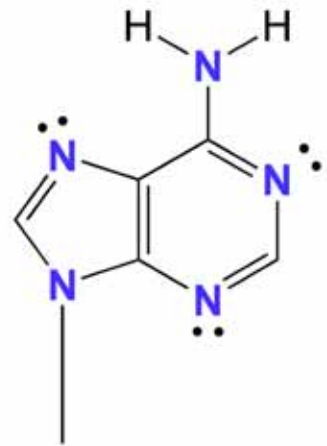
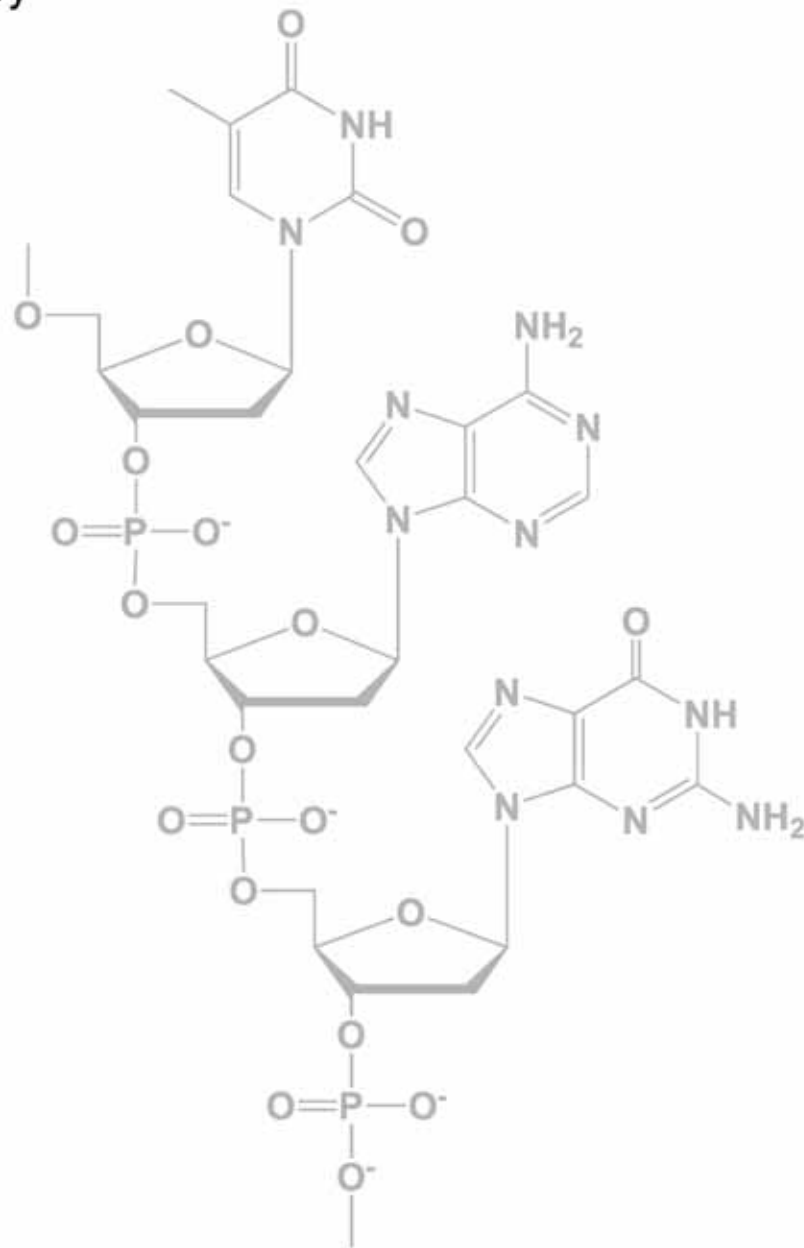
DNA

A look at the Chemistry



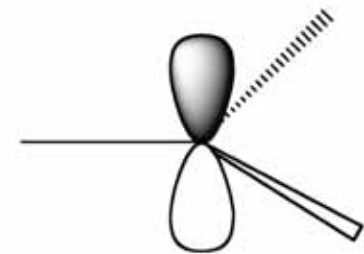
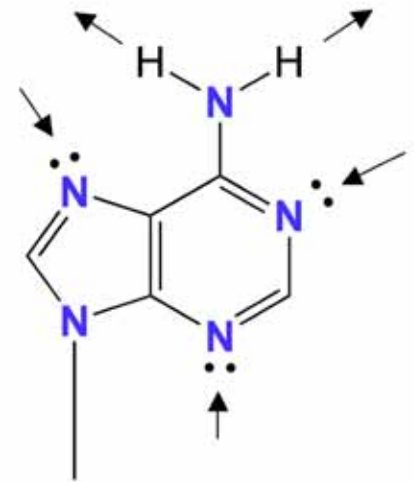
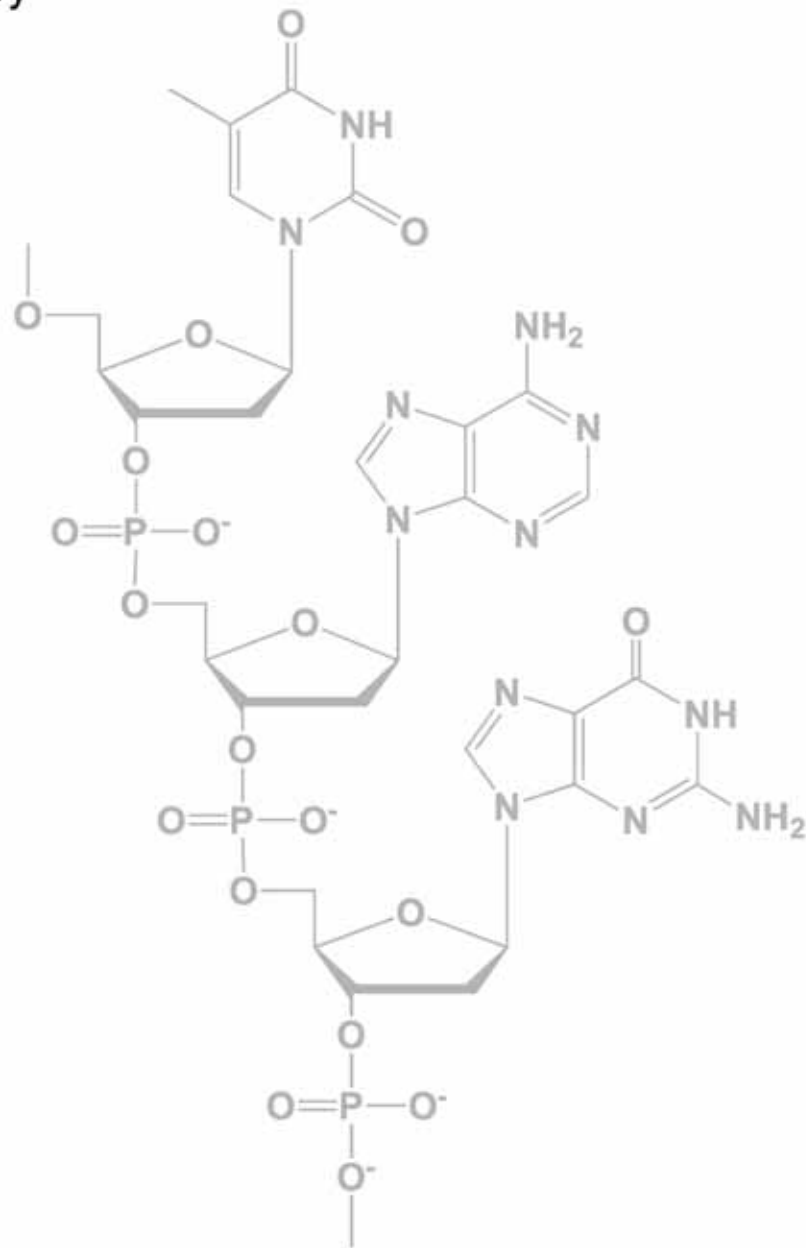
DNA

A look at the Chemistry



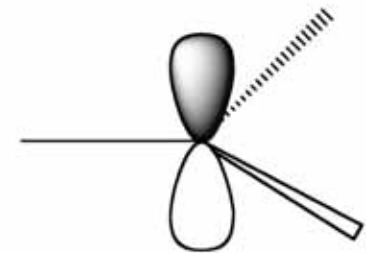
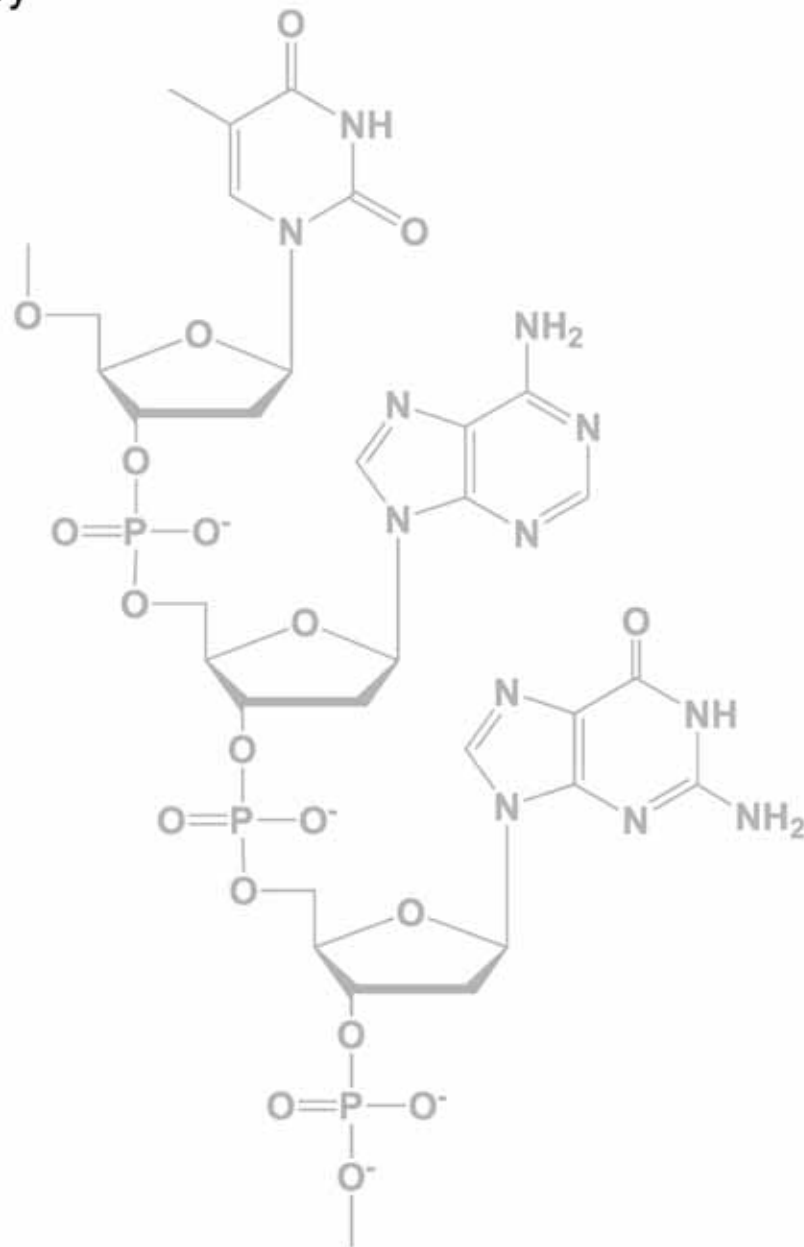
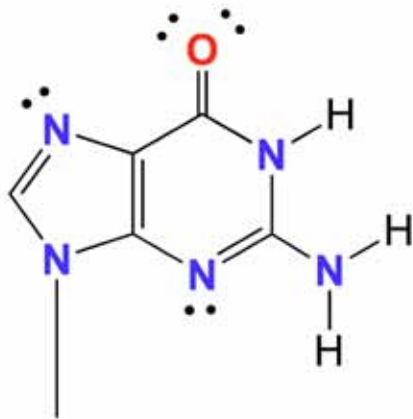
DNA

A look at the Chemistry



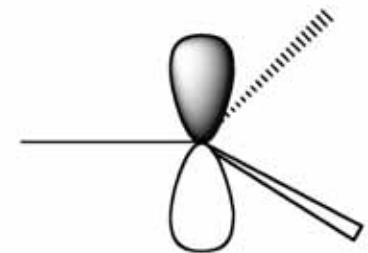
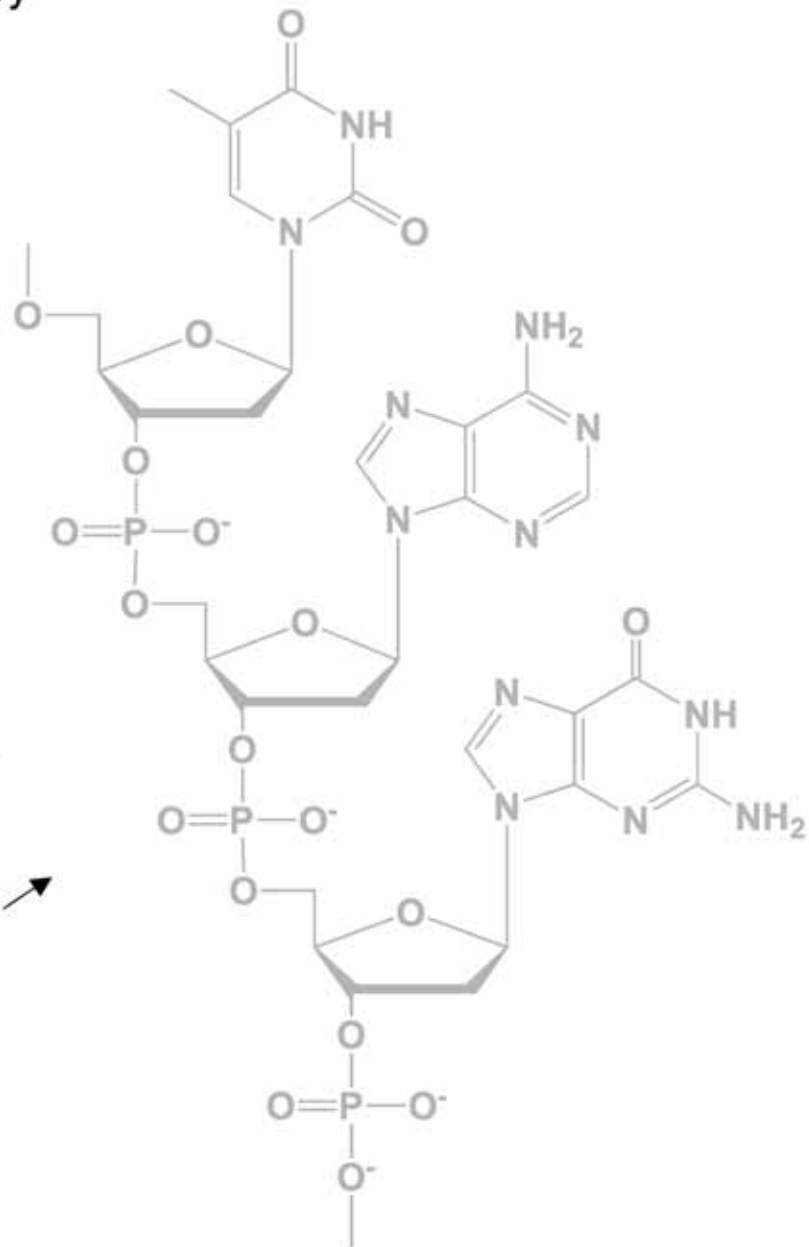
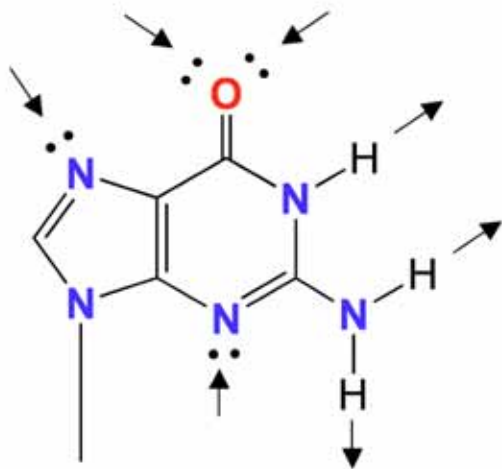
DNA

A look at the Chemistry

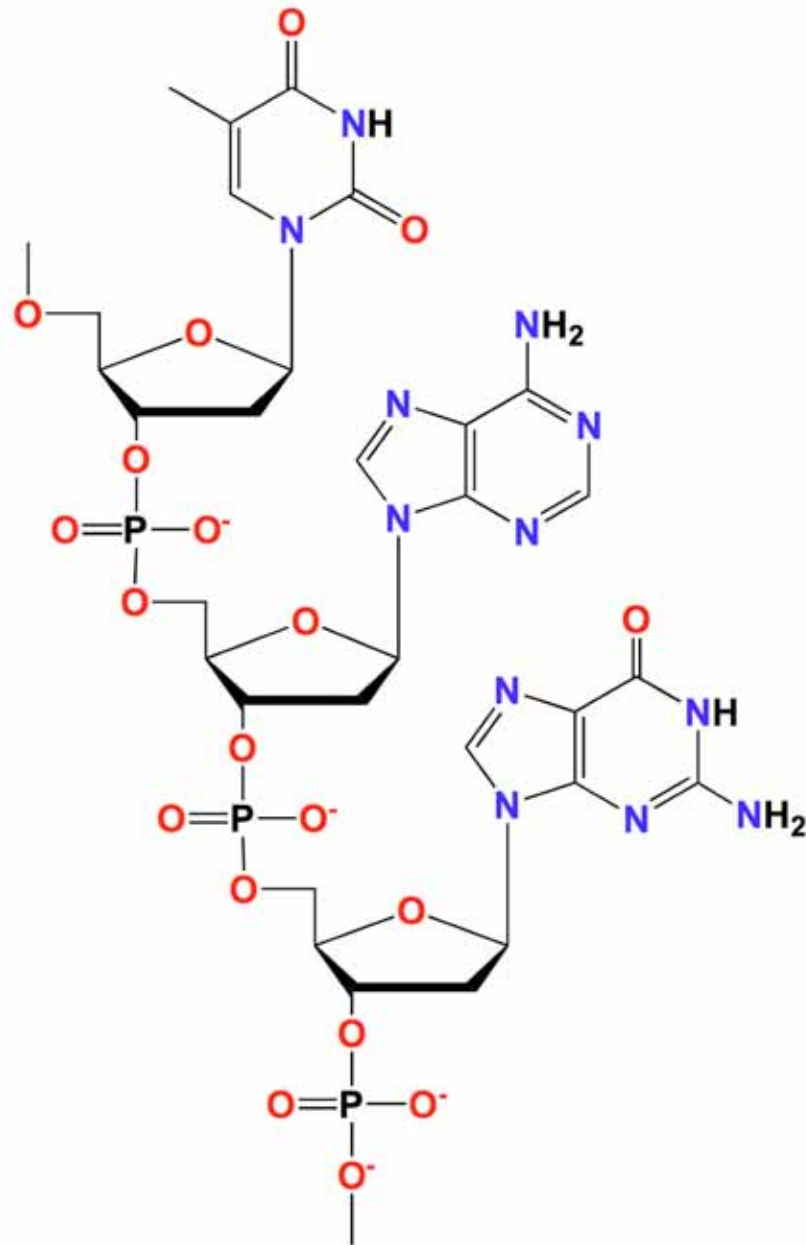


DNA

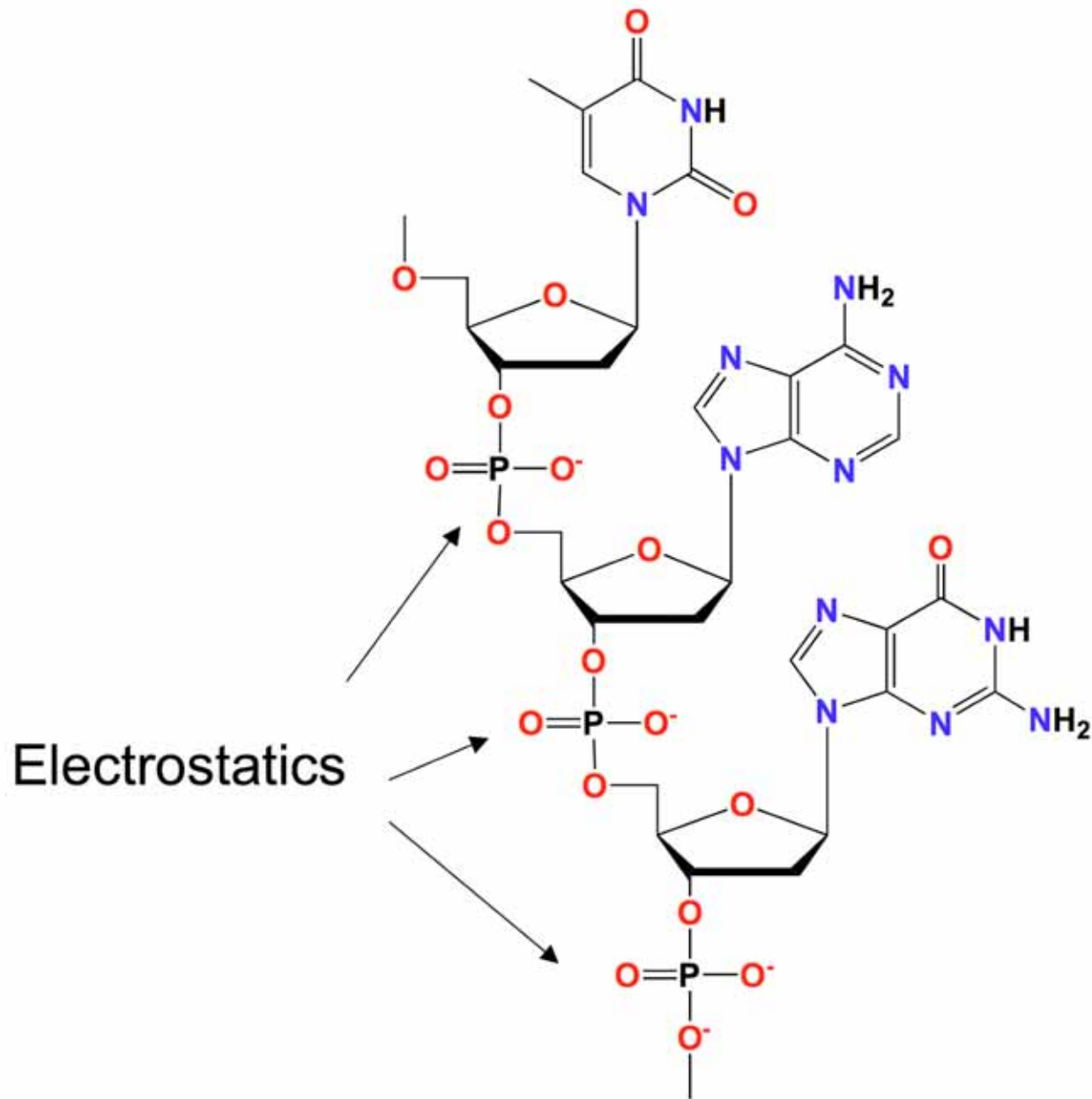
A look at the Chemistry



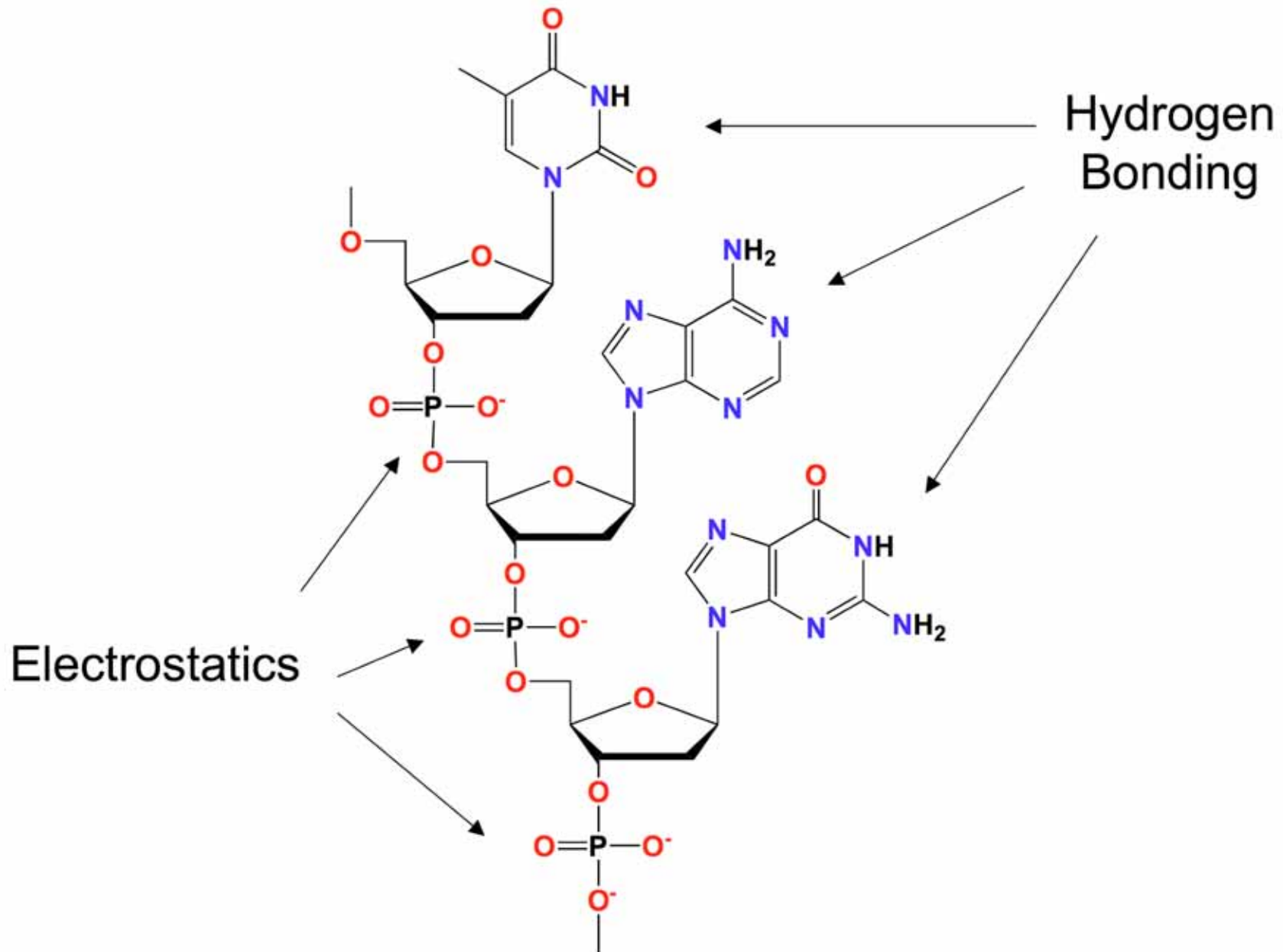
What forces are important?



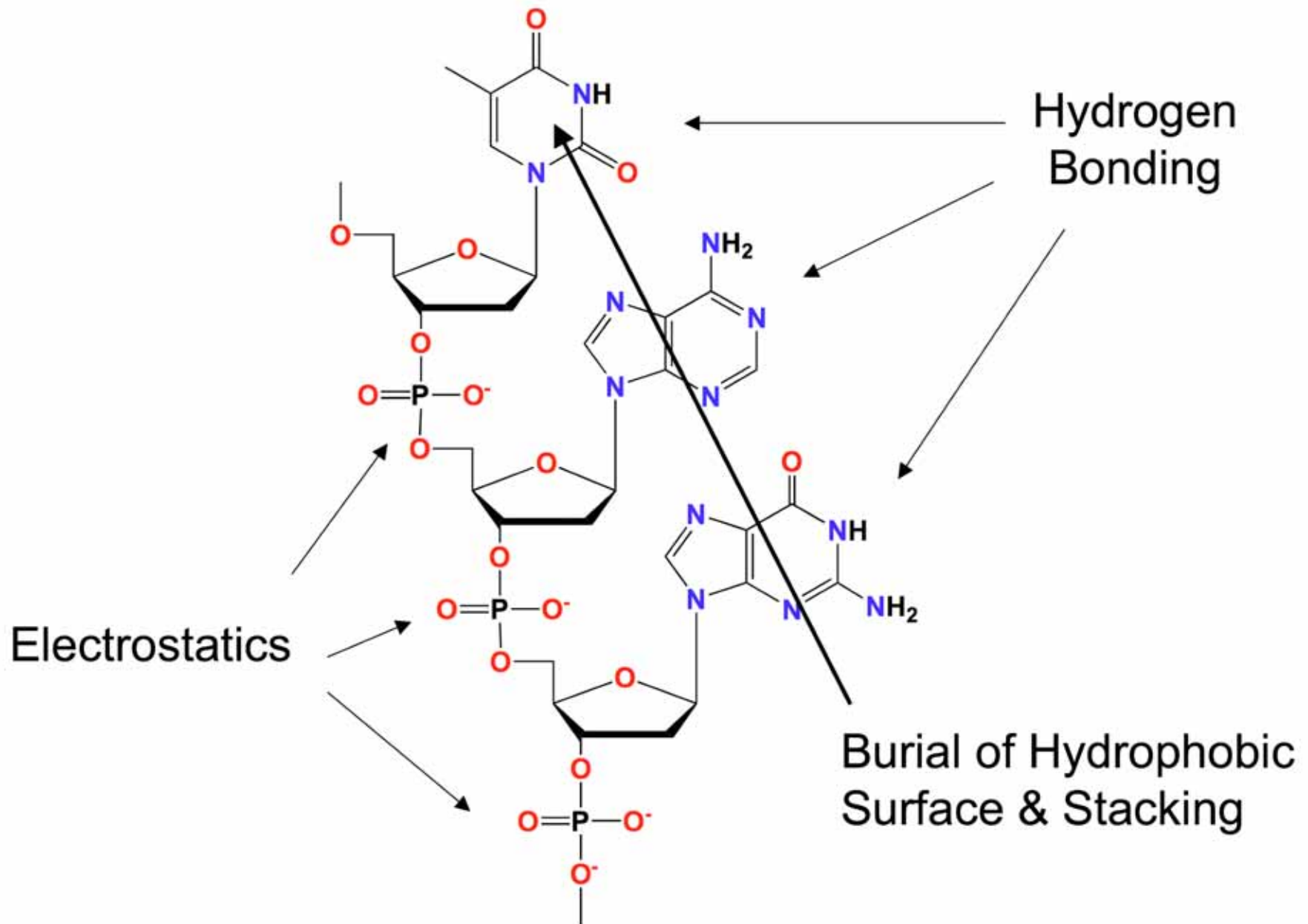
What forces are important?



What forces are important?



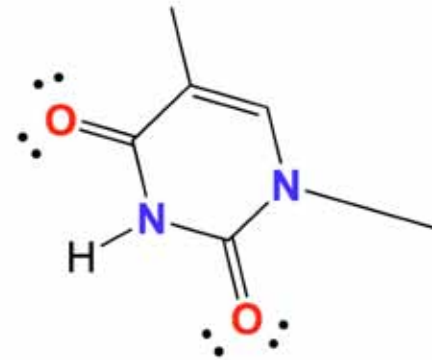
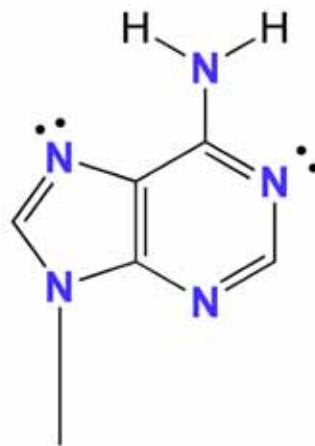
What forces are important?



Base Pairing

(Donors matched to Acceptors)

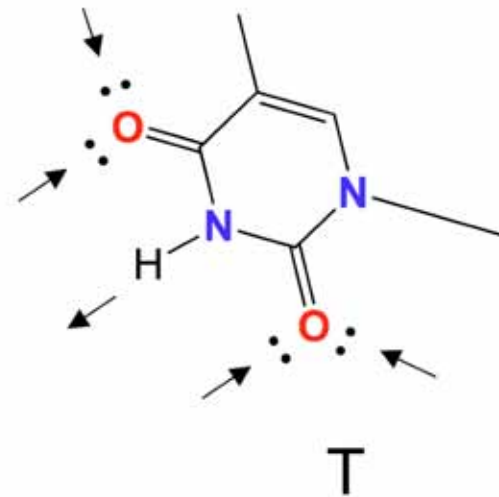
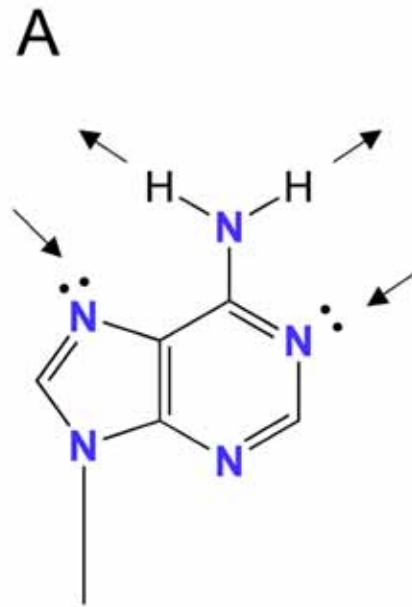
A



T

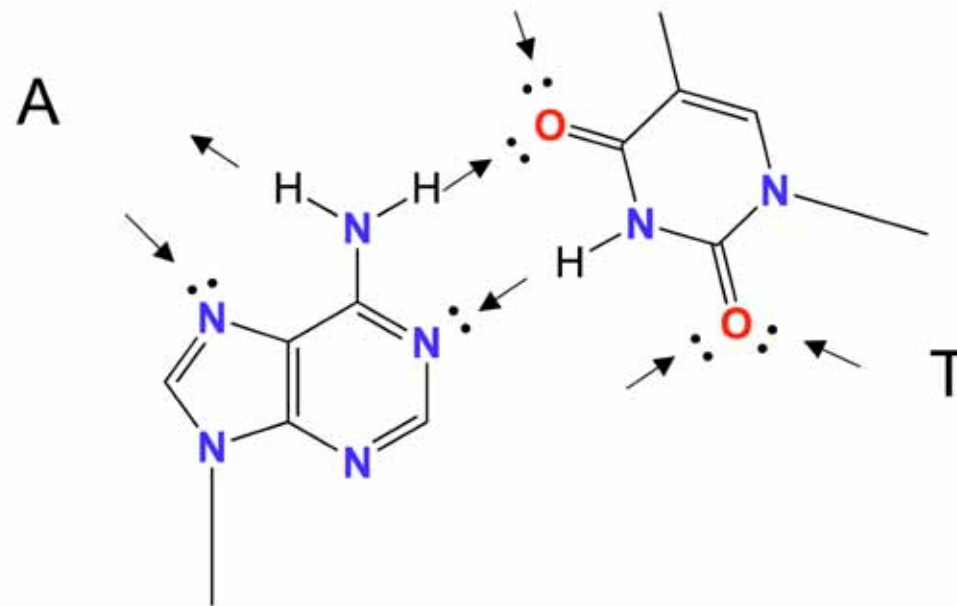
Base Pairing

(Donors matched to Acceptors)



Base Pairing

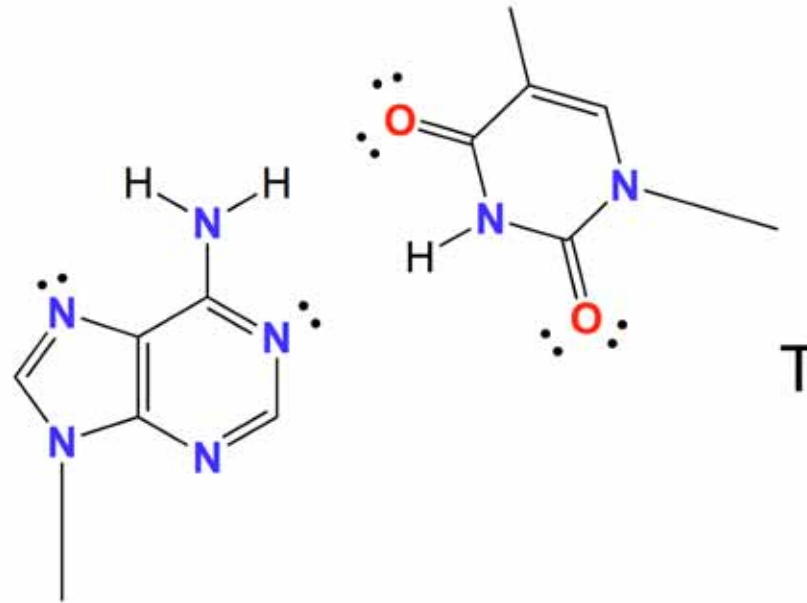
(Donors matched to Acceptors)



Base Pairing

(Donors matched to Acceptors)

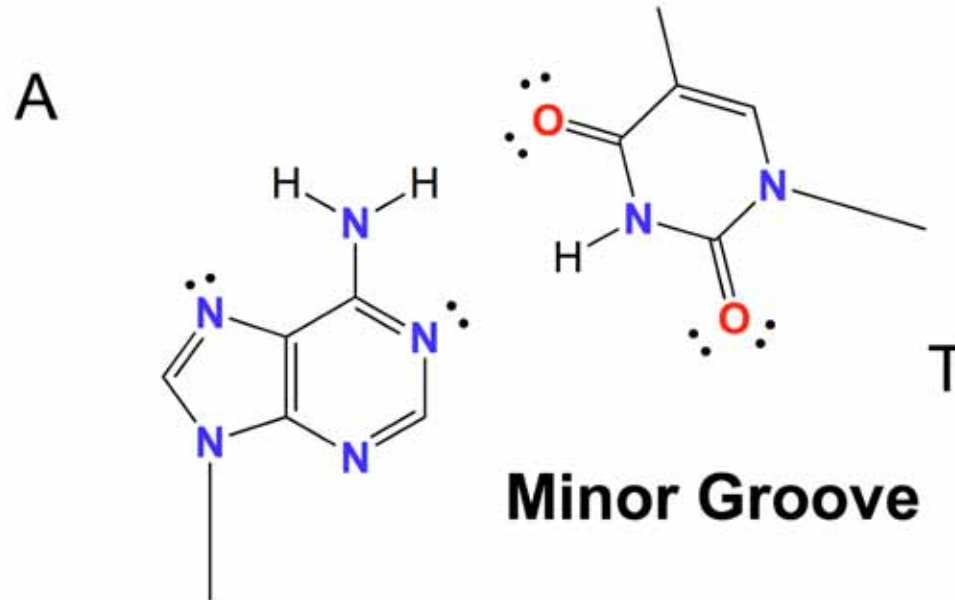
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Base Pairing

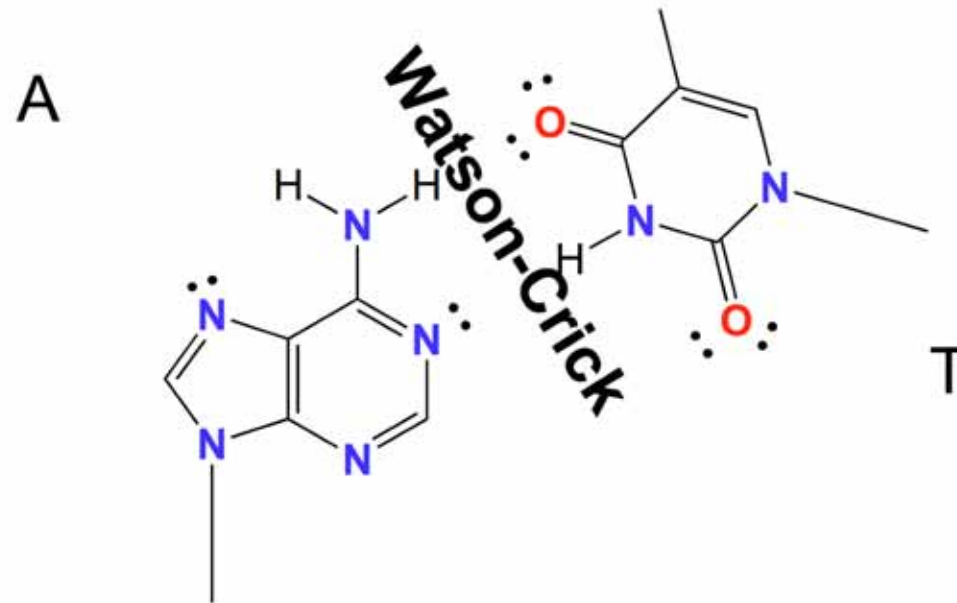
(Donors matched to Acceptors)

Major Groove



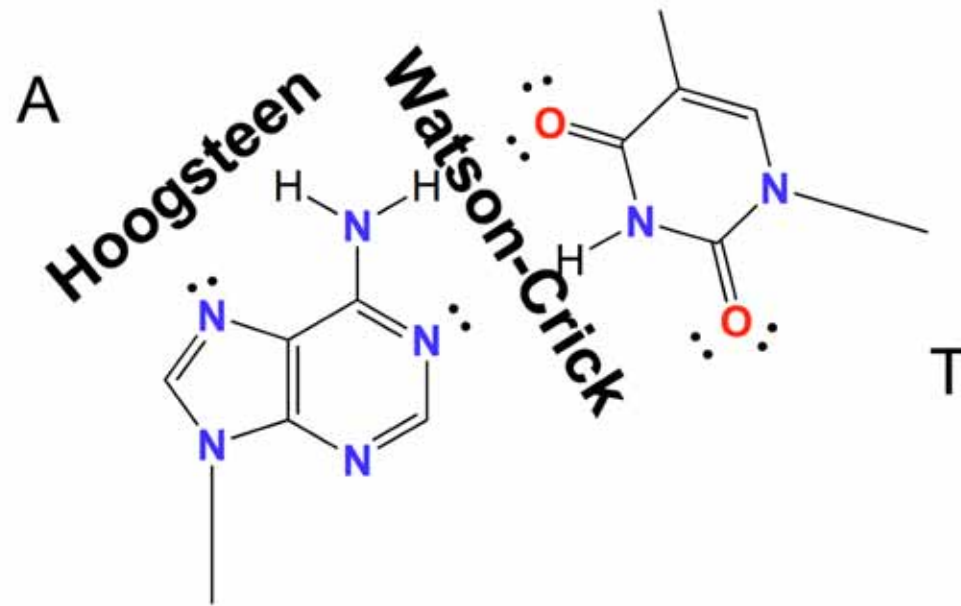
Base Pairing

(Donors matched to Acceptors)



Base Pairing

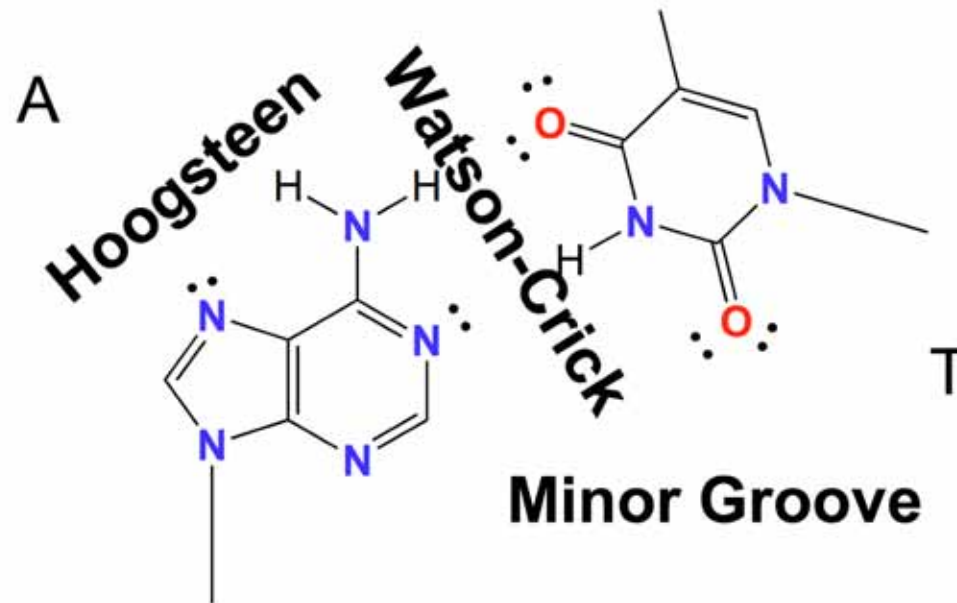
(Donors matched to Acceptors)



Base Pairing

(Donors matched to Acceptors)

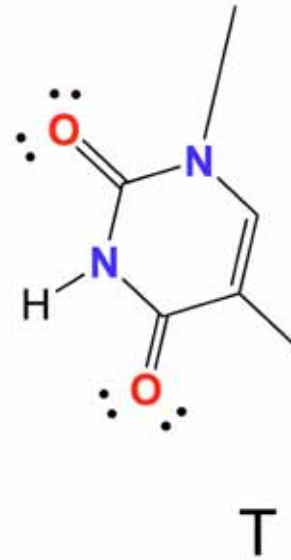
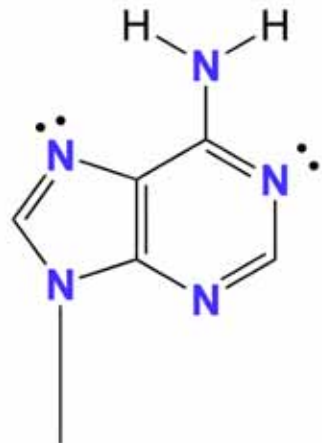
Major Groove



Base Pairing

(Donors matched to Acceptors)

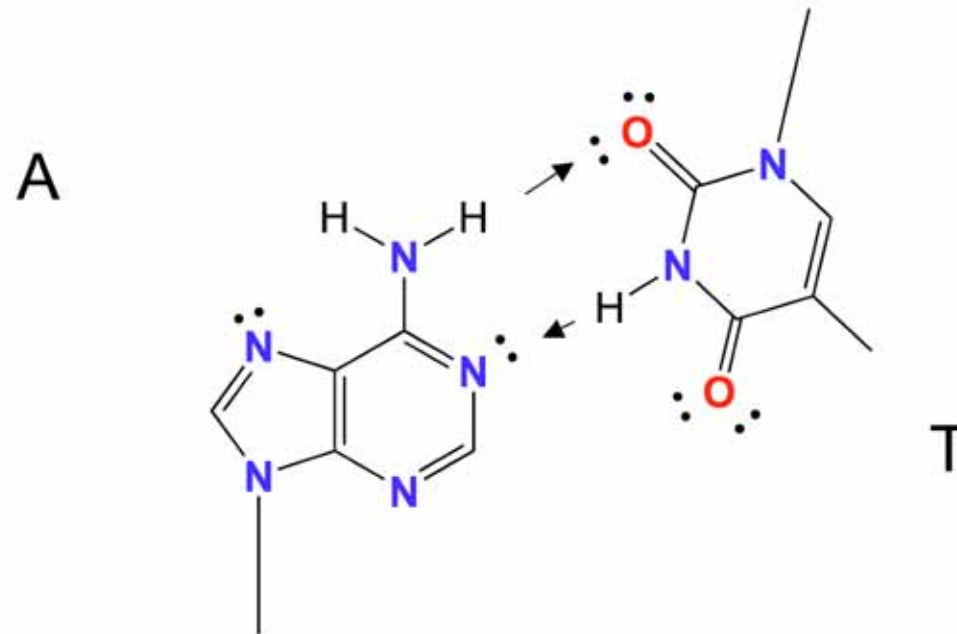
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T

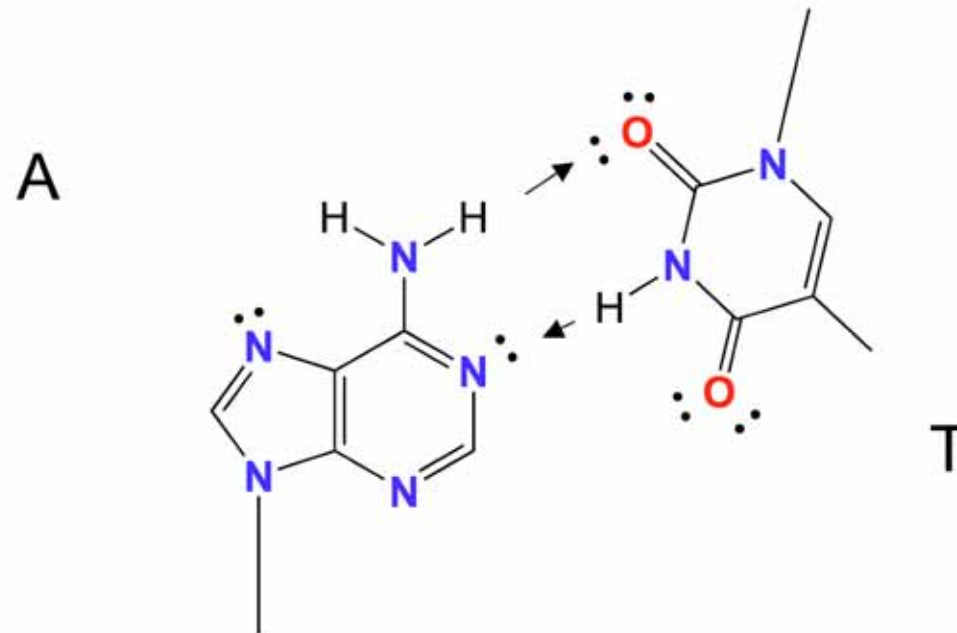
Base Pairing

(Donors matched to Acceptors)



Base Pairing

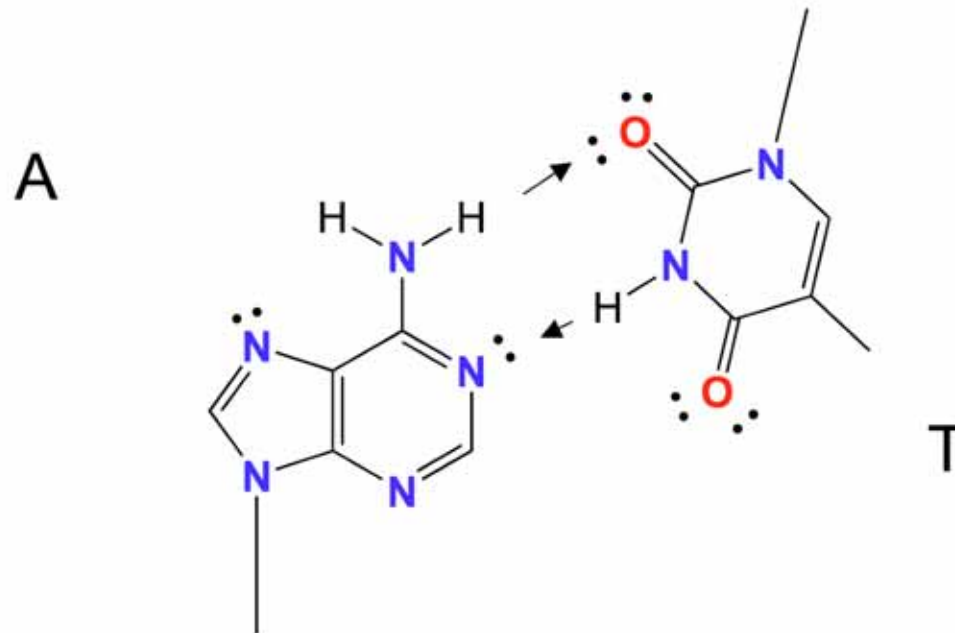
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Good base pairing

Base Pairing

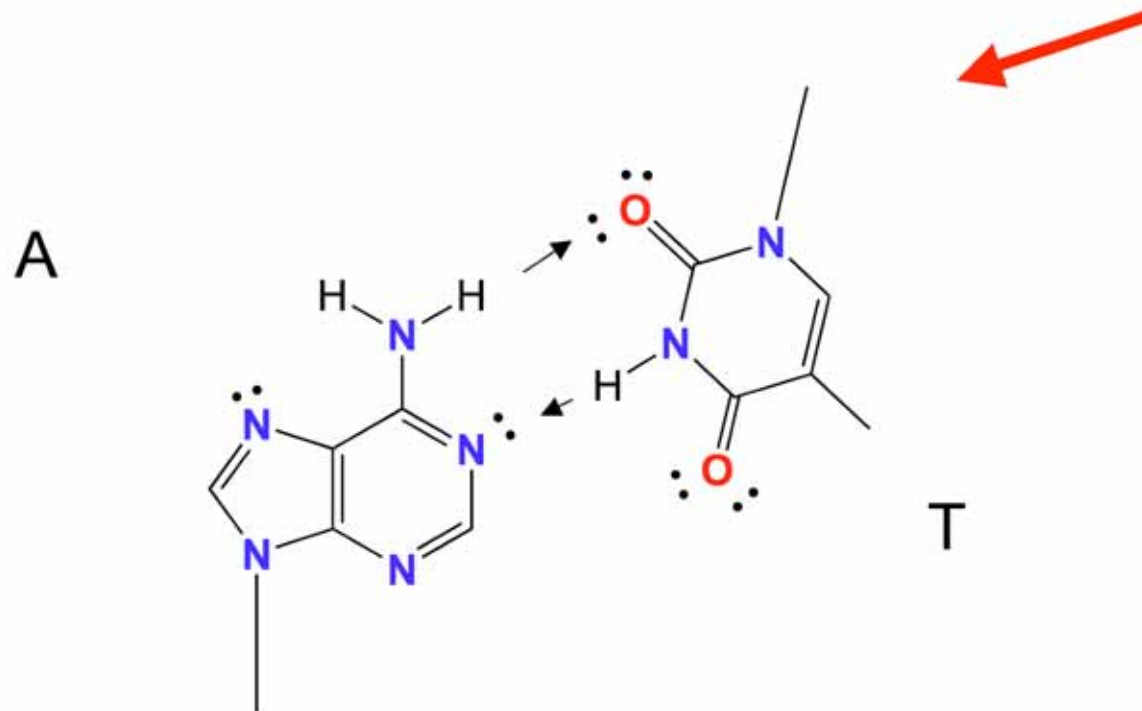
(Donors matched to Acceptors)



Good base pairing
Watson-Crick facing

Base Pairing

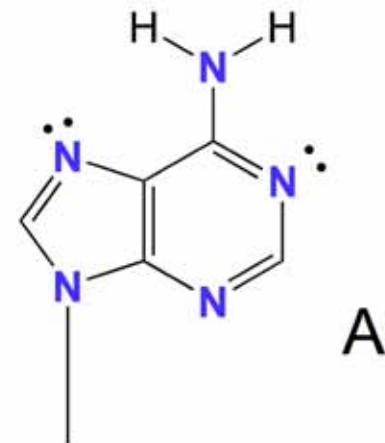
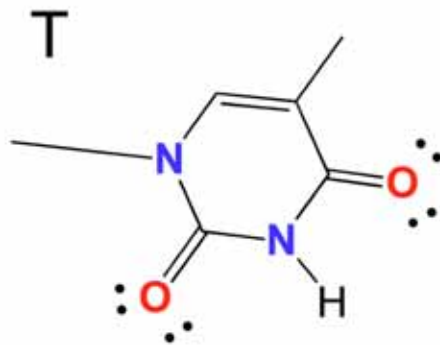
(Donors matched to Acceptors)



Good base pairing
Watson-Crick facing
but *Anti-Watson-Crick* orientation

Base Pairing

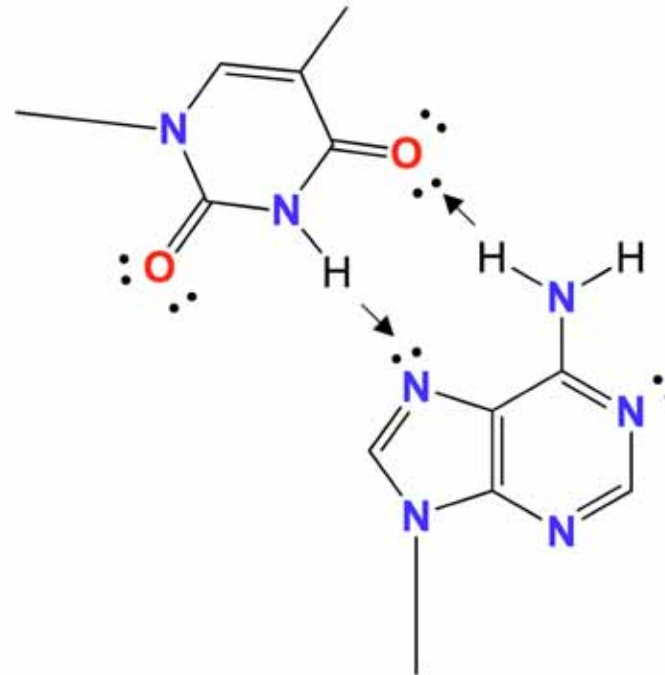
(Donors matched to Acceptors)



Base Pairing

(Donors matched to Acceptors)

T

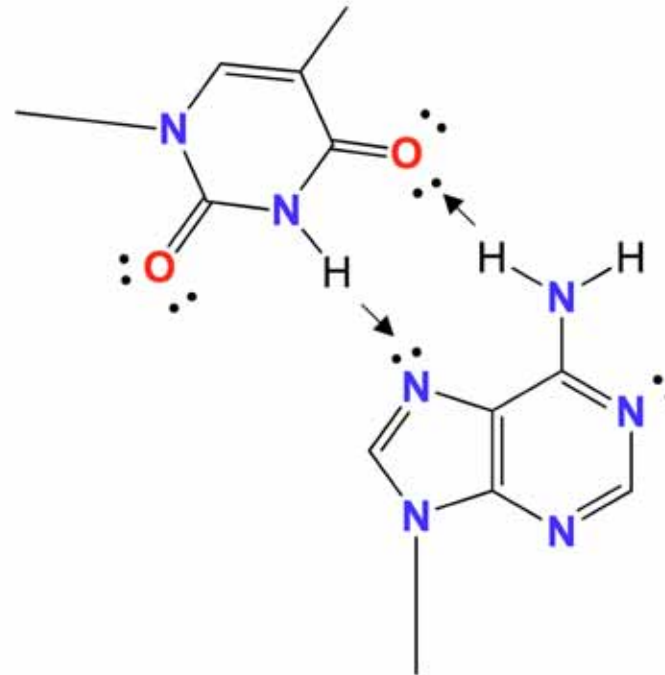


A

Base Pairing

(Donors matched to Acceptors)

T



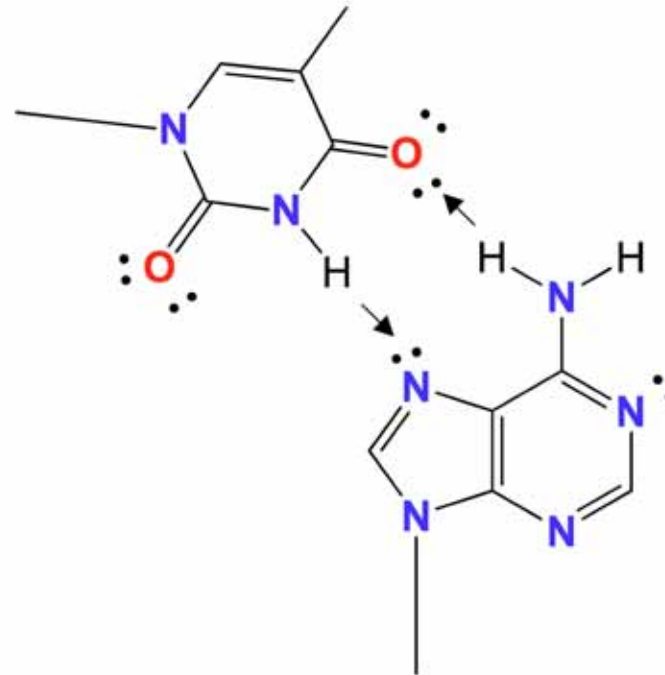
A

Good base pairing

Base Pairing

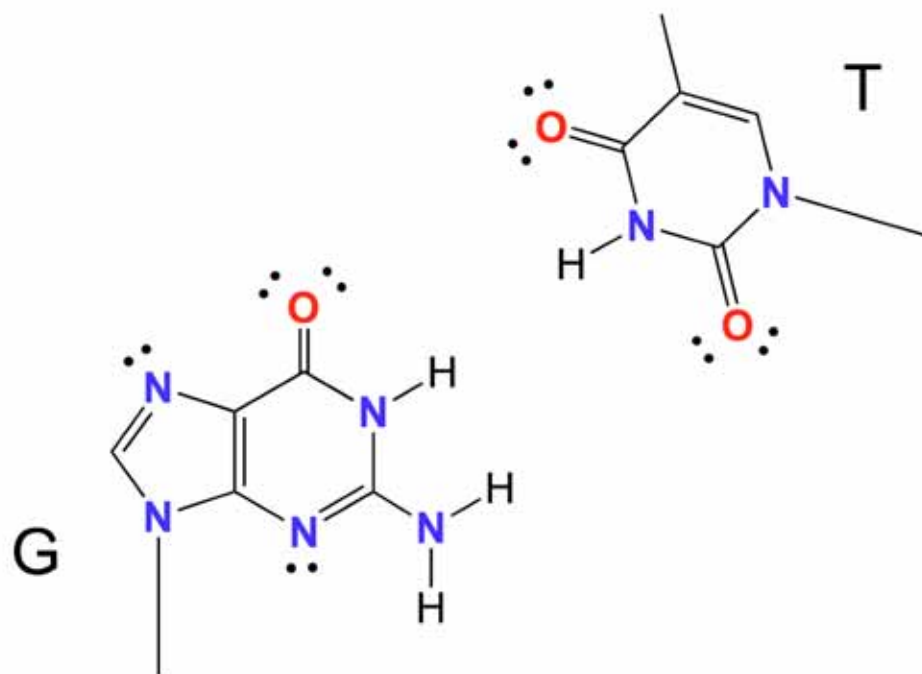
(Donors matched to Acceptors)

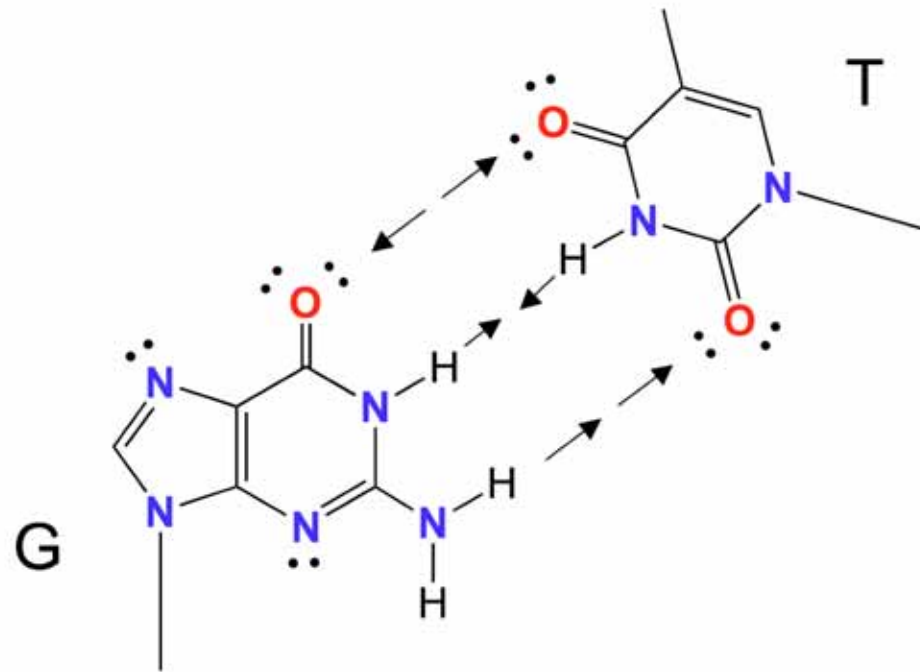
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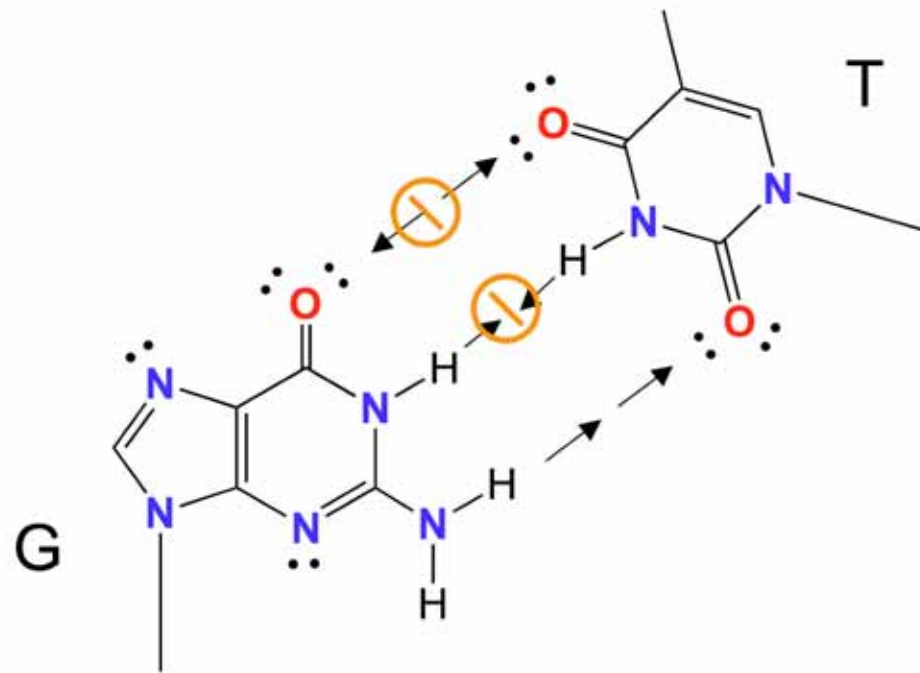


A

Good base pairing
WC-Hoogsteen facing

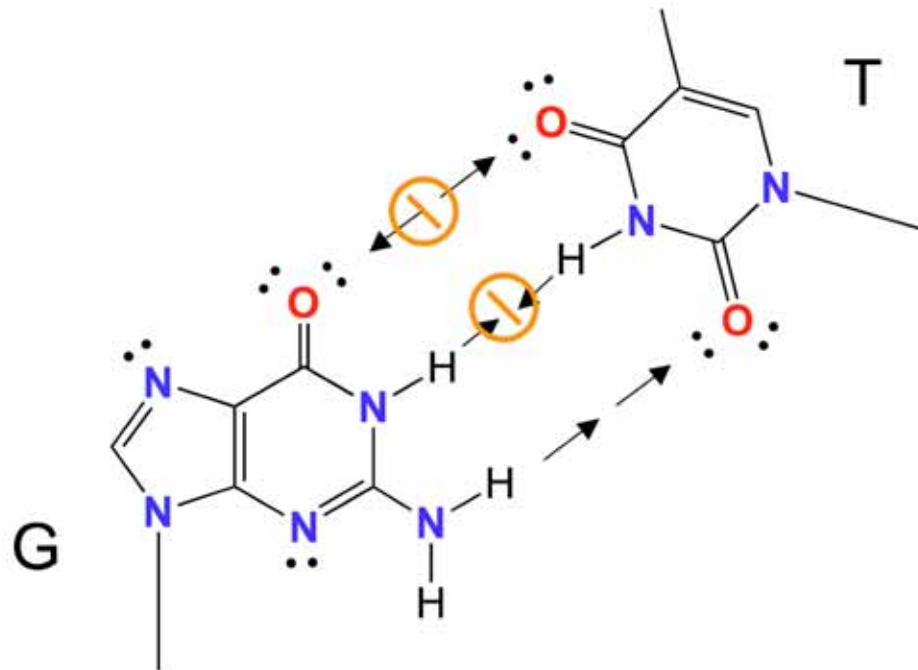


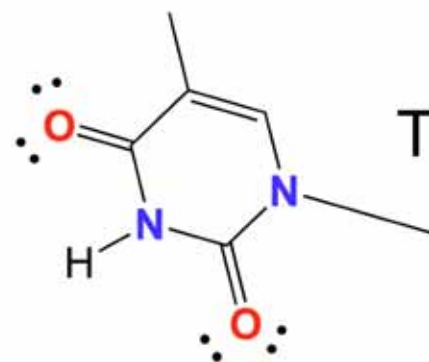
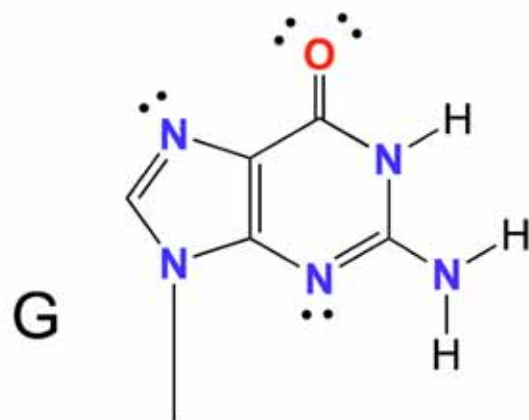


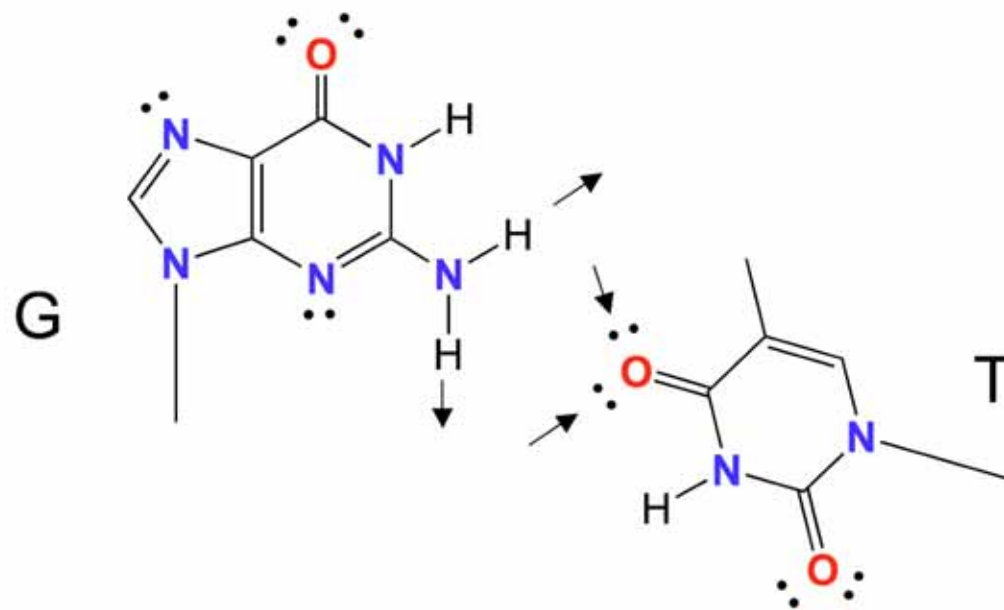


Bad Base Pairing

(Donors **not** matched to Acceptors)

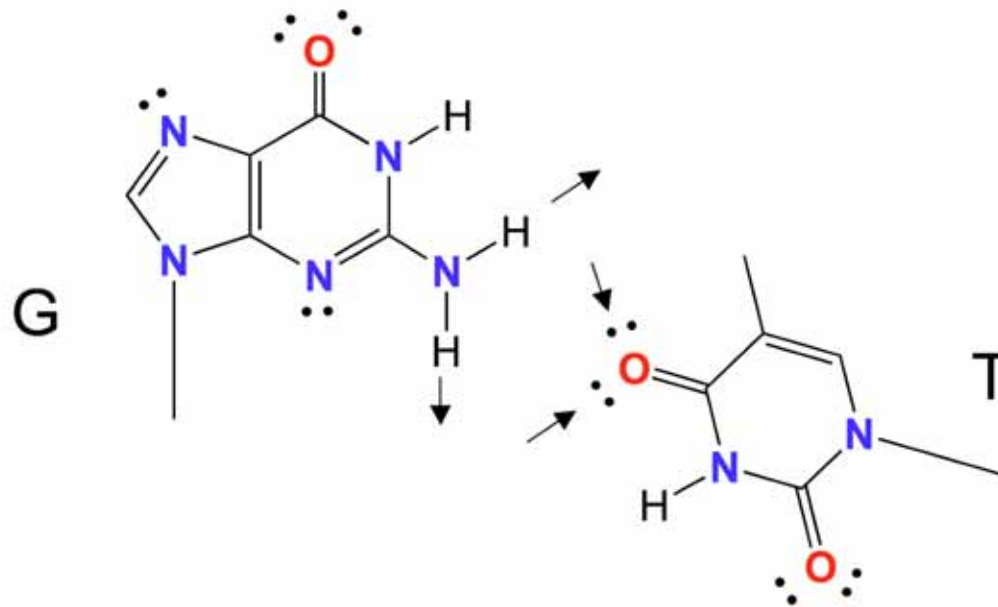




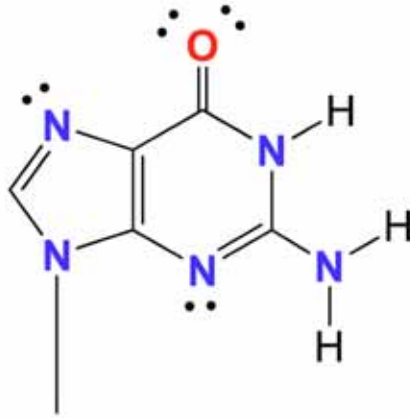


Bad Base Pairing

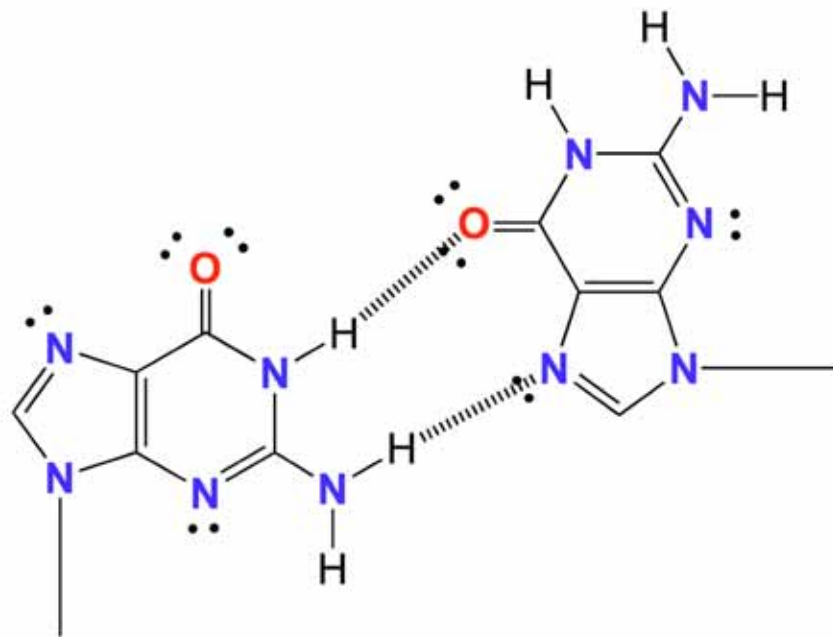
(Donors to Acceptors with *terrible angles*)



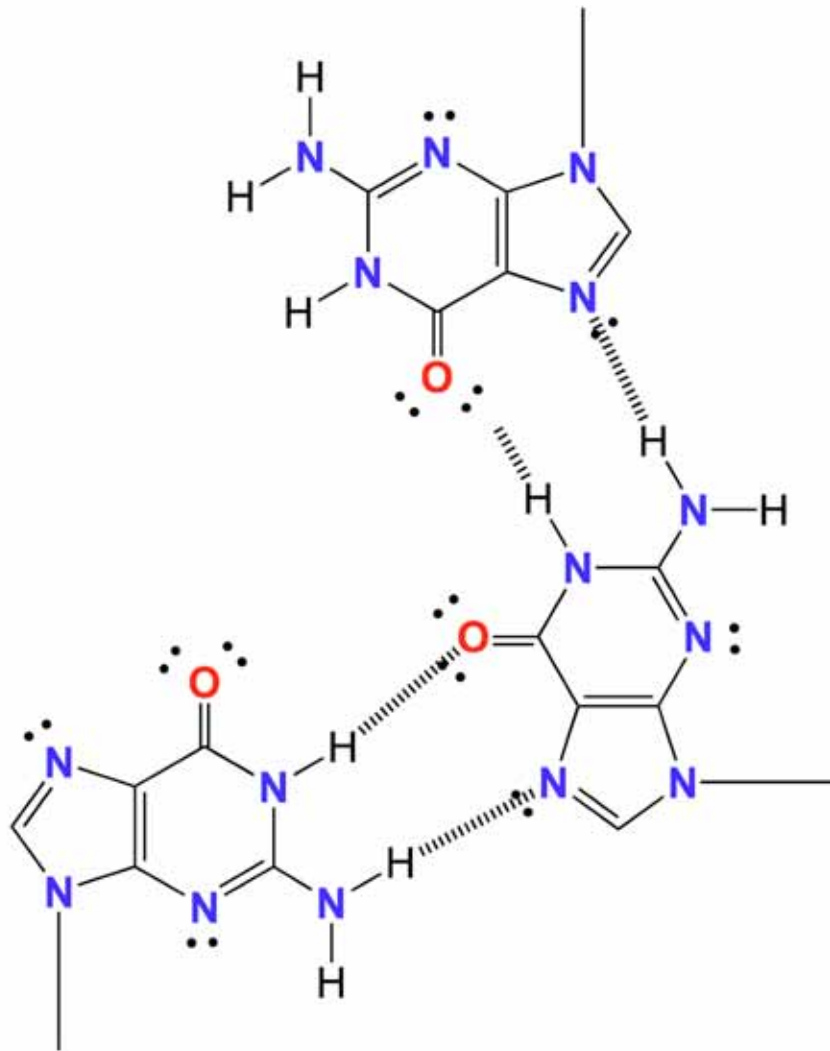
Wild (but good) Base Pairing



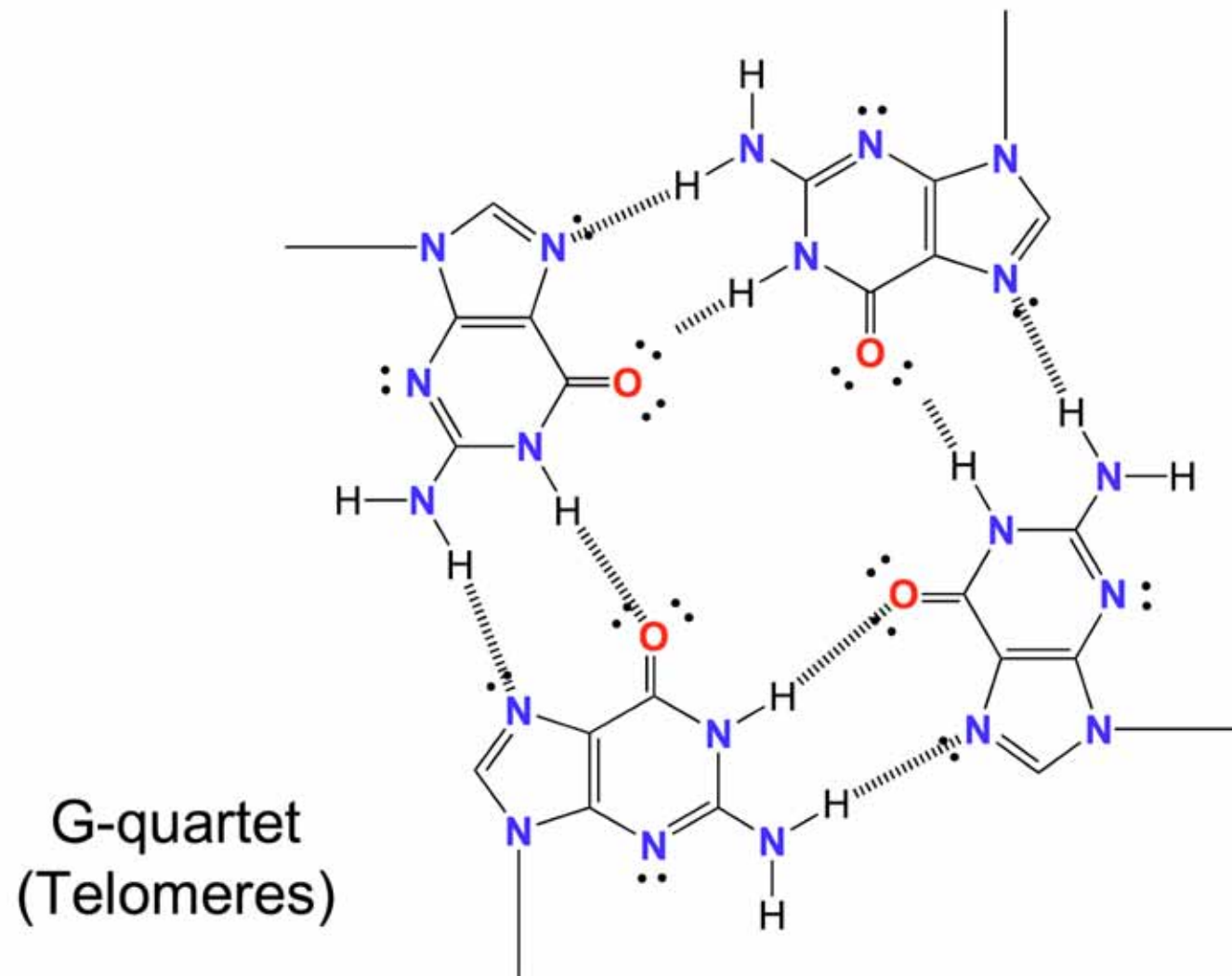
Wild (but good) Base Pairing



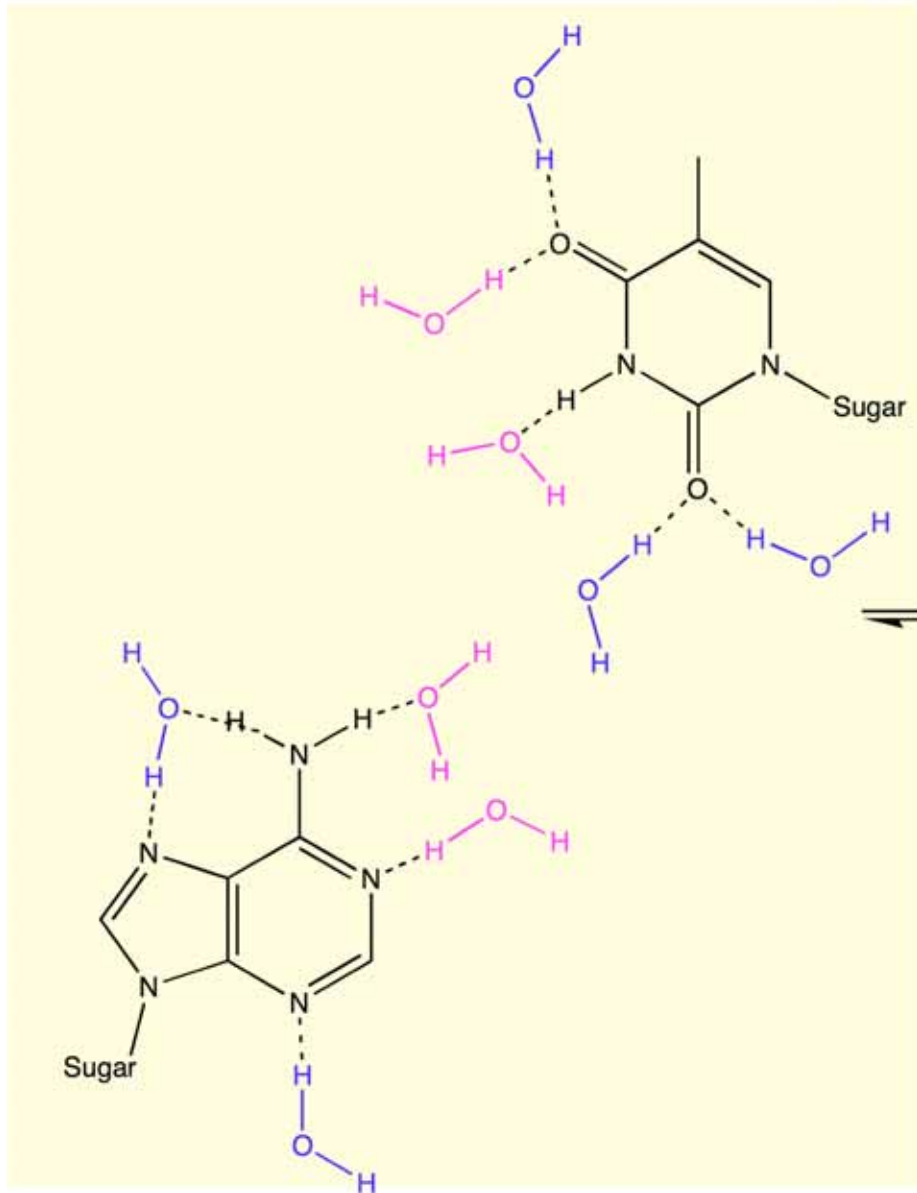
Wild (but good) Base Pairing



Wild (but good) Base Pairing

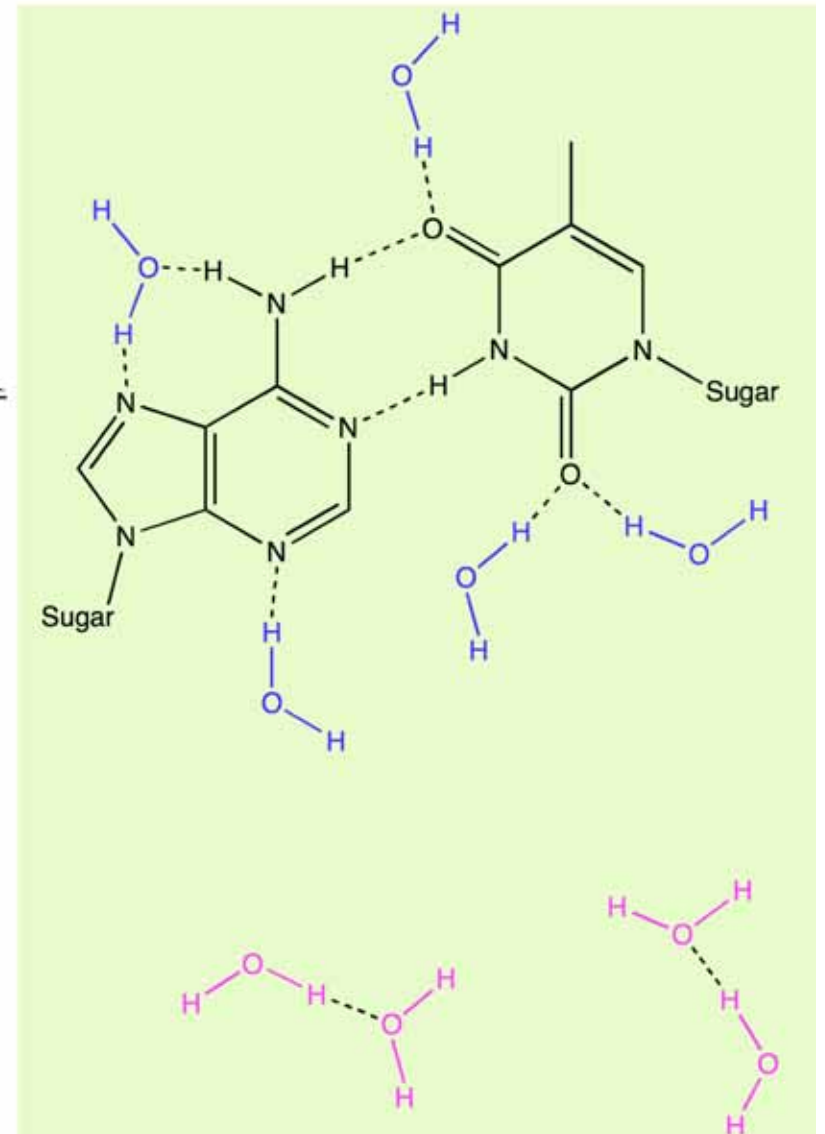


AT Base Pair

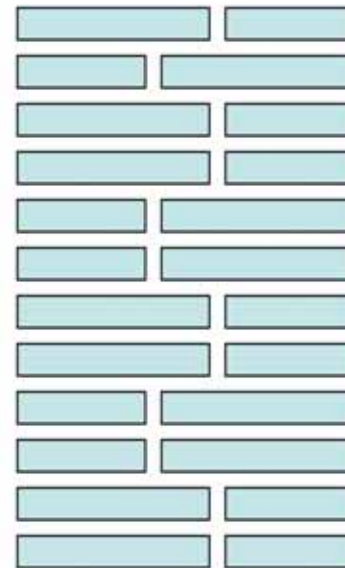
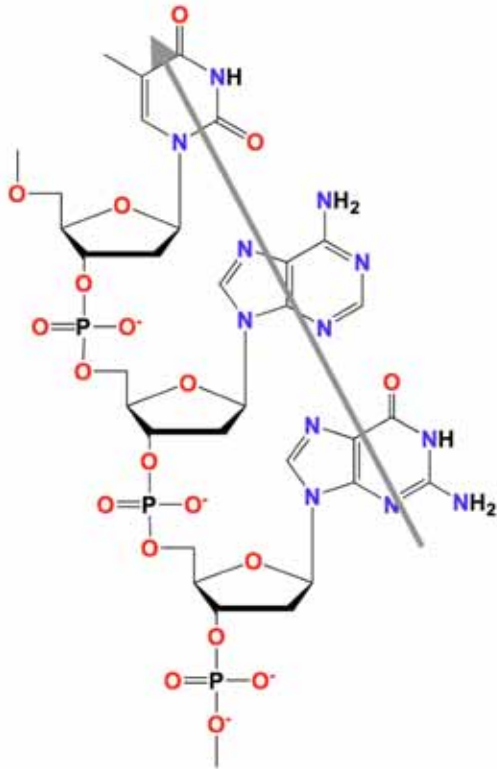


Ten H-Bonds

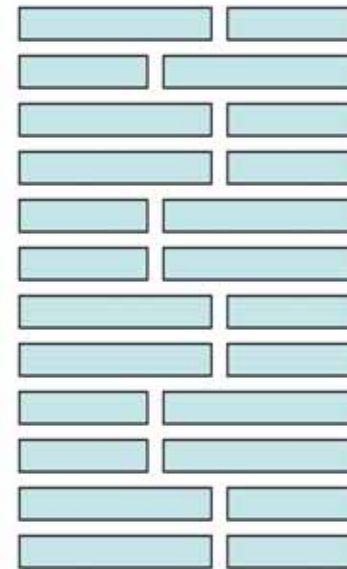
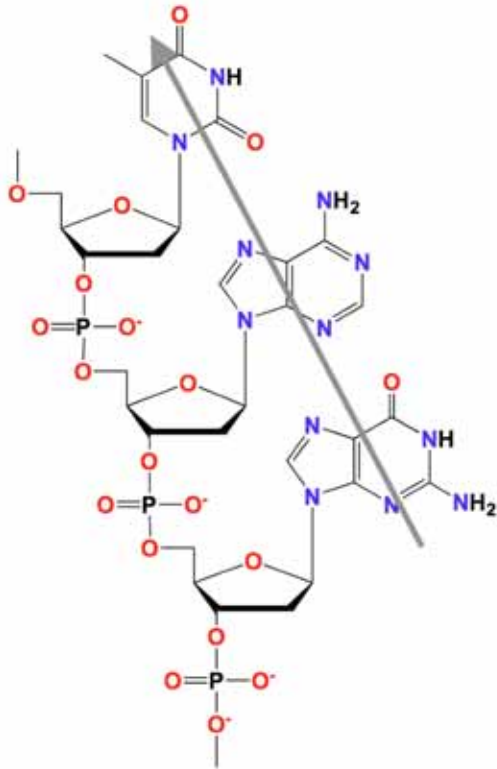
Ten H-Bonds



Burial of hydrophobic surface drives helix formation
(hydrophobic core / stacking interactions)

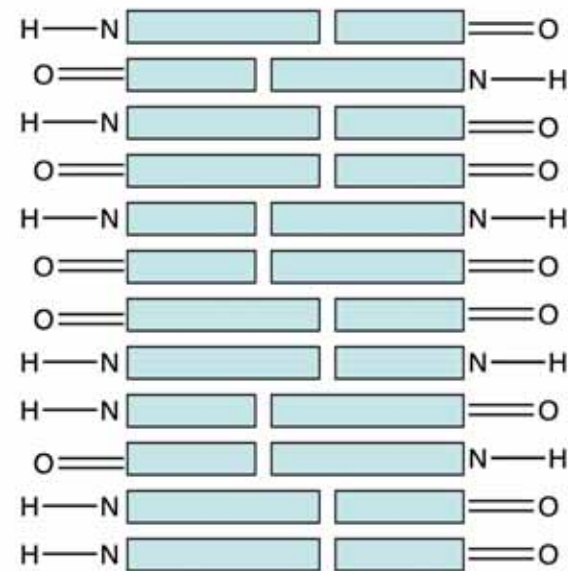
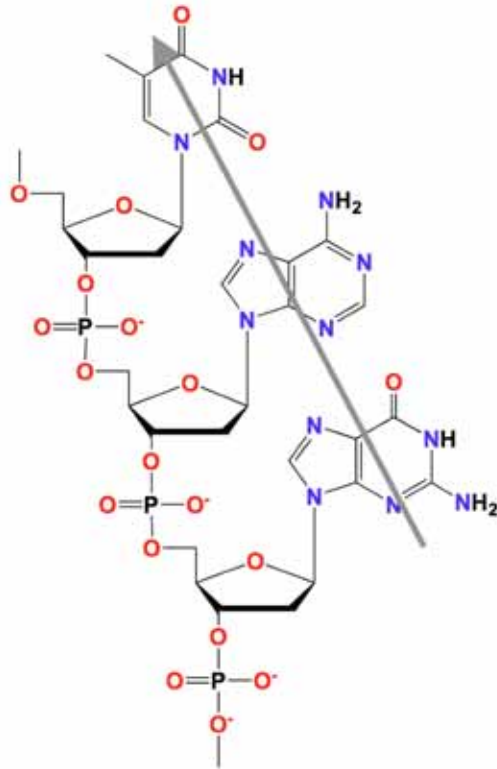


Burial of hydrophobic surface drives helix formation
(hydrophobic core / stacking interactions)



Flat faces are nonpolar

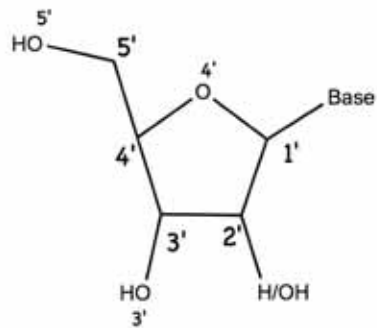
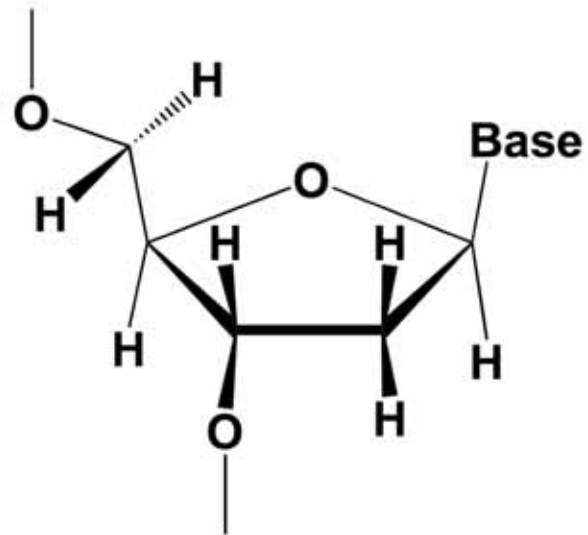
Burial of hydrophobic surface drives helix formation (hydrophobic core / stacking interactions)



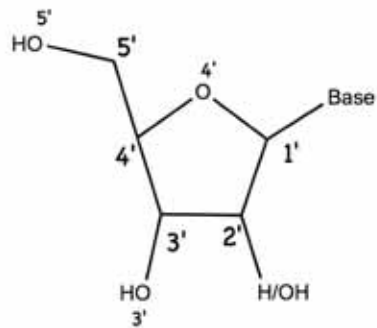
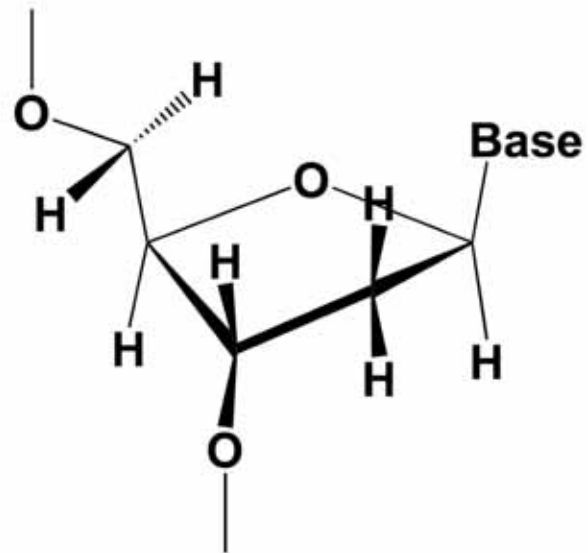
Flat faces are nonpolar
Edges are very polar (can H-bond)

Other chemical constraints

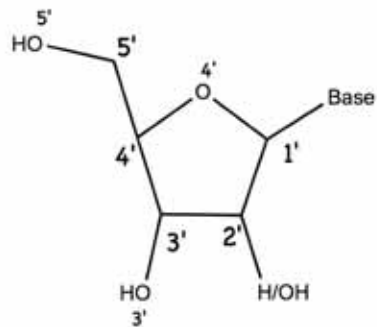
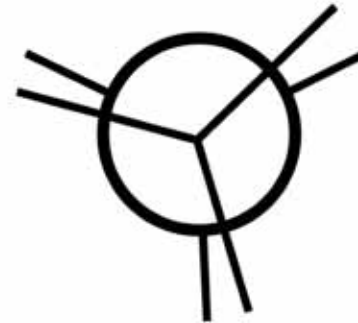
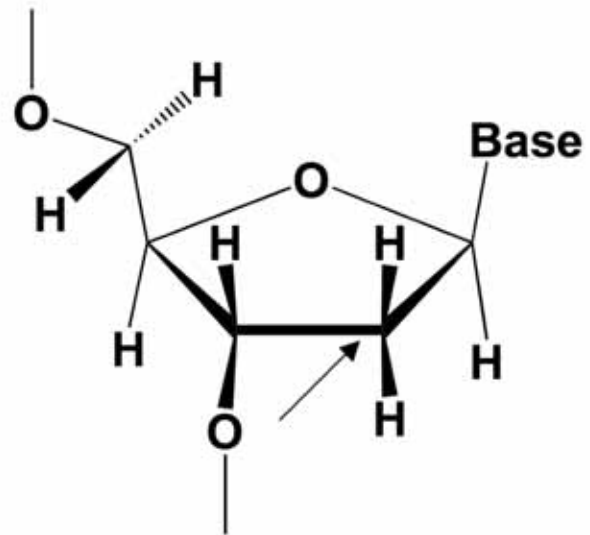
Furanose Sugar Ring



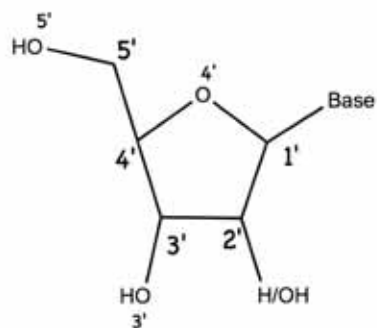
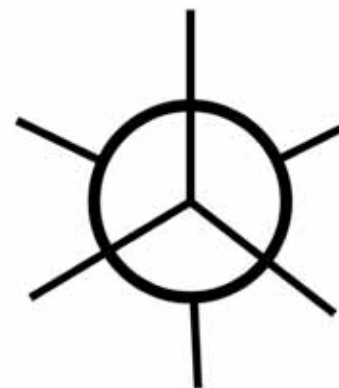
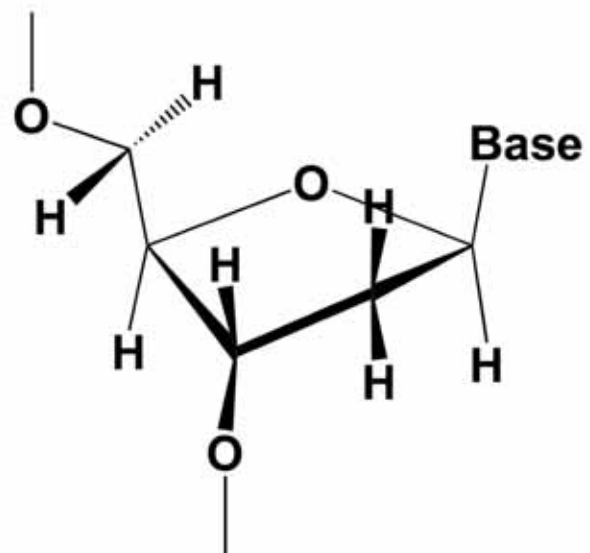
Furanose Sugar Ring



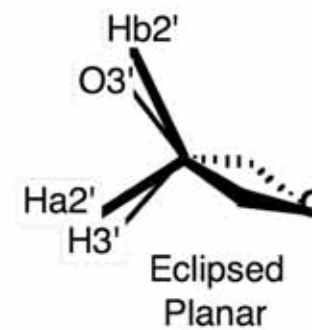
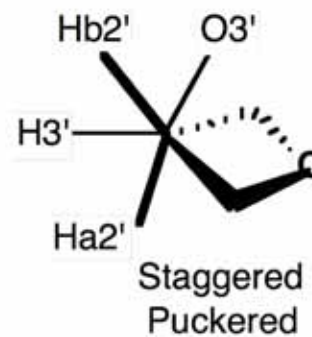
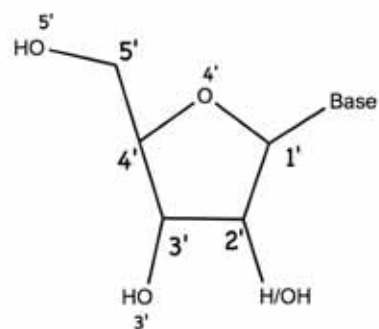
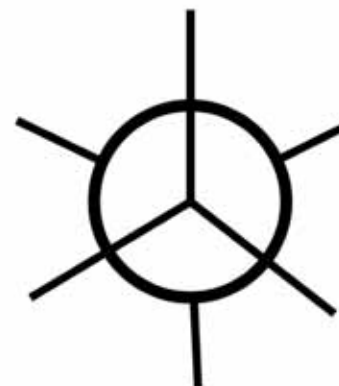
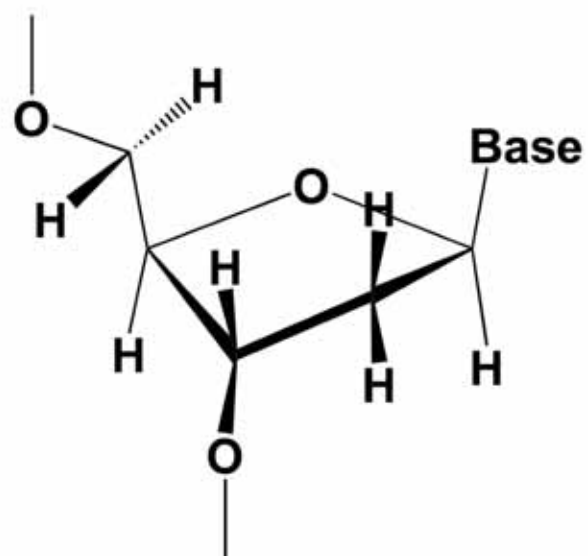
Furanose Sugar Ring



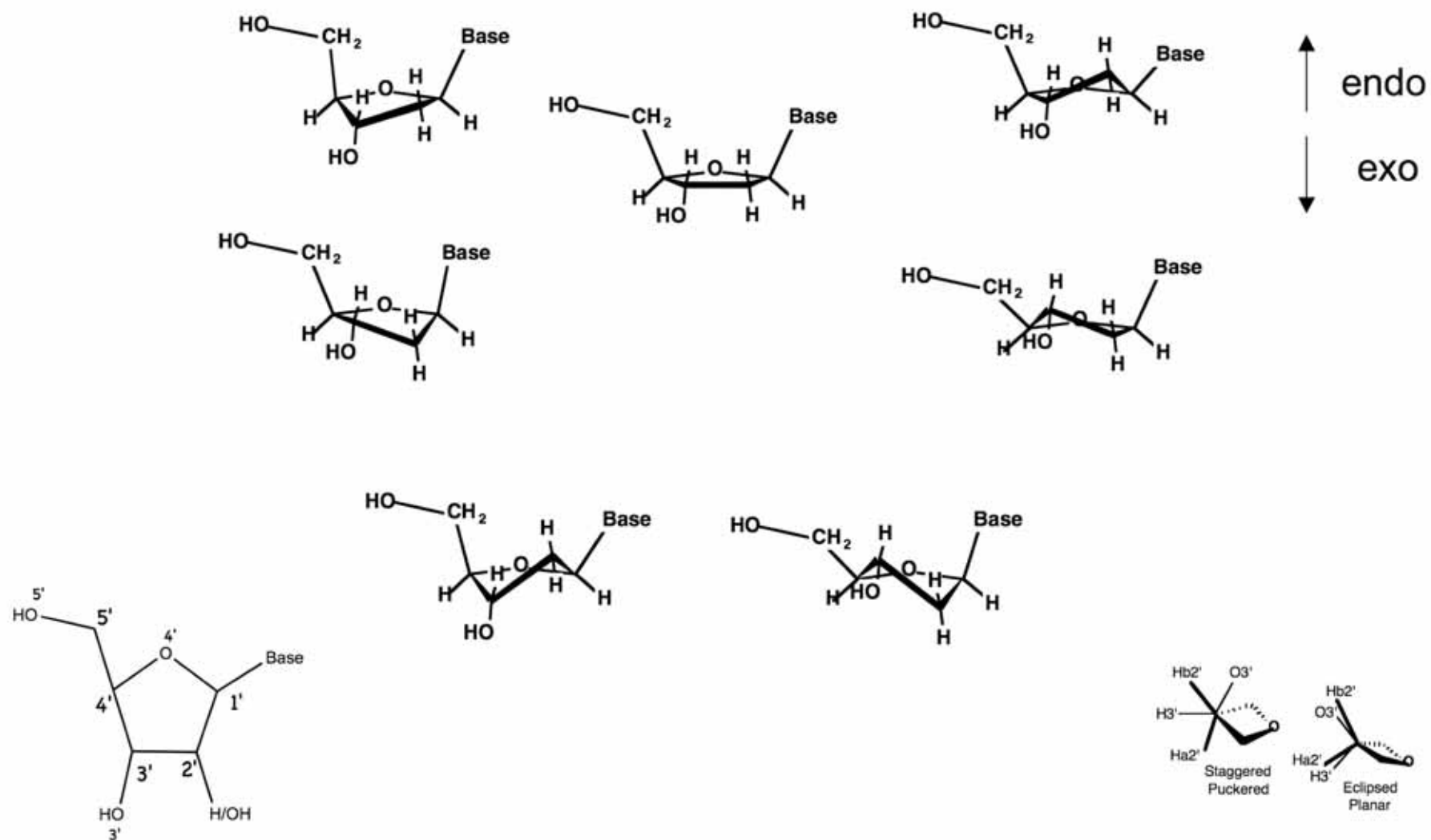
Furanose Sugar Ring



Furanose Sugar Ring



Furanose Sugar Ring



Replication by human DNA polymerase- ϵ occurs by Hoogsteen base-pairing

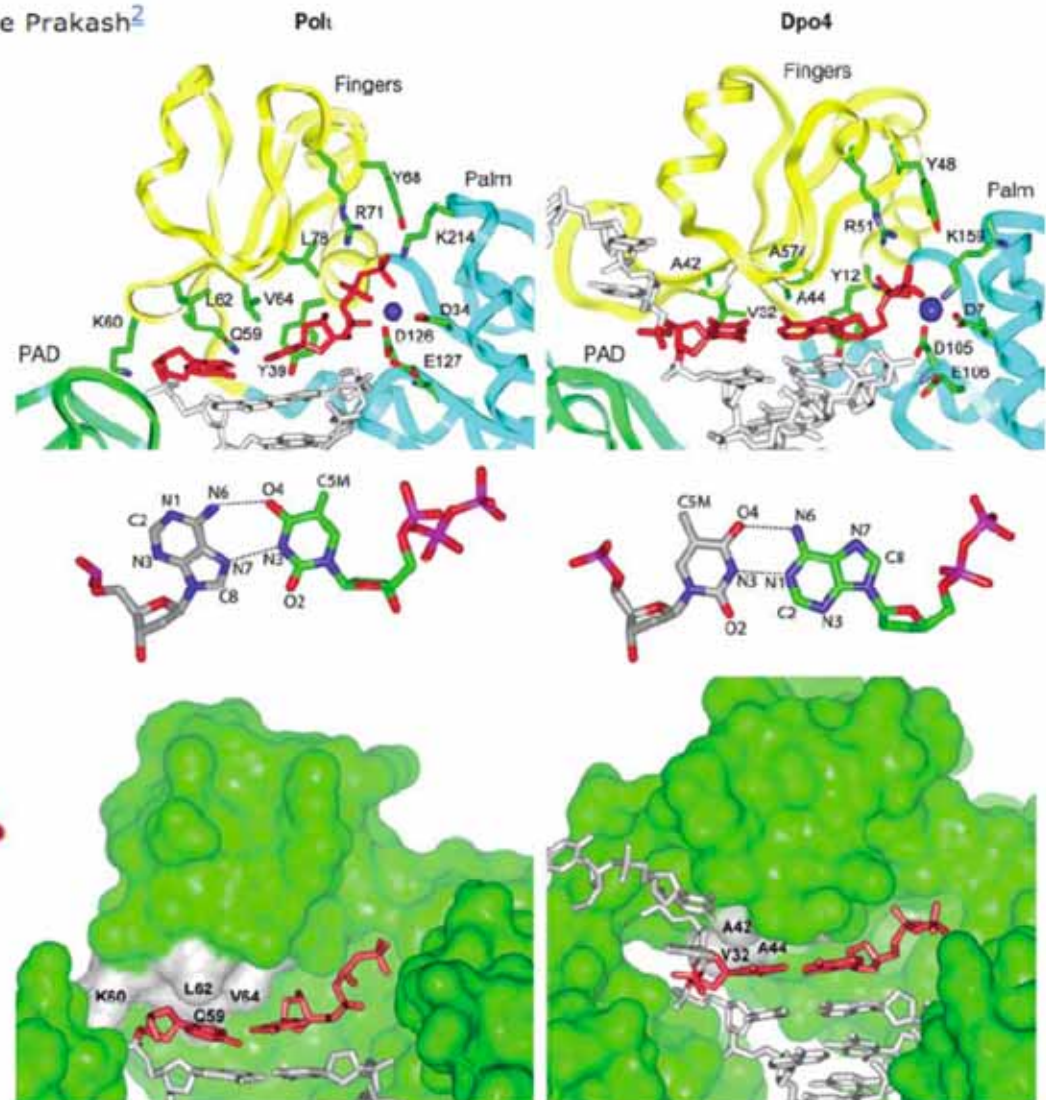
Deepak T. Nair¹, Robert E. Johnson², Satya Prakash², Louise Prakash² and Aneel K. Aggarwal¹

A crystal structure from July 2004 shows that an error correcting (and error-prone) DNA polymerase uses Hoogsteen-WC base pairing to recognize the incoming substrate dNTP.

This supports biochemical studies that had been suggesting such a pairing.

This polymerase is good at bypassing lesions in the DNA (eg, damaged Watson-Crick face of G).

Hoogsteen vindicated



Letters to Nature

Nature **430**, 377-380 (15 July 2004) | doi: 10.1038/nature02692

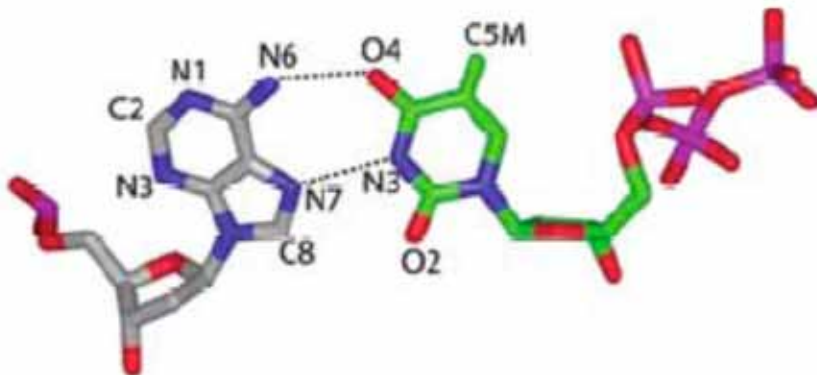
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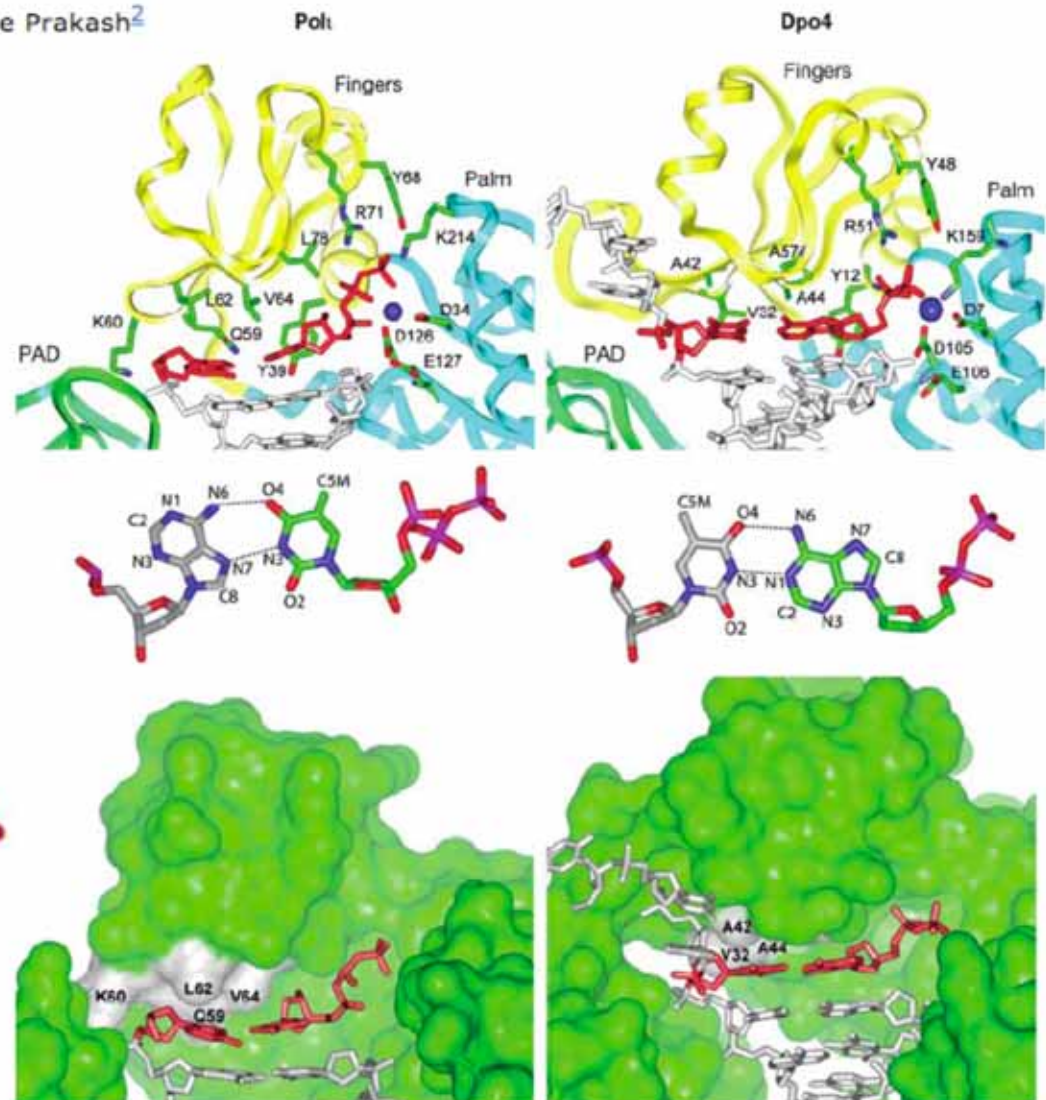
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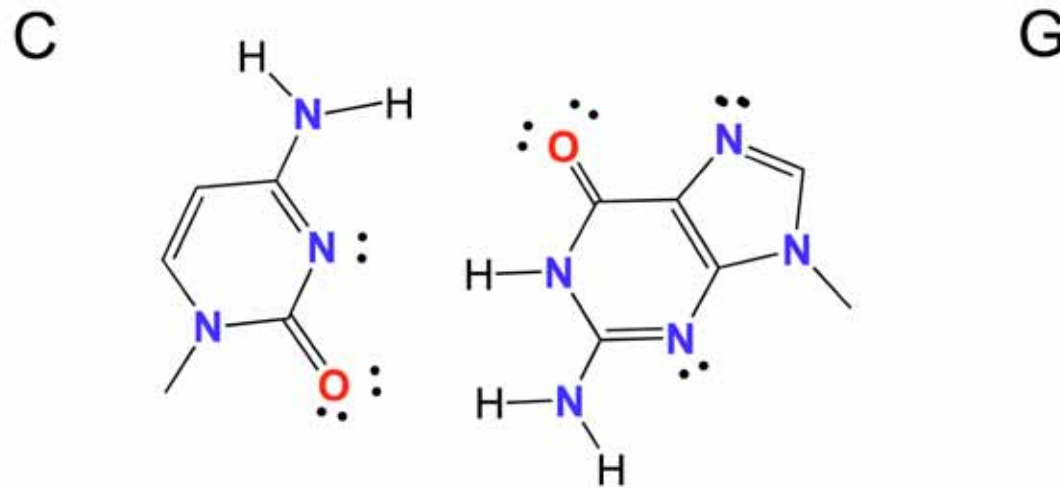


Hoogsteen vindicated

But... perhaps not...

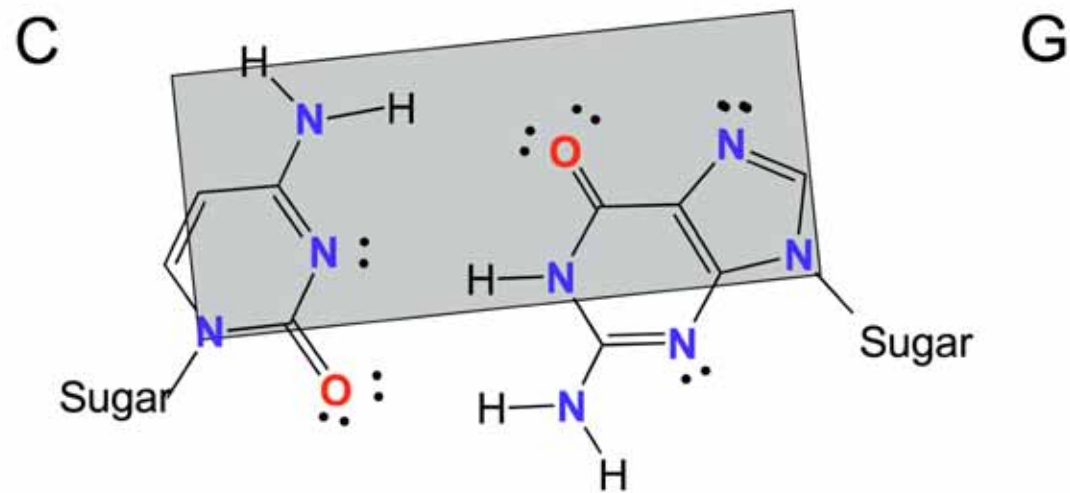


Why is Watson-Crick so good?



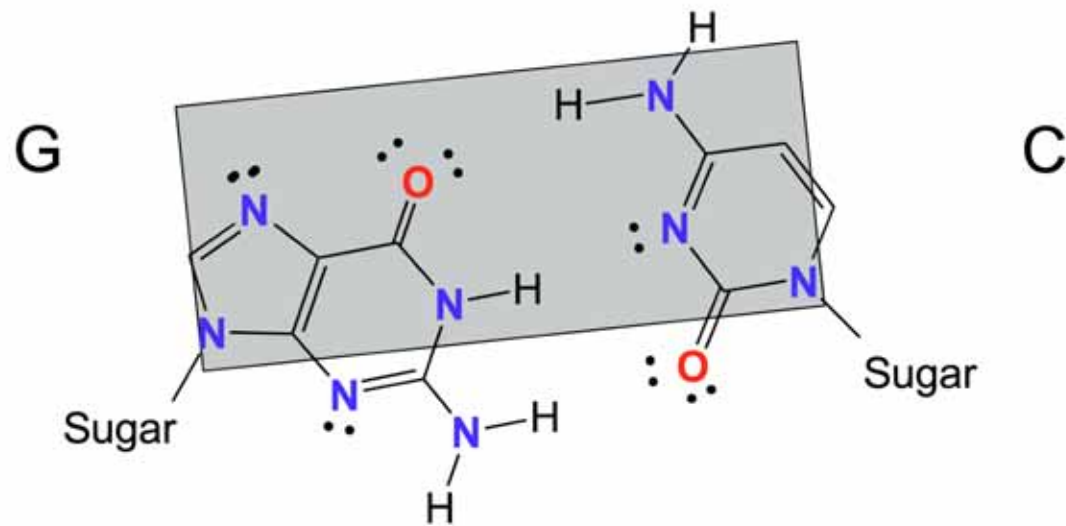
All four WC base pairs
are
isosteric

Why is Watson-Crick so good?



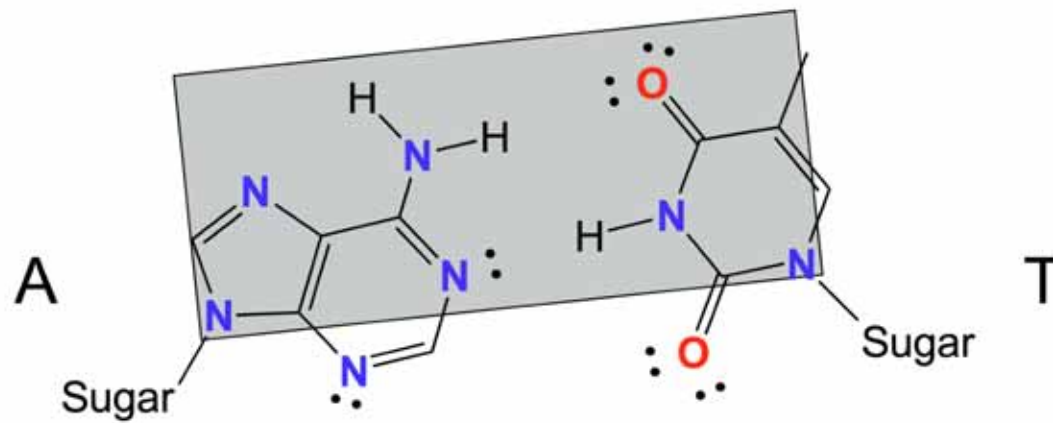
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Why is Watson-Crick so good?



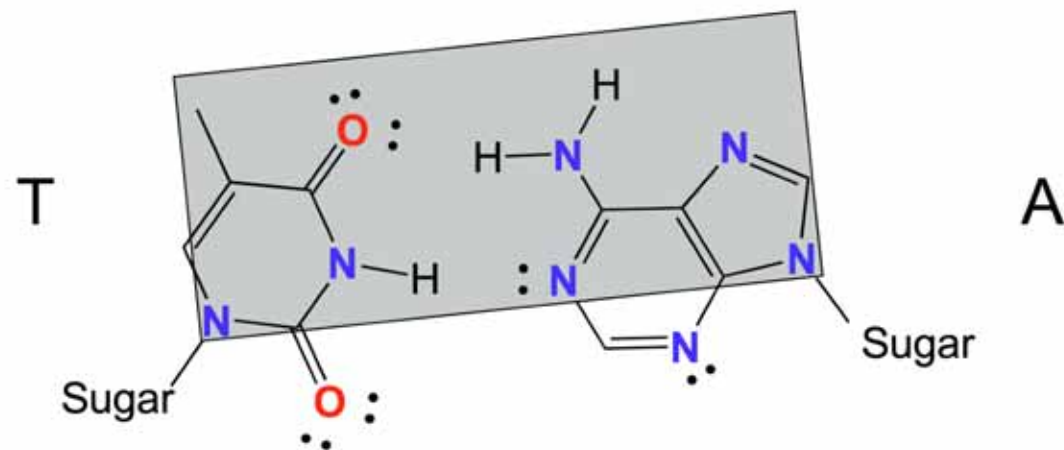
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Why is Watson-Crick so good?



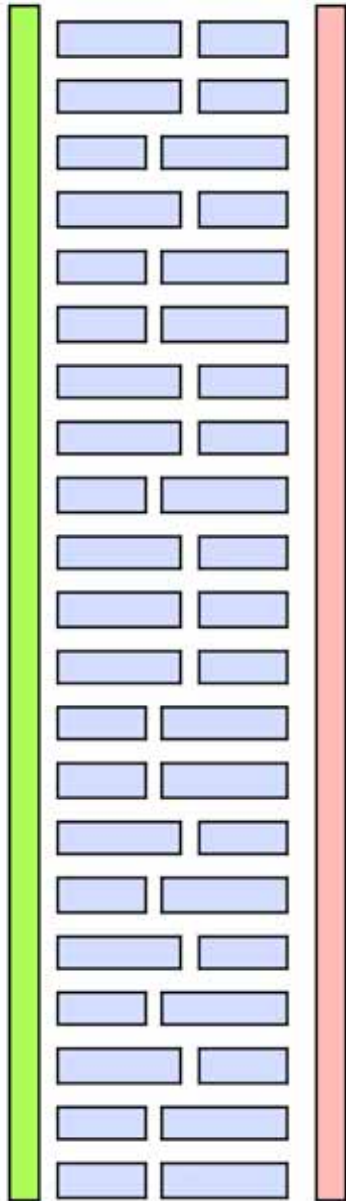
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Why is Watson-Crick so good?

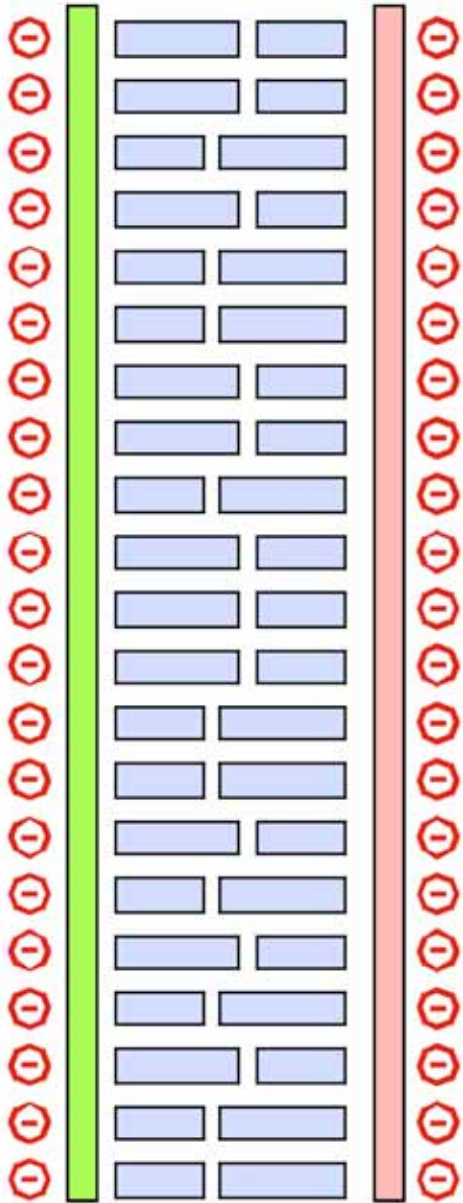


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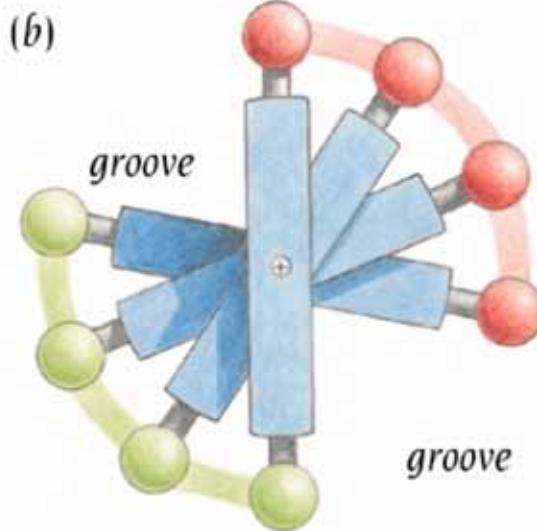
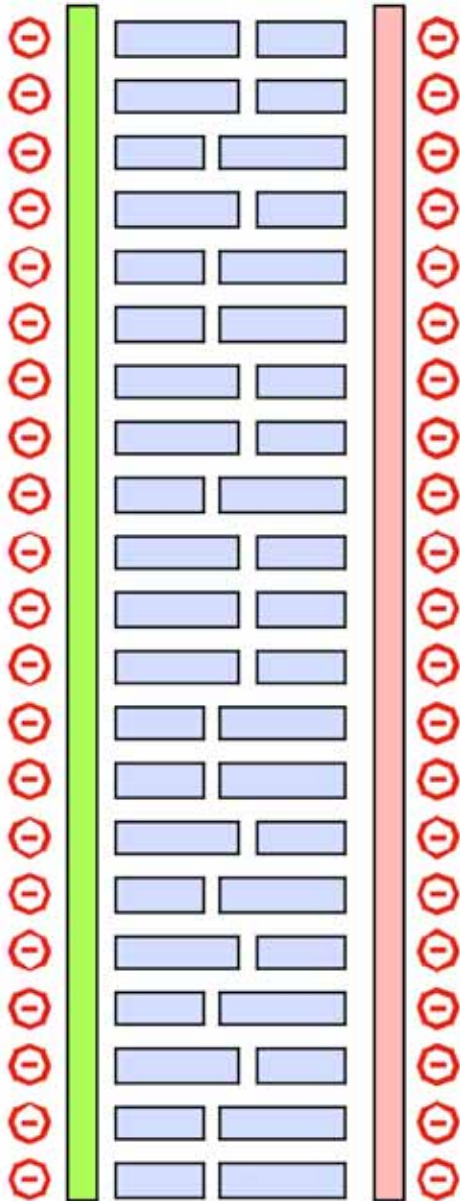
Why *a helix*?



Why *a helix*?

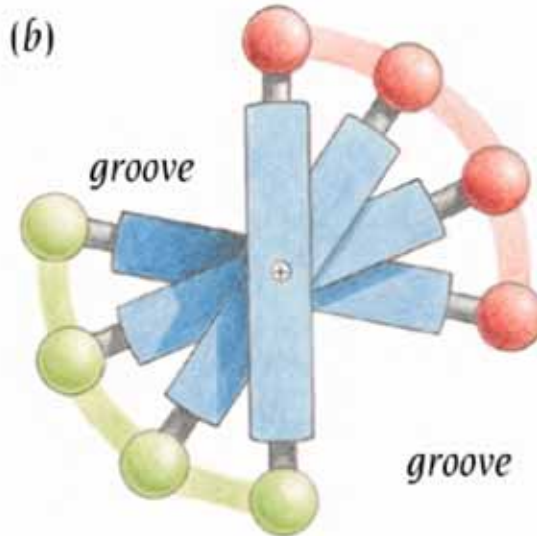
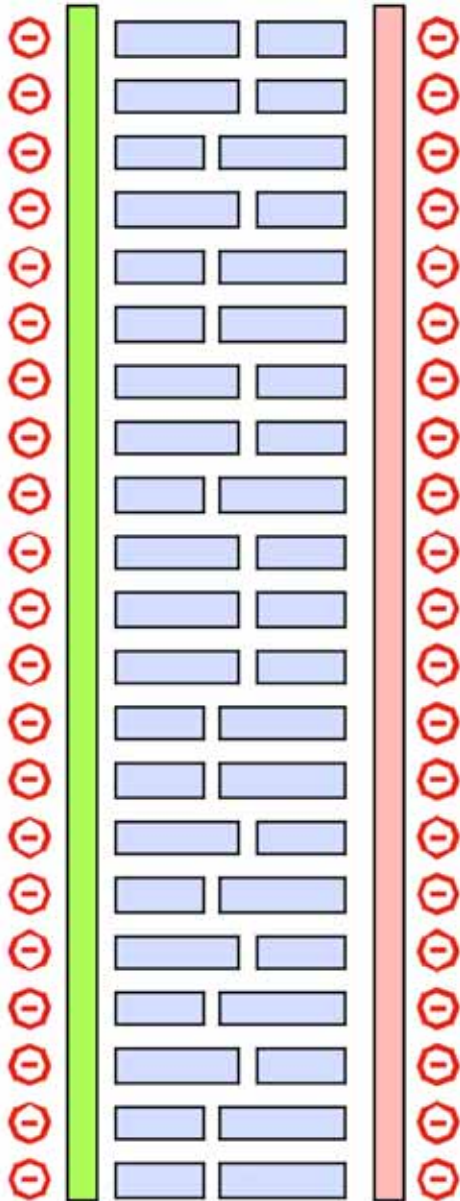


Why *a helix*?



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