

STRUCTURE OF NUCLEIC ACIDS

Course: Molecular Biology (02022312)

Instructor: Dr. M A Srouf

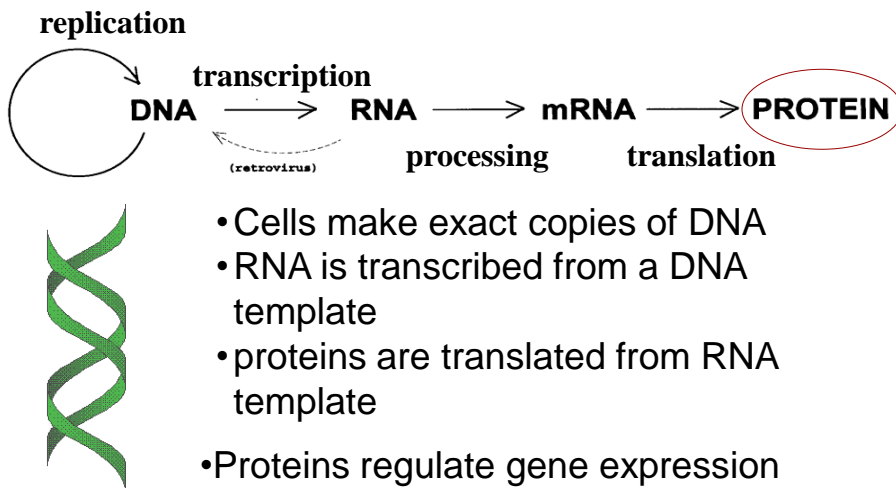
Textbook:

Watson J, Baker TA, Bell SP, Gann A, Levine M, Losick R (2008). Molecular Biology of the Gene, 6th ed. [Chap 6 / pp.101-33](#)

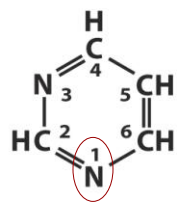
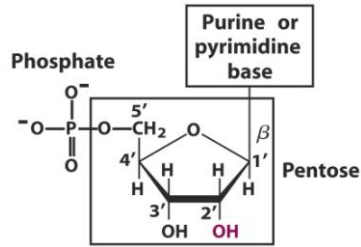
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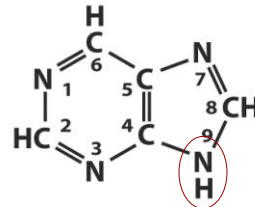
The Central Dogma



Nucleotide structure

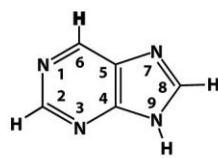


Pyrimidine

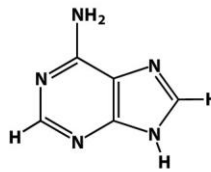


Purine

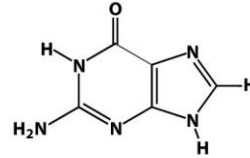
PURINES



Purine

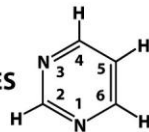


Adenine

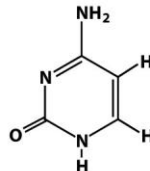


Guanine

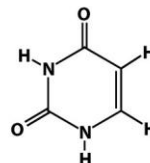
PYRIMIDINES



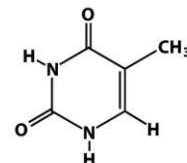
Pyrimidine



Cytosine



Uracil



Thymine

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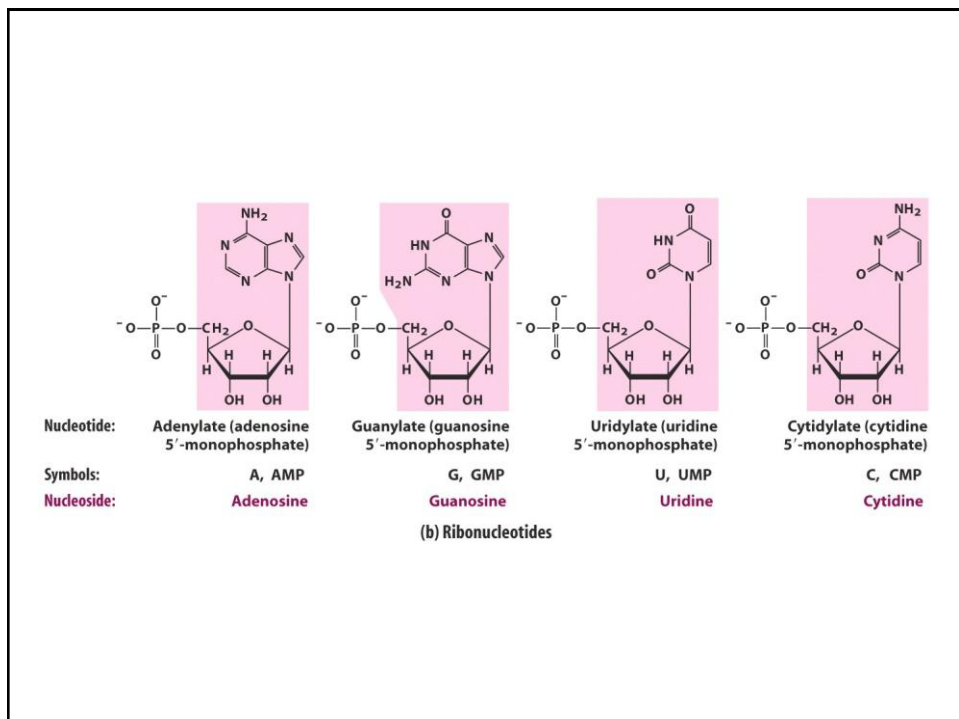
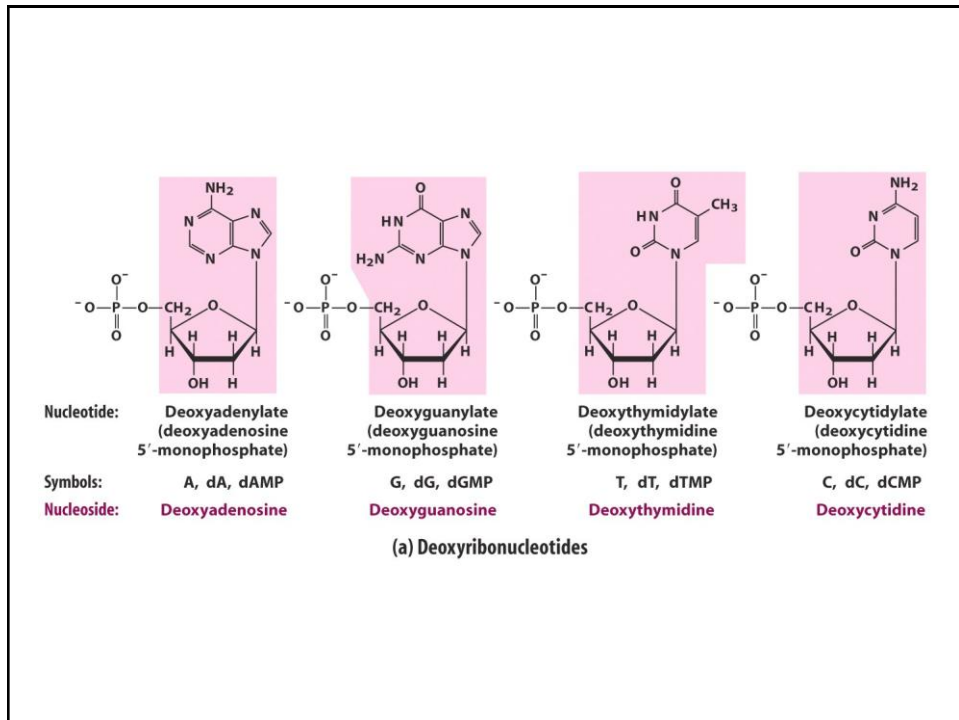
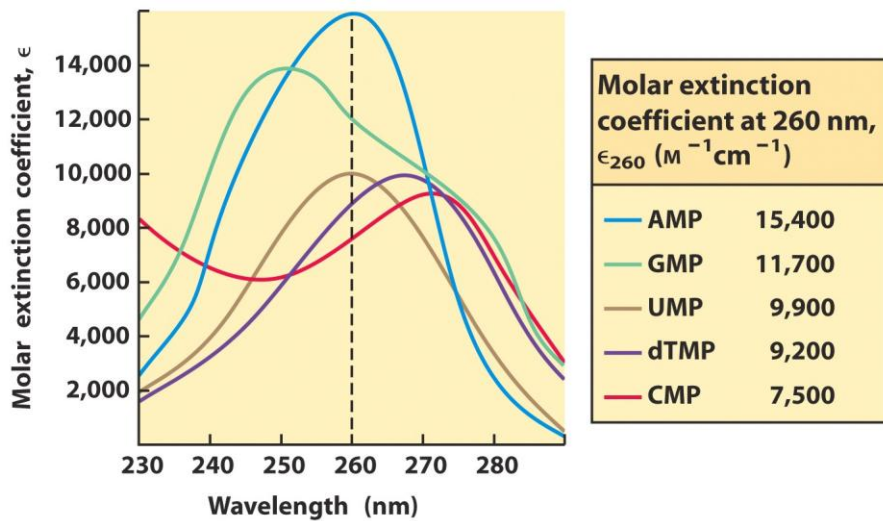
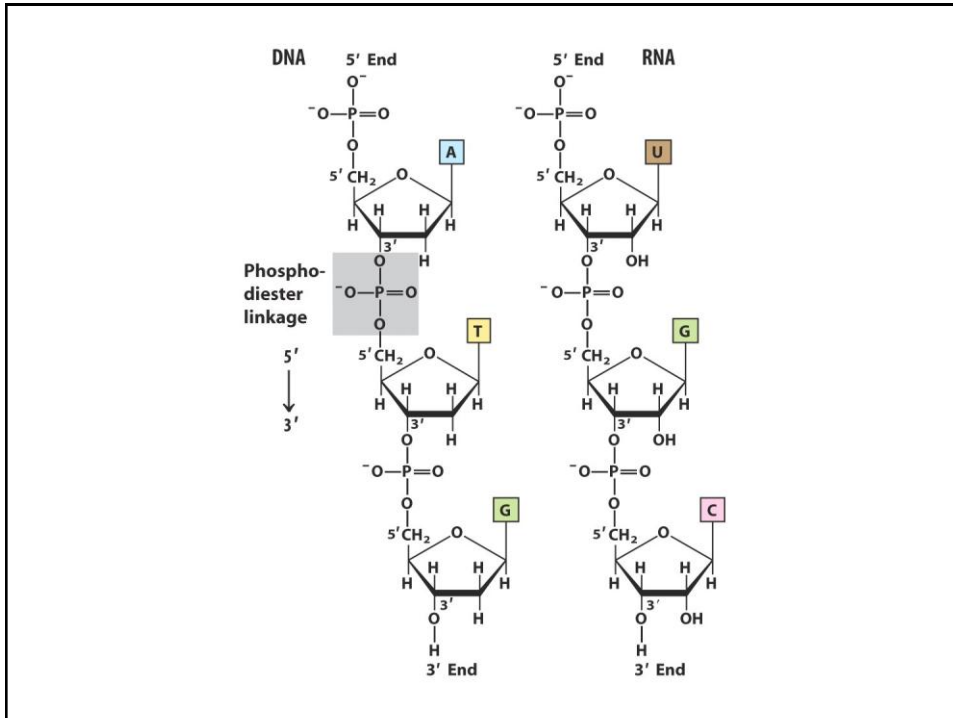


TABLE 8-1 Nucleotide and Nucleic Acid Nomenclature

Base	Nucleoside	Nucleotide	Nucleic acid
Purines			
Adenine	Adenosine	Adenylate	RNA
	Deoxyadenosine	Deoxyadenylate	DNA
Guanine	Guanosine	Guanylate	RNA
	Deoxyguanosine	Deoxyguanylate	DNA
Pyrimidines			
Cytosine	Cytidine	Cytidylate	RNA
	Deoxycytidine	Deoxycytidylate	DNA
Thymine	Thymidine or deoxythymidine	Thymidylate or deoxythymidylate	DNA
Uracil	Uridine	Uridylate	RNA





Why did DNA evolve to be the carrier of genetic information in the cell as opposed to RNA?

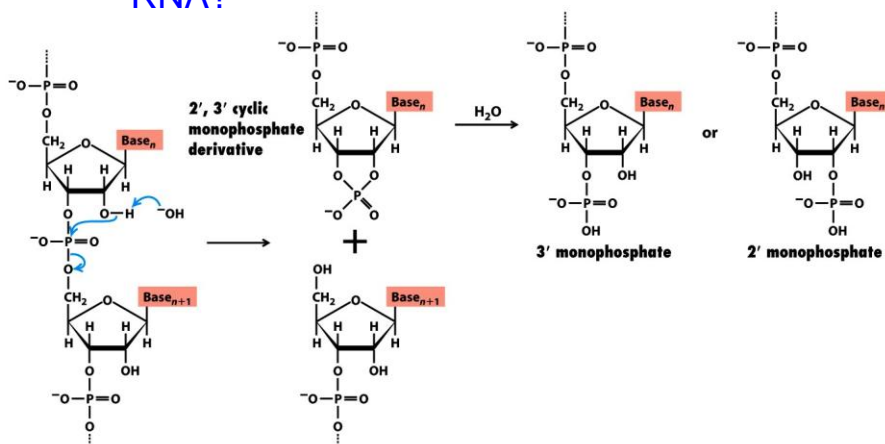


Figure 4-6
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A slow (OH⁻) catalyzed hydrolysis of RNA at neutral pH

Chemical directionality of a nucleic acid strand

(5'→3')

Backbone: S-P-S...

Phosphat is ionized at pH~7.0

Oligonucleotides
<50 nts

Polynucleotides >50 nts

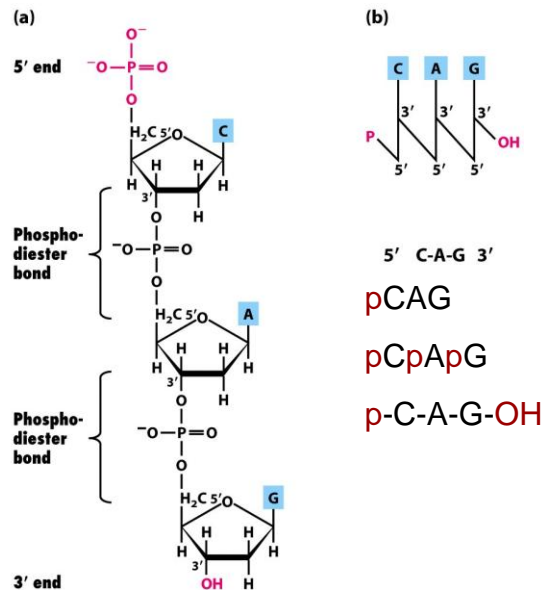
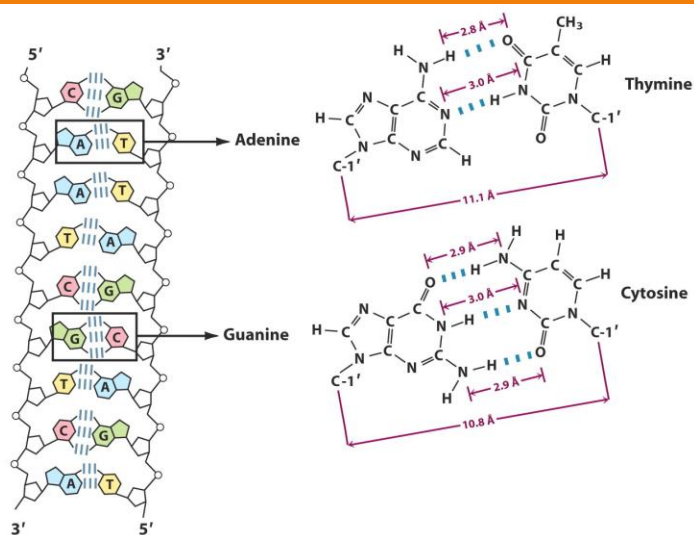


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Watson –Crick model of DNA: hydrogen bonding



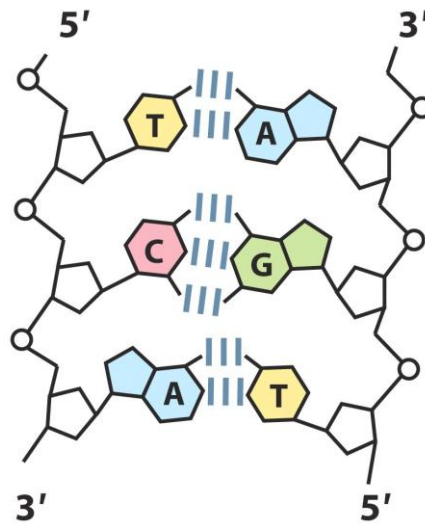
DNA:

Watson-Crick base
pairs > G.C & A.T

Nonstandard base-
pairing > distort the
double helix:

G.T & C.T in DNA

G.U in RNA



X-ray diffraction pattern of DNA:
reveals a helical structure with two periodicities of
0.34 & 34 nm

3.4-Å spacing

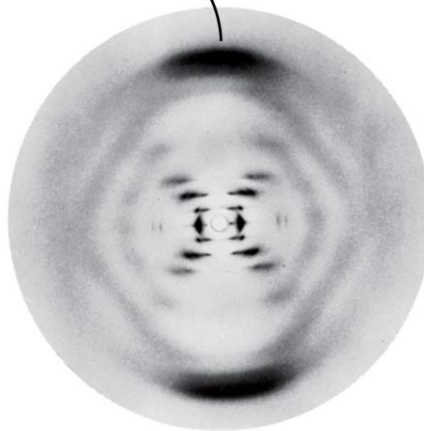


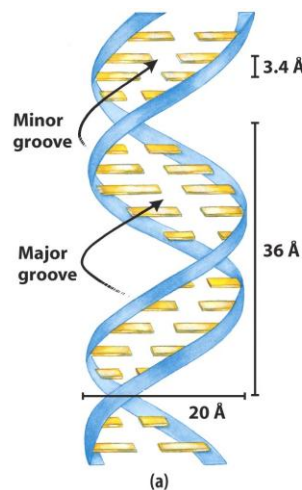
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DNA is a double Helix: Watson-Crick model

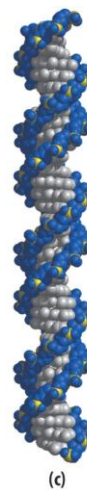
- Rosalind Franklin & Maurice Wilkins, early 1950s> X-diffraction pattern of DNA> DNA has a helical structure with 2 periodicities 0.34 nm & 3.4 nm
- Chargaff's rules: $A=T$ & $G=C$, $A+G=T+C$,
- W&C: right-handed double helix, antiparallel & complementary strands, nt are 0.34 nm apart and 10nt/turn (3.4nm)

W-C model of DNA: a double helix of two complementary antiparallel strands

Chemical structure of DNA double helix



Space-filling model of B DNA



Native DNA: a double helix of two complementary antiparallel strands

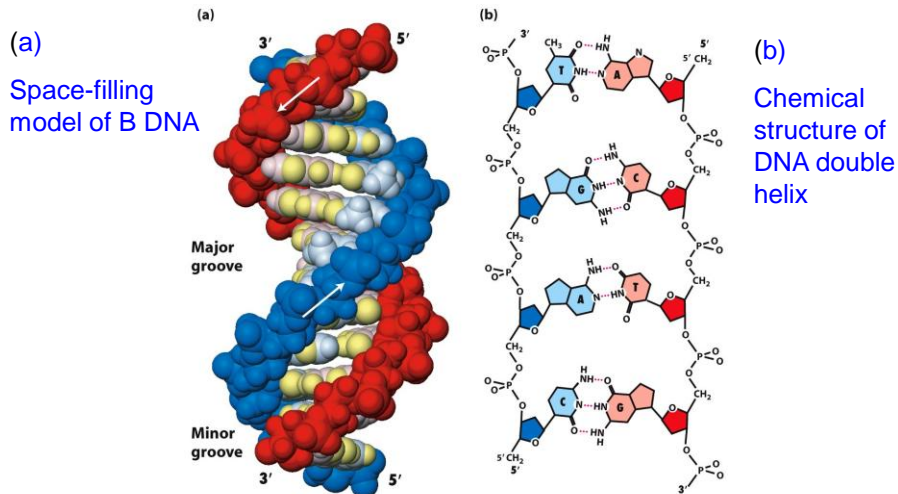
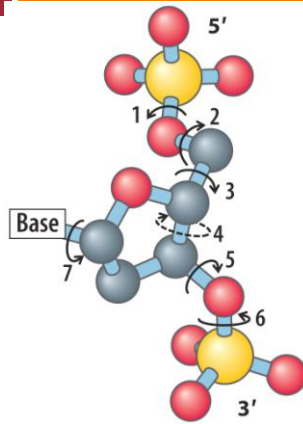


Figure 4-3
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DNA can occur in different 3D forms

- DNA is flexible
- Considerable rotation is possible around a number of bonds in the S-P backbone
- Thermal fluctuations can produce bending, stretching and unpairing (denaturation or melting) of the strands
- Significant deviation in conformation from W-C model exist

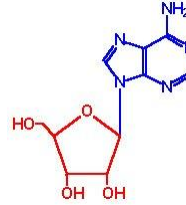
Structural variation in DNA



Rotation around different
bonds in DNA



Syn-Adenosine

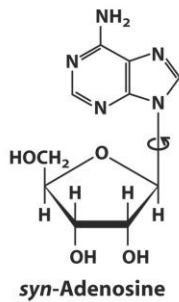


anti-Adenosine

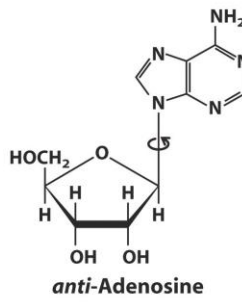
Purines occur in *anti* or *syn* conformation.

Pyrimidines occur only in *anti* conformation.

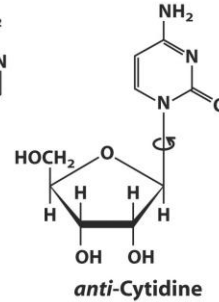
Structural variation in DNA



syn-Adenosine

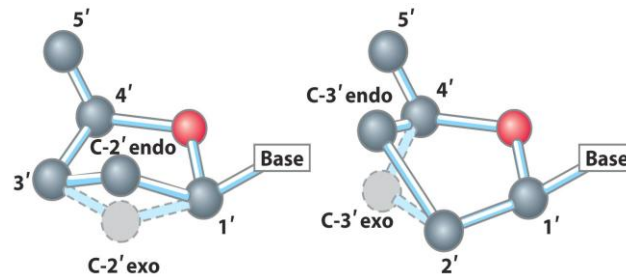


anti-Adenosine

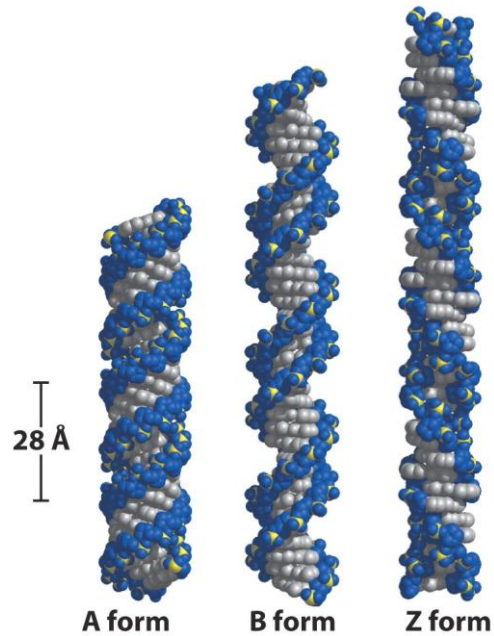


anti-Cytidine

Different puckered Conformations of Ribose



Exo: opposite; Endo: same plane.



DNA can occur in different 3D forms

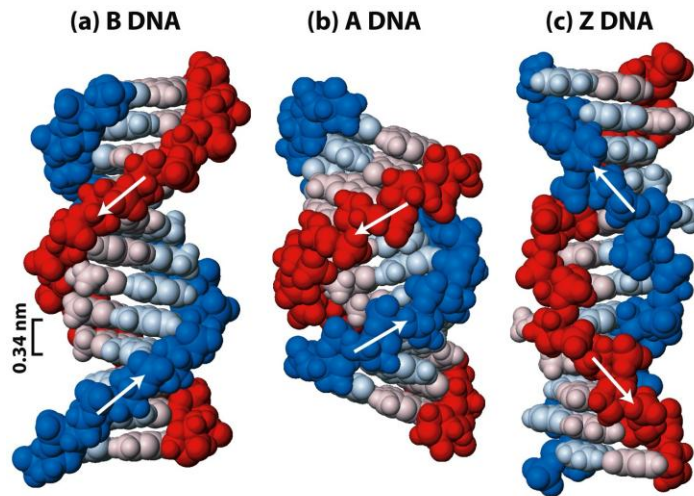


Figure 4-4
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	<i>A form</i>	<i>B form</i>	<i>Z form</i>
Helical sense	Right handed	Right handed	Left handed
Diameter	~26 Å	~20 Å	~18 Å
Base pairs per helical turn	11	10.5	12
Helix rise per base pair	2.6 Å	3.4 Å	3.7 Å
Base tilt normal to the helix axis	20°	6°	7°
Sugar pucker conformation	C-3' endo	C-2' endo	C-2' endo for pyrimidines; C-3' endo for purines
Glycosyl bond conformation	Anti	Anti	Anti for pyrimidines; syn for purines

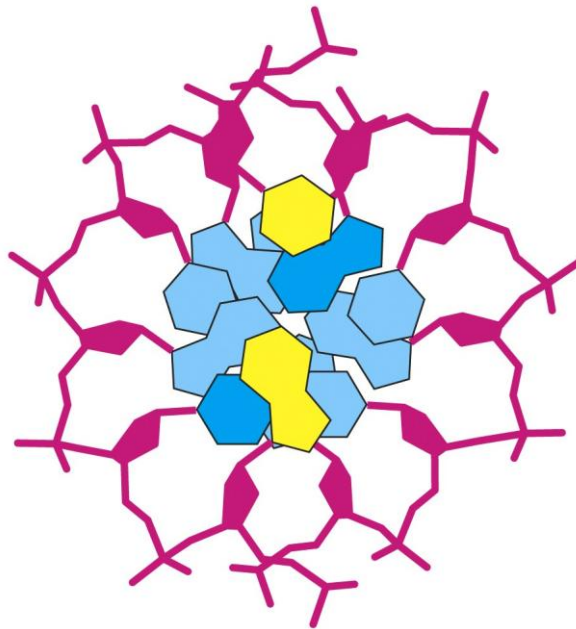
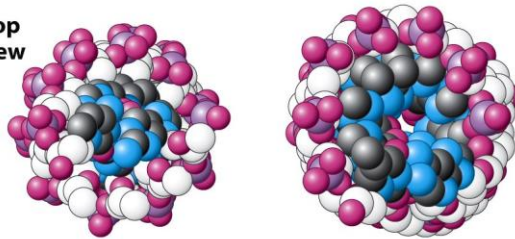
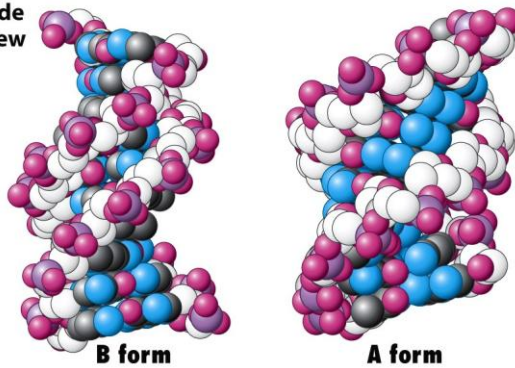


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**Top
 view**



**Side
 view**



B form

A form

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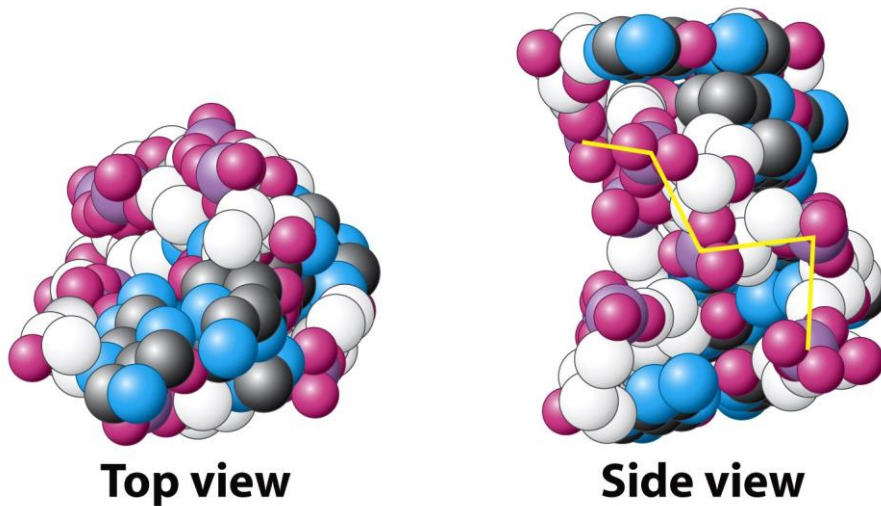


Figure 28-8
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	<i>A form</i>	<i>B form</i>	<i>Z form</i>
Helical sense	Right handed	Right handed	Left handed
Diameter	~26 Å	~20 Å	~18 Å
Base pairs per helical turn	11	10.5	12
Helix rise per base pair	2.6 Å	3.4 Å	3.7 Å
Base tilt normal to the helix axis	20°	6°	7°
Sugar pucker conformation	C-3' endo	C-2' endo	C-2' endo for pyrimidines; C-3' endo for purines
Glycosyl bond conformation	Anti	Anti	Anti for pyrimidines; syn for purines

DNA can undergo reversible strand separation: the hyperchromic effect

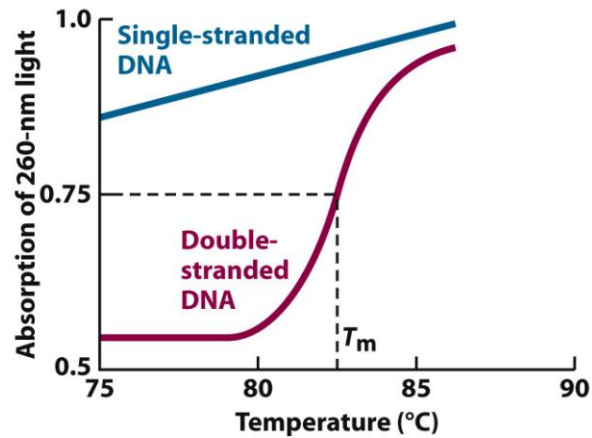


Figure 4-7a
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DNA can undergo reversible strand separation:
 T_m is affected by GC content

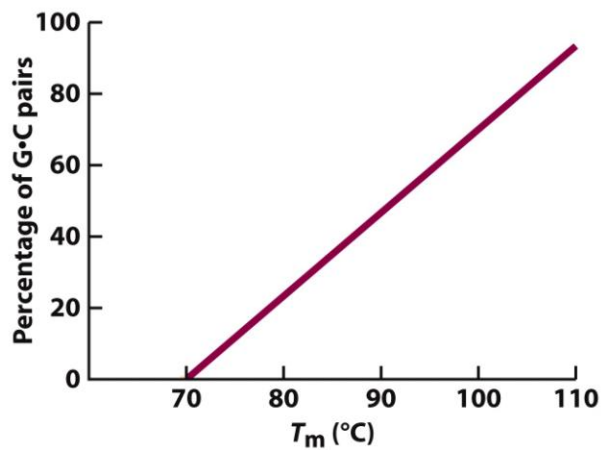


Figure 4-7b
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