

# **From Genes to Proteins**

**Structure-function in DNA and proteins** 

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**Department of Biochemistry** 

### **Aims and Schedule**

- 1) To understand basic principles in structural biology
- 2) To demonstrate a link between structure and function

### Lectures:

Monday – 13/Feb/2017: Genes to Proteins

Friday – 17/Feb/2017: Principles of Structural Biochemistry



### DNA in a 3D world

	Experiment	Conclusion
Late 19th century	Removal of the nucleus resulted in cell death, but removal of an equal volume of cytoplasm did not	The nucleus is key to cell survival
Late 19 <sup>th</sup> century	Removal and transplantation of a nucleus can change cells shape and function	Nucleus is linked to phenotype
Griffiths – 1929	Virulent strains of <i>Pneumococcus</i> were able to transform non-pathogenic <i>Pneumococcus</i> into a disease causing organism	"Some" material has been exchanged between the two cells that is able to change the phenotype of the cells
Avery and MacLeod – 1944	DNA extracted from virulent <i>Pneumococcus</i> . On transformation to non-virulent strains, it conferred pathogenicity. Transfomation not affected by proteases, but by Dnase	DNA is suggested as the transforming agent
Hershey and Chase – 1953	Infected <i>E. coli</i> with T2 phage. DNA: 32P labelled, coat protein: 35S labelled. Only 32P detected inside cells	DNA alone is responsible for cells phenotype, no proteins involved

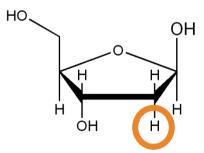
Molecular Structure of Nucleic Acids: A Structure for Deoxyribose Nucleic Acid, - Francis Crick, James D. Watson, Nature, **171**, 1953



## **Building DNA**

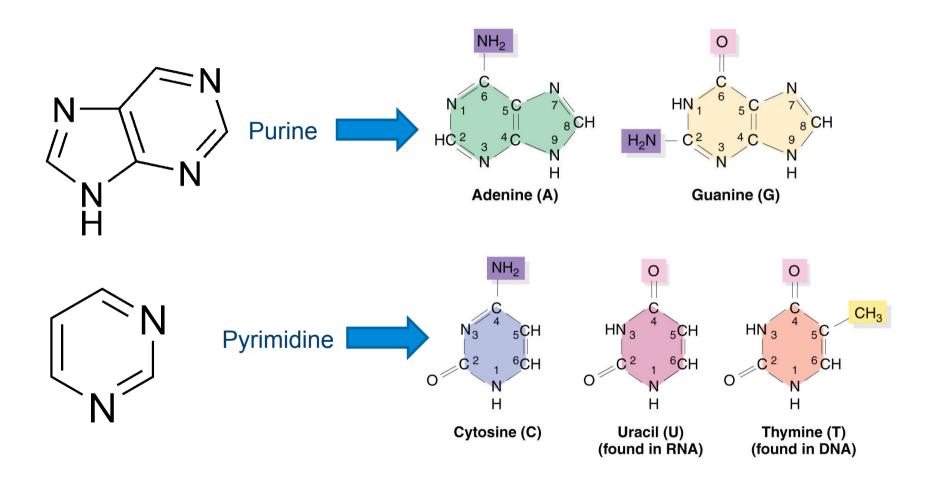
- Both DNA and RNA contain:
  - 1. Pentose sugars (5C)
  - 2. Organic bases
  - 3. Inorganic phosphate

Ribose (Forms ribonucleic acid – RNA)

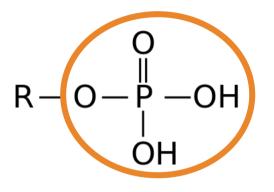


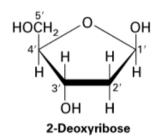
Deoxyribose (Forms <u>deoxy</u>ribonucleic acid – DNA)

### Bases



### **Nucleotides**





Phosphate group

Base	Nucleoside	(Deoxy)ribonucleotide
Adenine	Adenosine	(Deoxy)adenosine 5'-mono/di/triphosphate
Guanine	Guanosine	(Deoxy)guanosine 5'-mono/di/triphosphate
Cytosine	Cytidine	(Deoxy)cytidine 5'-mono/di/triphosphate
Thymine	Thymidine	(Deoxy)thymidine 5'-mono/di/triphosphate
Uracil	Uridine	Uridine 5'-mono/di/triphosphate

Ribonucleoside Deoxyribonucleoside

Glycoside link



### Regulatory nucleotides

 Nucleotides play other roles in a range of processes, including metabolism and cell signalling

Cyclic AMP (cAMP)

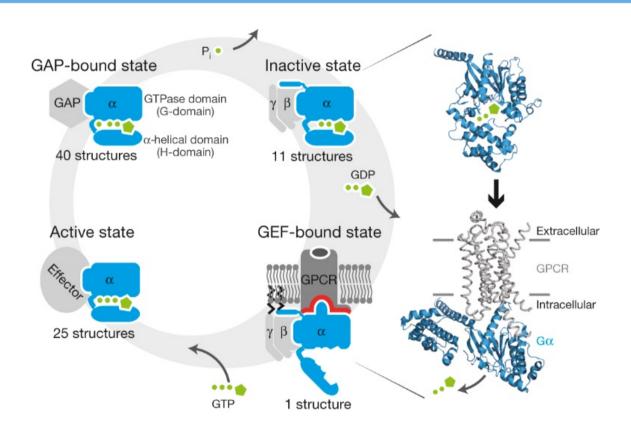
Second messenger GPCR→PKA Cyclic GMP (cGMP)

Second messenger

 $Photoreceptor {\rightarrow} phosphodiesterase$ 

→cGMP gated Na channel →hyperpolarisation

## **GPCRs and G-proteins**



Whole system of signalling based on GTP/GDP exchange



### Regulatory nucleotides

Nucleotides play other roles in a range of processes, including

metabolism and cell signalling

NH<sub>2</sub> HO-P=O HO-P=O ÓН

ppGpp

Bacterial alarmone,

Stringent response

Cyclic AMP (cAMP)

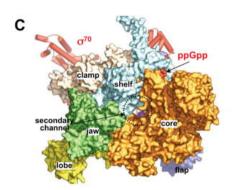
Second messenger GPCR→PKA Cyclic GMP (cGMP)

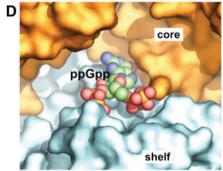
Second messenger

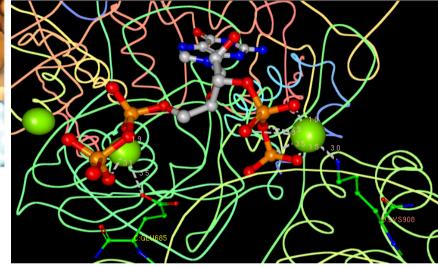
Photoreceptor—phosphodiesterase

→cGMP gated Na channel →hyperpolarisation

### ppGpp







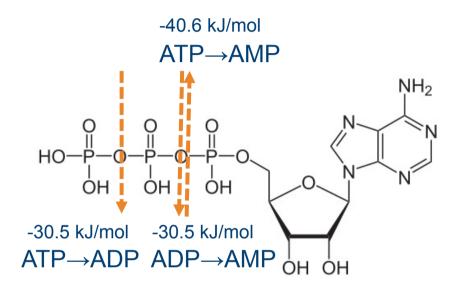
ppGpp in E. coli (PDB: 4JKR)...

...and in *Thermus thermophilus* (PDB: 1SMY). Which is right?

- 1. https://www.ncbi.nlm.nih.gov/pmc/articles/PMC3677725/
  - The Mechanism of *E. coli* RNA Polymerase Regulation by ppGpp Is Suggested by the Structure of Their Complex
  - Mol Cell. 2013 May 9; 50(3): 430-436.
- 2. https://www.ncbi.nlm.nih.gov/pubmed/15109491
  - **Structural Basis for Transcription Regulation by Alarmone ppGpp** Cell. 2004 Apr 30;117(3):299-310.

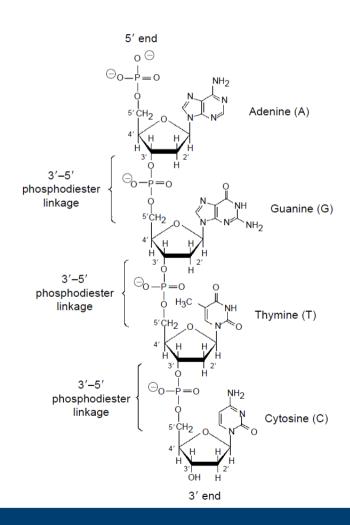


### **Nucleotide chains make oligonucleotides**



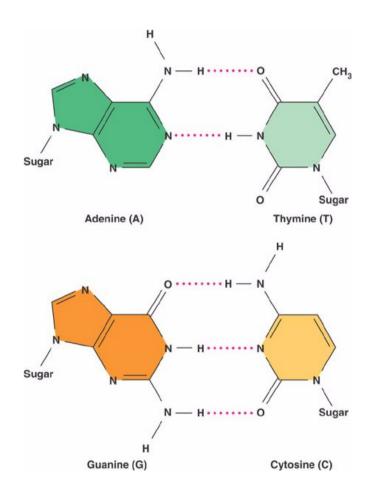
DNA has 5'-3' directionality, always written:

5'-NNNNN-3'





### **Base pairing**



- Purine pairs a Pyrimidine
  - A=T, G≡C
- Chargaff's rules
  - A+G = C+T

Bacteria A: 32% Adenine

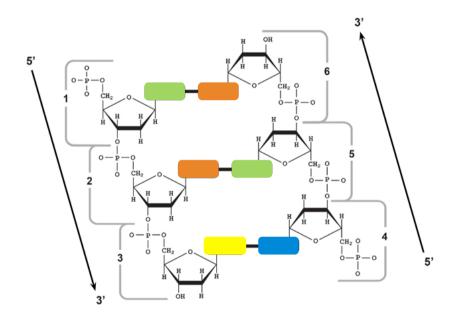
Bacteria B: 17% Adenine

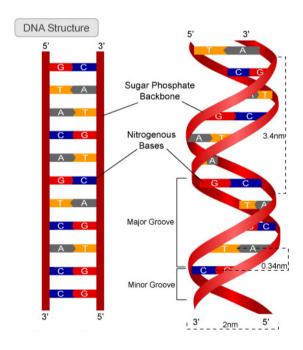
What's the proportion of C, G and T? One of these was isolated from a hot

spring. Which one?

# **Antiparallel DNA helix**

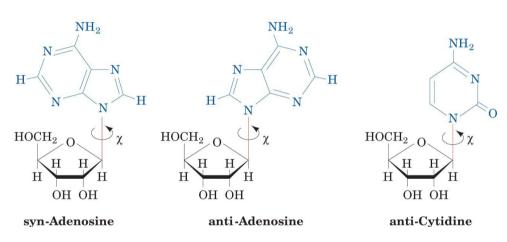
• Two strands: 5'-3': 3'-5'





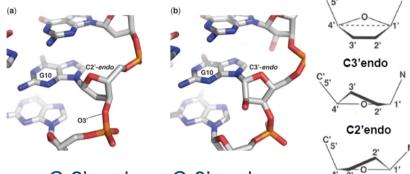
### From sequence to 3D structure

1) Rotation of C1'-N-Glycoside bond



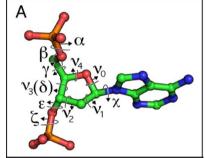
- Purines (A, G) can adopt both syn and antiorientation of the base
- Pyrimidines: =O in steric hindrance with ribose O

2) Sugar puckering



• C-2' endo or C-3' endo

3) Bond rotations in deoxyribose backbone



Rotations mainly about 7 bonds
 Only ribose ring constrained

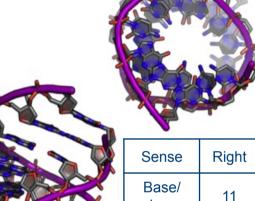


### Forms of the double helix

Alternative forms of DNA exist: some artificially, some naturally

### Dehydrated form

A-DNA



turn

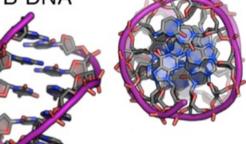
Rise/ bp

2.6 Å

	Base tilt to helix	20°
Lack of water	Glycosyl bond	Anti
Shallow minor groove Deeper major groove	Sugar pucker	C-3' endo

#### Watson-Crick form

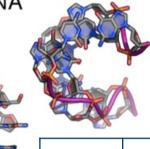
B-DNA



	Sense	Right
•	Base/ turn	10.5
	Rise/ bp	3.4 Å
	Base tilt to helix	6°
	Glycosyl bond	Anti
	Sugar pucker	C-2' endo

#### Rare but observed

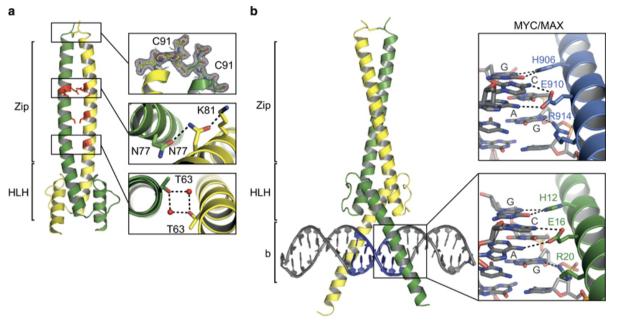
**Z-DNA** 



Sense	Left
Base/ turn	12
Rise/ bp	3.7 Å
Base tilt to helix	7°
Glycosyl bond	Pyr:Anti Pur:Syn
Sugar pucker	Pyr: C-3' endo Pur: C-2' endo



### **Implications of DNA structure**

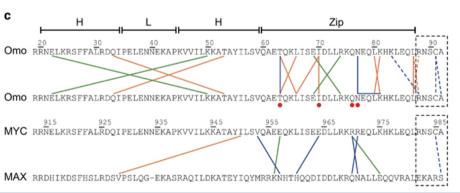


### Myc

Essential role in development and tumorogenesis.
Transcription factor

### **OmoMYC**

Dominant negative variant. Tumour suppressor



doi: 10.1038/onc.2016.354

mutationsalt bridgehydrophobic IAhydrogen bond

disulfide bond



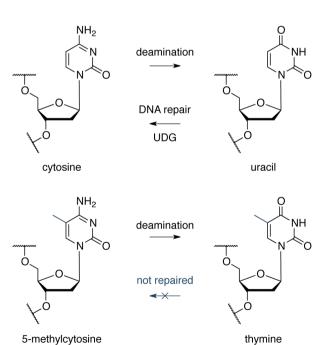
### **DNA** can be modified

- DNA methylation
  - Enzymatic process
- $N_6$  methyl adenine  $C_5$  methyl cytosine
- NHC H<sub>3</sub>

  N 3 5

  O N N H
- N<sub>4</sub> methyl cytosine
- OOC STORY OH OH

- Adenine, Cytosine > Guanine, Thymine
- CH3 source: always S-adenosylmethionine
- Bacteria: Defence and Repair (Dam methylase in GATC)
- Eukaryotes: 5% methylcytidine CpG islands



# **Epigenetics**

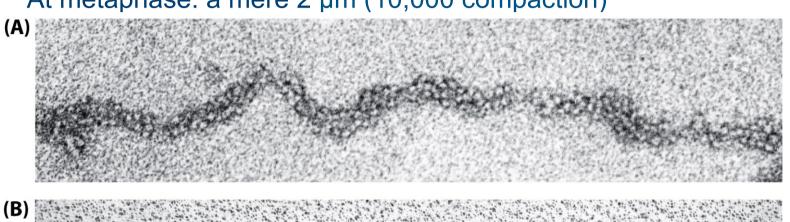
"Stably heritable phenotype resulting from changes in a chromosome without alterations in the DNA sequence"

Cold Spring Harbor, 2008

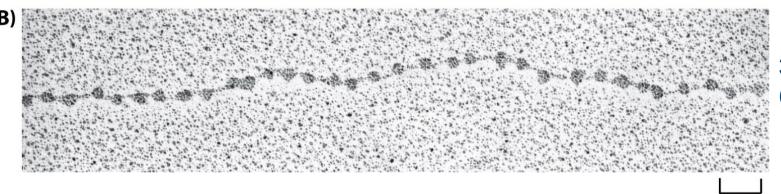


# **DNA** organisation

Chromosome 22: stretched, linear DNA: 1.5 cm At metaphase: a mere 2 µm (10,000 compaction)



30 nm fibre

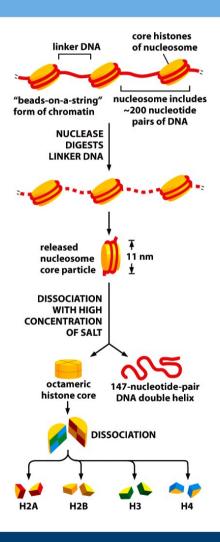


30 nm fibre (decondensed)

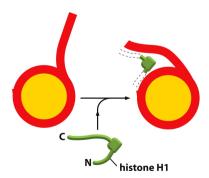




## The chromatin (DNA+Histones)

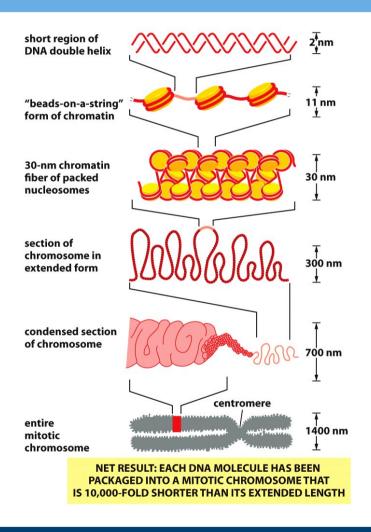


- Chromatin unwound
- Linker DNA digested
- DNA dissociated with high salt
- 147 bp: enough to be wrapped 2x round
- Histone: hetero-octamer + H1



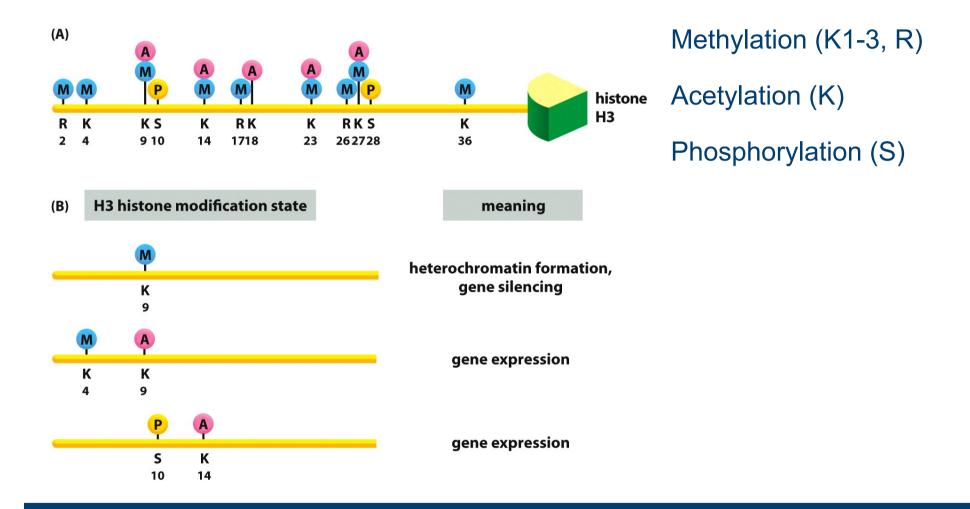


# **DNA** organisation on multiple levels





### **DNA** modification at the chromatin level



### **Summary**

- DNA (genetic information) has transforming effect on cells/organisms
- DNA is constructed of nucleotides
- 2x 5'-3' antiparallel strands form double helix
- DNA is recognised by binding elements
- DNA can be modified, which has functional implication
- DNA is tightly packaged into chromosomes, with a further opportunity for epigenetic variation



### Papers for next session

- https://www.ncbi.nlm.nih.gov/pmc/articles/PMC2938214/
   A crystallographic and modelling study of a human telomeric RNA (TERRA) quadruplex
   Nucleic Acids Res. 2010 Sep; 38(16): 5569–5580
- https://www.ncbi.nlm.nih.gov/pmc/articles/PMC3677725/
   The Mechanism of E. coli RNA Polymerase Regulation by ppGpp Is Suggested by the Structure of Their Complex
   Mol Cell. 2013 May 9; 50(3): 430–436.
- 3. http://www.sciencedirect.com/science/article/pii/S0092867416317391 Structures of the Human HCN1 Hyperpolarization-Activated Channel Cell. 2017 Jan 12;168(1-2):111-120.e11.

