A comprehensive cheat sheet covering DNA structure, experiments proving DNA as genetic material, replication processes, and key figures in its discovery.

#### **Evidence for DNA as Genetic Material**

Griffith's Experiment (1928)	Avery, MacLeod, & McCarty (1944)	Hershey & Chase Experime
<b>Objective:</b> To understand the difference between virulent and non-virulent strains of <i>Streptococcus pneumoniae</i> .	Objective: To identify the molecule responsible for transformation in Griffith's experiment.	Objective: To determine whe
	Experiment: Repeated Griffith's experiment using purified cell extracts from S strain.	Bacteriophages: Viruses that
S Strain (Virulent): Causes pneumonia and kills mice.	Procedure:	Experiment:
R Strain (Non-Virulent): Does not cause pneumonia and does not kill mice.	not cause pneumonia and does not kill mice.	
<b>Experiment 1:</b> Live S strain $\rightarrow$ Mouse dies.	<ol> <li>Removed proteins - Transformation still occurred.</li> <li>Removed RNA - Transformation still occurred.</li> <li>Removed DNA - Transformation did NOT occur.</li> </ol>	Labeled phage DINA with     A sheled phage protein w
<b>Experiment 2:</b> Live R strain $\rightarrow$ Mouse lives.		2. Labeled phage protein w
Experiment 3: Heat-killed S strain → Mouse lives.		Procedure:
Experiment 4: Heat-killed S strain + Live R strain → Mouse dies.	Conclusion: DNA is the genetic material responsible for transformation, at least in bacteria.	1. Infected bacteria with la
<b>Conclusion:</b> Transformation occurred, where the R strain acquired virulence from the dead S strain. Griffith did not identify the transforming principle.		<ol> <li>Separated phages from</li> <li>Measured radioactivity in</li> </ol>
		<b>Results:</b> Radioactive phosph

# remained outside.

**Conclusion:** DNA is the genetic material that is injected into the bacteria and used to produce more bacteriophages. Protein is not the genetic material.

#### **DNA Structure and Components**

Nucleotide Components		DNA vs. RNA Nucleotides		Chargaff's Rules
DNA is a nucleic acid composed of nucleotides:		DNA:	RNA:	Chargaff's Rules:
Deoxyribose: A 5-carbon sugar.		Contains deoxyribose sugar.	Contains ribose sugar.	The amount of Adenine (A) e
Phosphate Group (PO <sub>4</sub> ): Attached to the 5' carbon of the sugar.		Uses Thymine (T) as a base.	Uses Uracil (U) instead of Thymine.	The amount of Cytosine (C)
Nitrogenous Base: Adenine (A), Thymine (T), Cytosine (C), Guanine (G).		Phosphodiester Bonds		The ratio of A-T and G-C var
Hydroxyl Group (-OH): Attached at the 3' carbon of the sugar.				
Purines vs. Pyrimidines		Phosphodiester Bond: Bond between adjacen	t nucleotides.	
		Formed between the phosphate group of one nucleotide and the 3' -OH of the next nucleotide.		
Purines:	Pyrimidines:	Creates a chain of nucleotides with a 5'-to-3' orientation.		
Two-ringed structures (Adenine and Guanine).	Single-ringed structures (Cytosine and Thymine).			

#### **DNA Structure and Replication**

Watson and Crick Model	Antiparallel Configuration	DNA Replication Models
Watson and Crick (1953):	Antiparallel: Each phosphodiester strand has inherent polarity based on the orientation of the sugar-	Conservative Model: Both st
Deduced the structure of DNA using evidence from Chargaff, Franklin, and others.	phosphate backbone.	molecules.
DNA molecule is made of two intertwined chains of nucleotides, forming a double helix structure.	One end terminates in 3' OH, and the other in 5' $PO_4$ .	Semiconservative Model: Da
Double Helix Structure	Strands have either 5'-to-3' or 3'-to-5' polarity.	(Correct model).
	The two strands of a single DNA molecule have opposite polarity to one another.	Dispersive Model: New DNA
Double Helix: Two strands arranged as a double helix.		
Forms two grooves: major groove and minor groove.		
Strands connected via hydrogen bonds between bases on opposite strands.		
Base-Pairing: A-T (2 hydrogen bonds), G-C (3 hydrogen bonds).		

Consistent diameter and stability due to thousands of low-energy hydrogen bonds.



### ent (1952)

ether DNA or protein is the genetic material in bacteriophages.
t infect bacteria, composed of DNA and protein.
h radioactive phosphorus ( <sup>32</sup> P).
vith radioactive sulfur ( <sup>35</sup> S).
ibeled phages.
bacteria.
nside the bacteria.

 $^{\rm 2}$  phosphorus ( $^{\rm 32}\text{P})$  was found inside the bacteria, while radioactive sulfur ( $^{\rm 35}\text{S})$ 

equals the amount of Thymine (T).

equals the amount of Guanine (G).

ries by species.

rands of parental DNA remain intact; new DNA copies consist of all new

aughter strands each consist of one parental strand and one new strand

is dispersed throughout each strand of both daughter molecules after

### **DNA Replication Process**

Requirements for DNA Replication	DNA Polymerase	Semi-Discontinuous Replication
1. Template: Parental DNA molecule to copy.	DNA Polymerase: Matches existing DNA bases with complementary nucleotides and links them to build	Semi-Discontinuous: DNA polymera
	new DNA strands.	Leading Strand: Synthesized contin
2. Enzymes: Proteins to do the copying.	Features:	Lagging Strand: Synthesized discor
3. Building Blocks: Nucleotide triphosphates to make the copy.	<ul> <li>Adds new bases to the 3' end of existing strands.</li> </ul>	Enzymes Involved in Lagging-Str
Stages of DNA Replication	Synthesizes in the 5'-to-3' direction.	DNA Pol III: Synthesizes Okazaki fra
	Requires a primer of RNA to initiate synthesis.	Primase: Makes RNA primer for each
1. Initiation: Replication begins at specific sites called origins of replication.		DNA Pol I: Removes all RNA primers
2. Elongation: New strands of DNA are synthesized by DNA polymerase.		DNA Ligase: Joins Okazaki fragmen
3. Termination: Replication is terminated, often at specific termination sites or when replication forks		
meet.		DNA Gyrase (Topoisomerase): Unlin

polymerase can only synthesize in the 5'-to-3' direction.

I continuously from an initial primer.

discontinuously with multiple priming events, creating Okazaki fragments.

## jing-Strand Synthesis

azaki fragments.

r for each Okazaki fragment.

primers and replaces them with DNA.

fragments to form complete strands.

se): Unlinks two copies of DNA at the termination site.