



NUCLEIC ACIDS

PAPER 2

UNIT 1.2

NUCLEIC ACIDS

- Hereditary determinants
- Macromolecules
- Free state
- Bound state– Nucleoproteins

Nucleotides are biopolymers of high molecular weight with mononucleotides as their repeating units

NUCLEIC ACIDS

DNA (Deoxyribonucleic acid)

RNA (Ribonucleic acid)

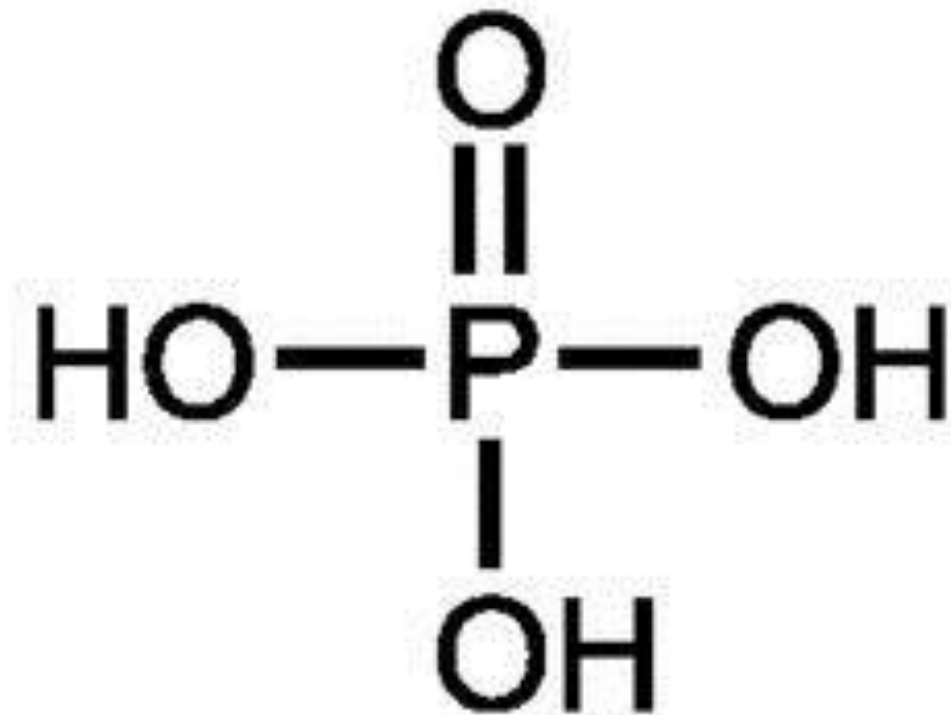
NUCLEIC ACIDS

- DNA--- Chromatin of cell nucleus
- RNA--- Cell cytoplasm, Nucleolus
- Extranuclear DNA--- Mitochondria, Chloroplast

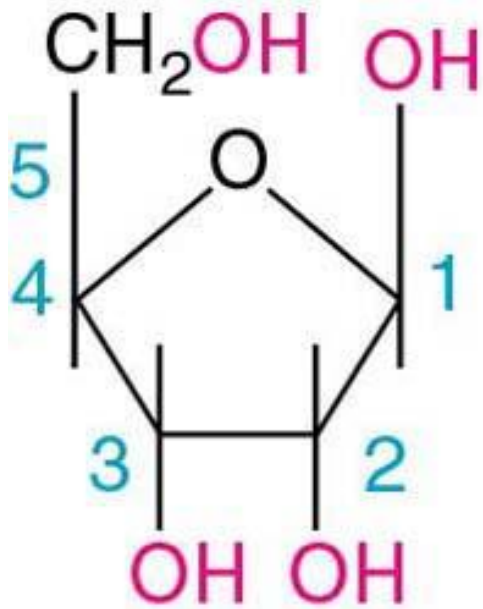
COMPOSITION OF NUCLEIC ACIDS

NUCLEIC ACID= PHOSPHORIC ACID + PENTOSE SUGAR+ NITROGENOUS
BASE

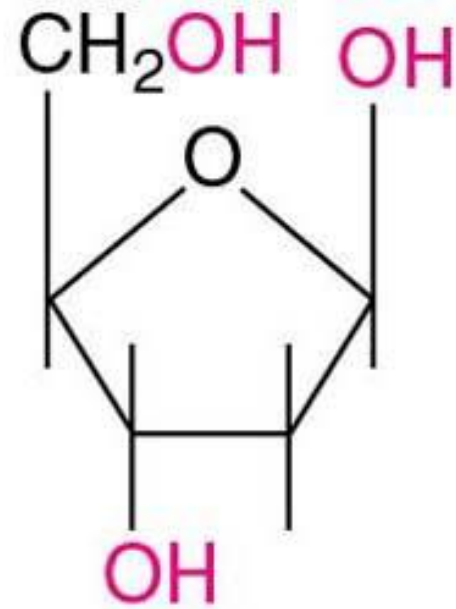
PHOSPHORIC ACID



PENTOSE SUGAR

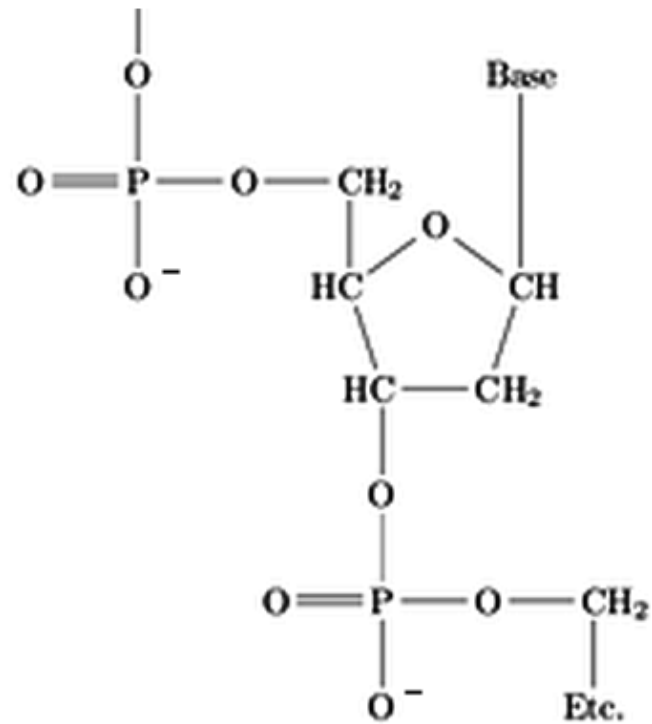


Ribose



2-deoxyribose

PHOSPHODIESTER BOND

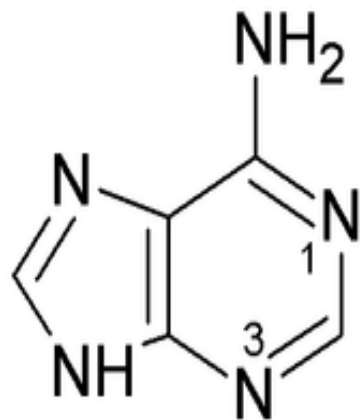


NITROGENOUS BASES

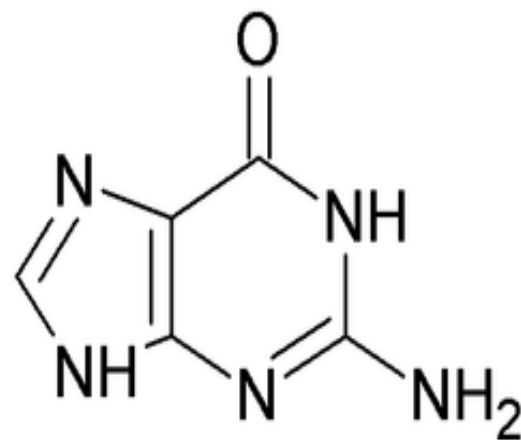
PYRIMIDINES

PURINES

Purines

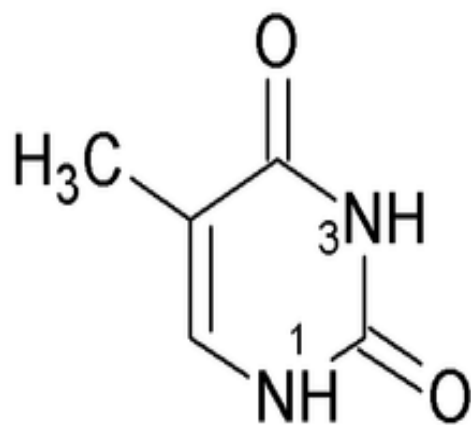


Adenine

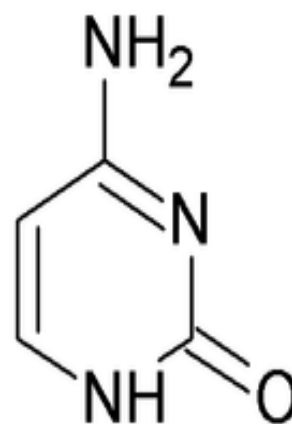


Guanine

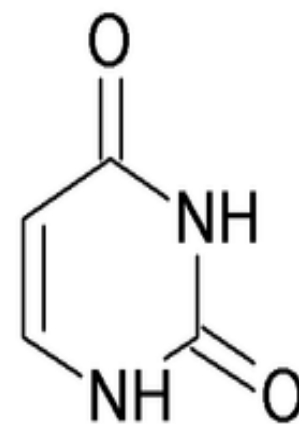
Pyrimidines



Thymine



Cytosine



Uracil

NUCLEOSIDES

NITROGENOUS BASES– PURINES OR PYRIMIDINES ARE CONJUGATED TO THE PENTOSE SUGARS RIBOSE OR DEOXYRIBOSE BY β -GLYCOSIDIC LINKAGE

NUCLEOSIDES

RIBONUCLEOSIDES

DEOXYRIBONUCLEOSIDES

RIBONUCLEOSIDES

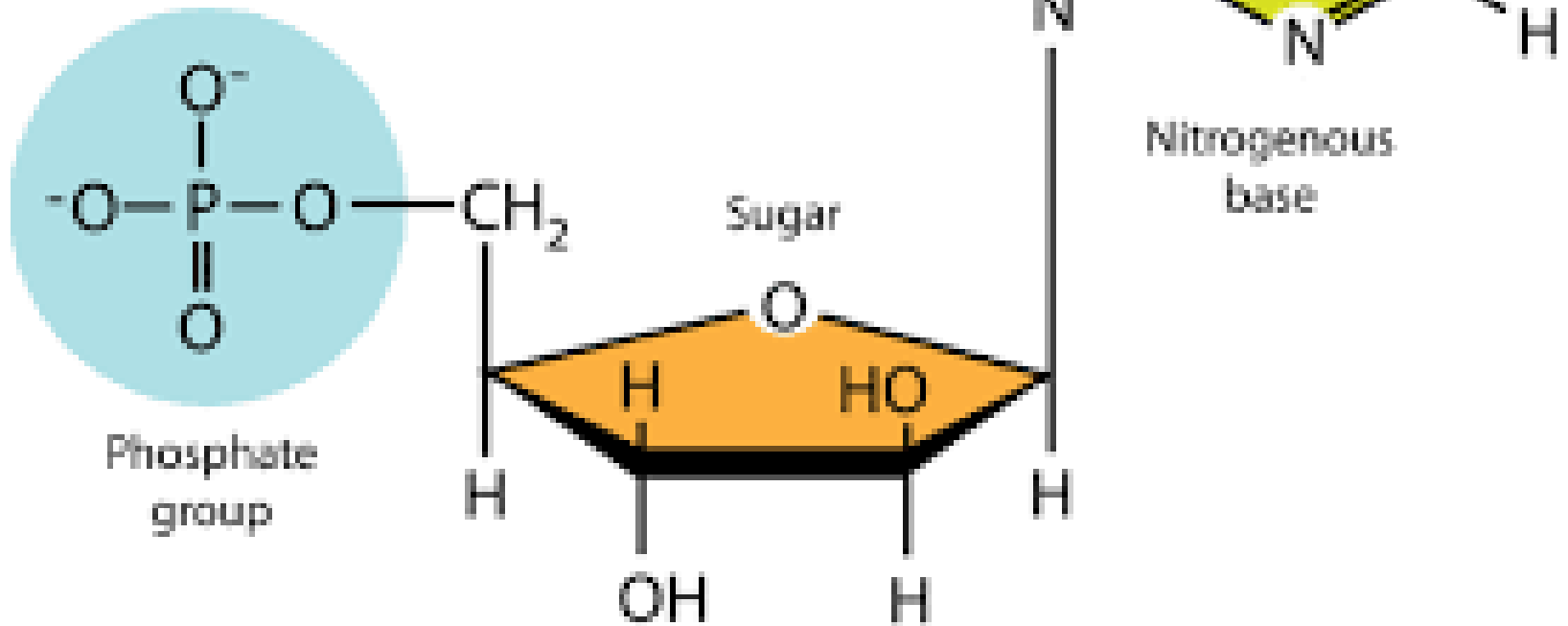
| BASE | NUCLEOSIDE | TRIVIAL NAME | ABBREVIATION |
|----------|----------------------------|--------------|--------------|
| ADENINE | ADENINE RIBONUCLEOSIDE | ADENOSINE | AR |
| GUANINE | GUANINE RIBONUCLEOSIDE | GUANOSINE | GR |
| CYTOSINE | CYTOSINE RIBONUCLEOSIDE | CYTIDINE | CR |
| THYMINE | THYMINE RIBONUCLEOSIDE | THYMIDINE | TR |
| URACIL | URACIL RIBONUCLEOSIDE | URIDINE | UR |

DEOXYRIBONUCLEOSIDES

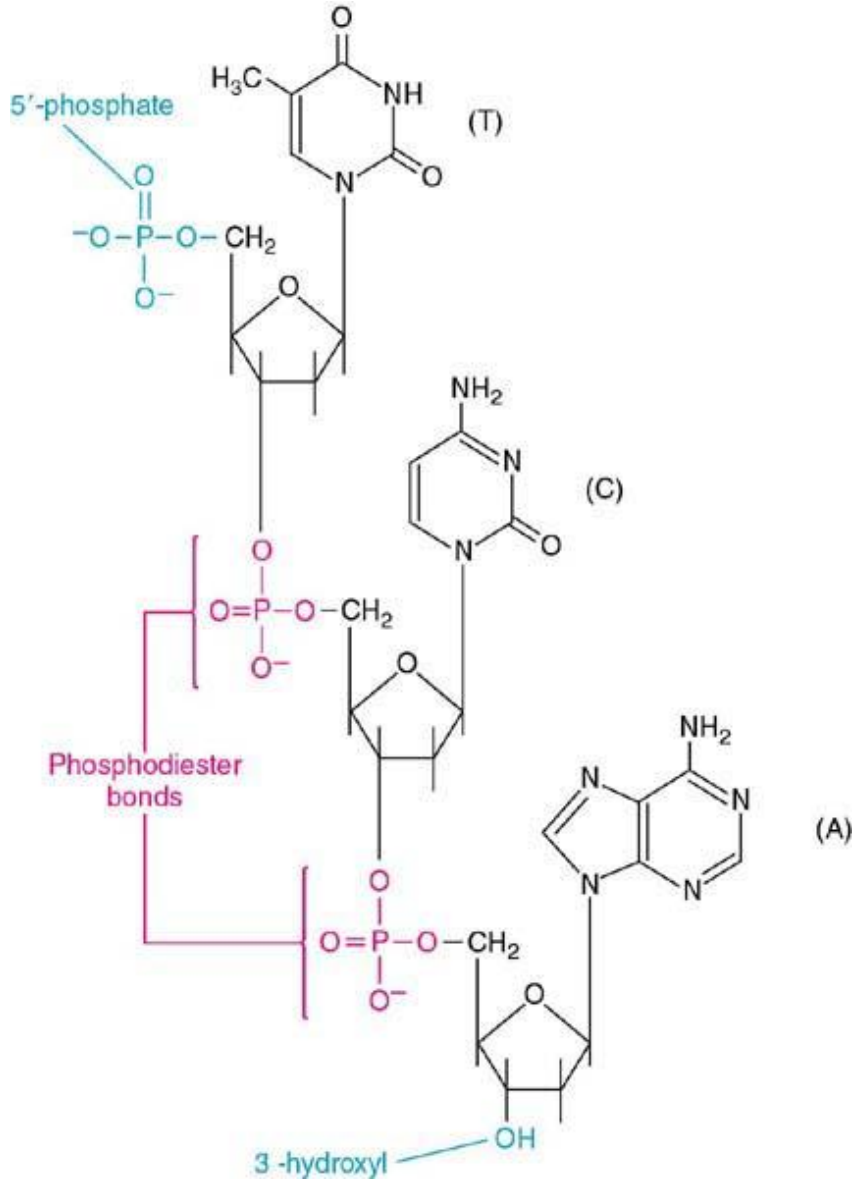
| BASE | NUCLEOSIDE | TRIVIAL NAME |
|----------|---------------------------------|----------------|
| ADENINE | ADENINE DEOXYRIBONUCLEOSIDE | DEOXYADENOSINE |
| GUANINE | GUANINE DEOXYRIBONUCLEOSIDE | DEOXYGUANOSINE |
| CYTOSINE | CYTOSINE DEOXYRIBONUCLEOSIDE | DEOXYCYTIDINE |
| THYMINE | THYMINE DEOXYRIBONUCLEOSIDE | DEOXYTHYMIDINE |
| URACIL | URACIL DEOXYRIBONUCLEOSIDE | DEOXYURIDINE |

NUCLEOTIDES ARE
PHOSPHORIC ACID ESTERS OF
NUCLEOSIDES

Nucleotide



NUCLEOTIDES

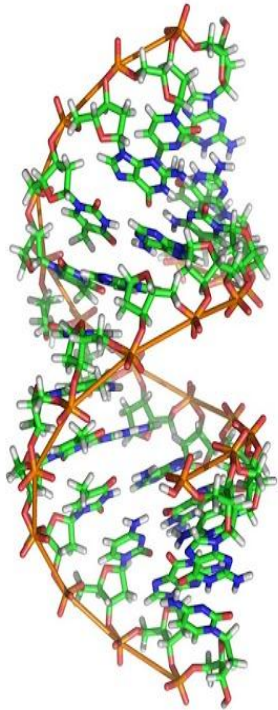


RIBONUCLEOTIDES

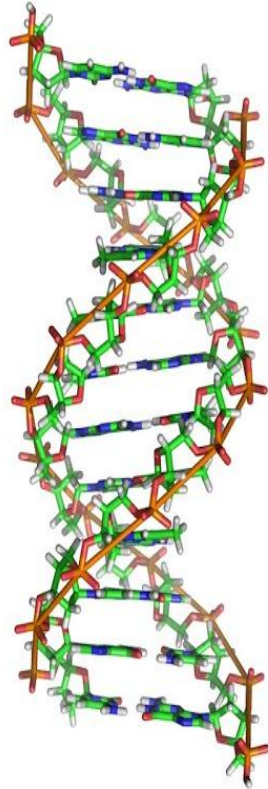
| RIBONUCLEOTIDES | TRIVIAL NAME | ABBREVIATIONS |
|-------------------------------|----------------|---------------|
| ADENOSINE 5' MONOPHOSPHATE | ADENYLIC ACID | AMP |
| GUANOSINE 5' MONOPHOSPHATE | GUANYLIC ACID | GMP |
| CYTIDINE 5' MONOPHOSPHATE | CYTIDYLIC ACID | CMP |
| URIDINE 5' MONOPHOSPHATE | URIDYLIC ACID | UMP |

2' DEOXYRIBONUCLEOTIDES

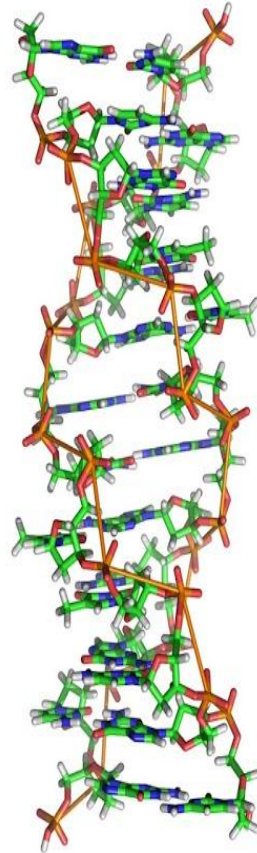
| RIBONUCLEOTIDES | TRIVIAL NAME | ABBREVIATIONS |
|------------------------------------|-----------------------|---------------|
| DEOXYADENOSINE 5' MONOPHOSPHATE | DEOXYADENYLIC ACID | AMP |
| DEOXYGUANOSINE 5' MONOPHOSPHATE | DEOXYGUANYLIC ACID | GMP |
| DEOXYCYTIDINE 5' MONOPHOSPHATE | DEOXYCYTIDYLIC ACID | CMP |
| DEOXYURIDINE 5' MONOPHOSPHATE | DEOXYURIDYLIC ACID | UMP |



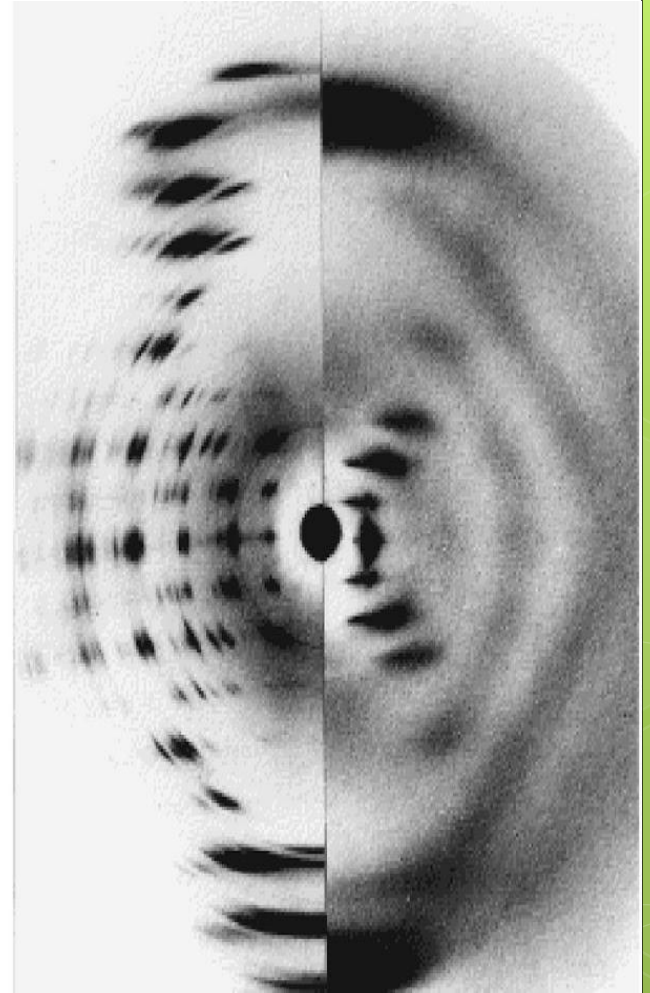
A-DNA



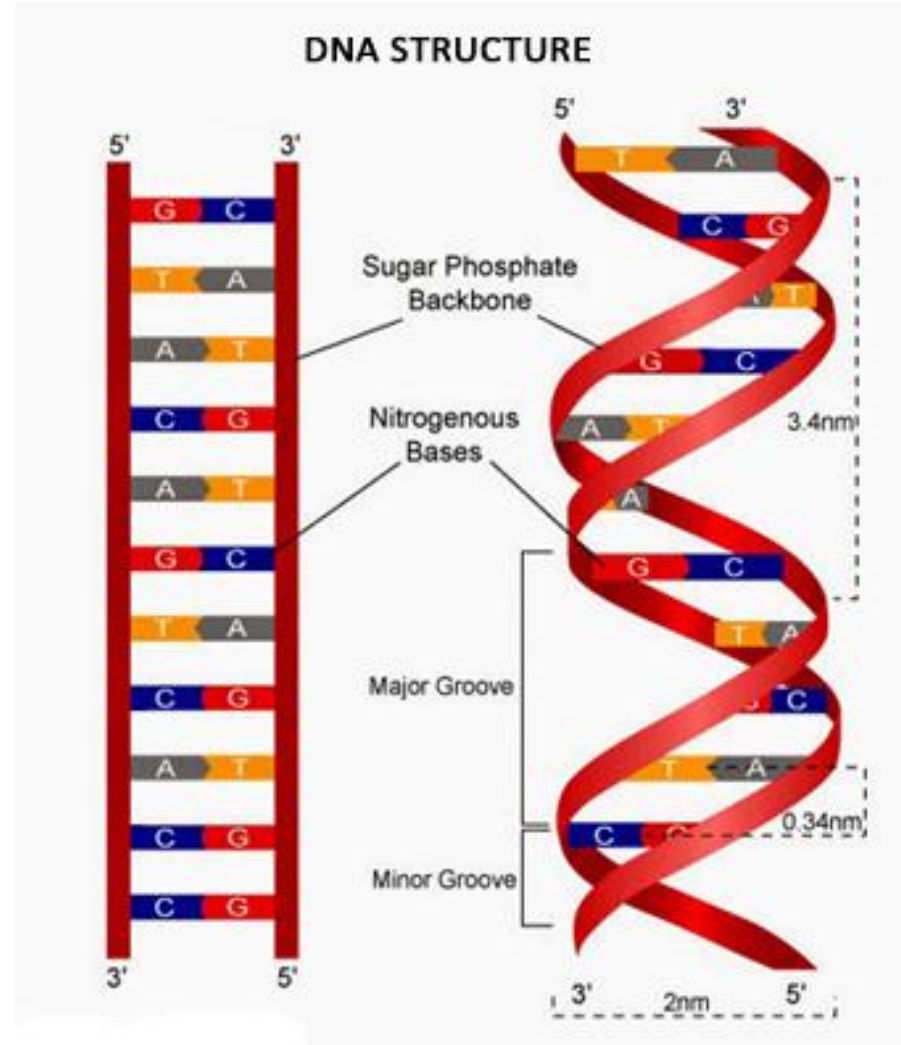
B-DNA



Z-DNA



WATSON AND CRICK MODEL OF DNA



SALIENT FEATURES OF WATSON-CRICK MODEL OF DNA

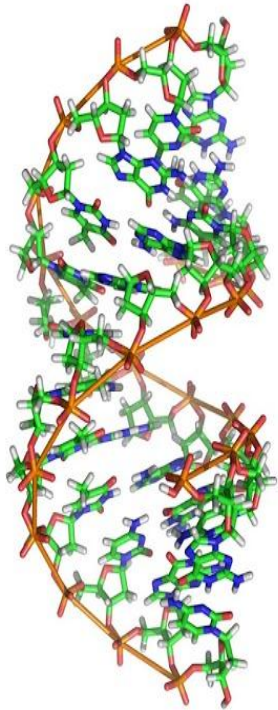
- 2 helical polynucleotide chains which are coiled around a common axis in the form of **Right handed Double helix**.
- Interchain spacings
- **Major groove** (Width 12 Å and 8.5 Å)
- **Minor groove** (Width 6 Å and depth 7.5 Å)
- Base pair are not diametrically opposite to each other
- **Protein interaction** with specific sequence of DNA

SALIENT FEATURES OF WATSON-CRICK MODEL OF DNA

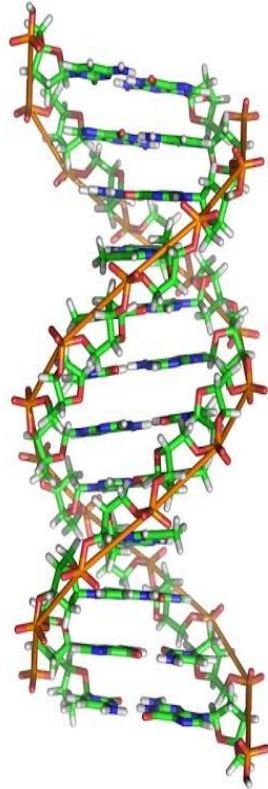
- Plectomic coils– coils interlocked around same axis
- Antiparallel
- Complementary
- Phosphate and deoxyribose --- periphery of the helix
- Purines and Pyrimidines --- centre

SALIENT FEATURES OF WATSON-CRICK MODEL OF DNA

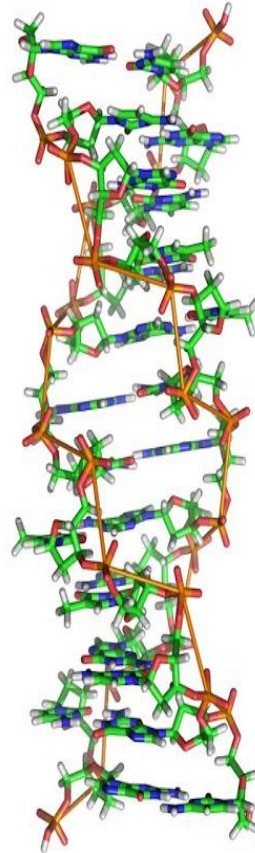
- Diameter of the helix– 20 Å⁰
- Bases--- 3.4 Å⁰
- Each turn of helix – 10 nucleotides
- Hydrogen bonds between base pairs
- A=T, G=C Base Complementarity
- Precise sequence of bases carries the genitic information



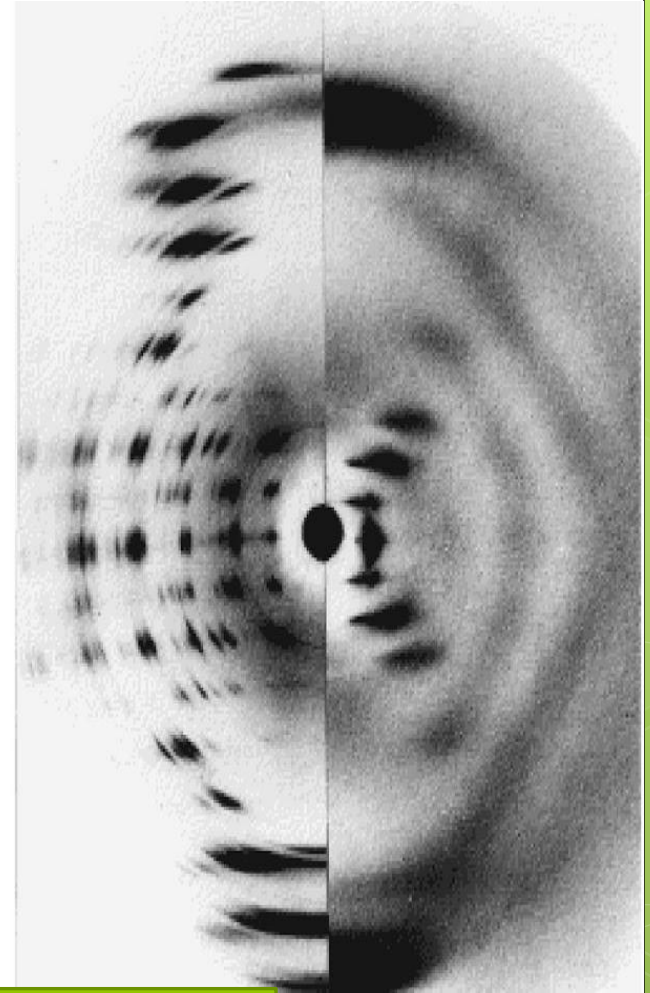
A-DNA



B-DNA



Z-DNA



RIGHT HANDED

LEFT HANDED

RNA types & functions

| Types of RNAs | Primary Function(s) |
|---------------------------------------------------|------------------------------------------------------------------------------------------|
| mRNA - messenger | translation (protein synthesis) regulatory |
| rRNA - ribosomal | translation (protein synthesis) < catalytic > |
| t-RNA - transfer | translation (protein synthesis) |
| hnRNA - heterogeneous nuclear | precursors & intermediates of mature mRNAs & other RNAs |
| scRNA - small cytoplasmic | signal recognition particle (SRP) tRNA processing < catalytic > |
| snRNA - small nuclear snoRNA - small nucleolar | mRNA processing, poly A addition < catalytic > rRNA processing/maturation/methylation |
| regulatory RNAs (siRNA, miRNA, etc.) | regulation of transcription and translation, |

RNA

Ribosomal RNA

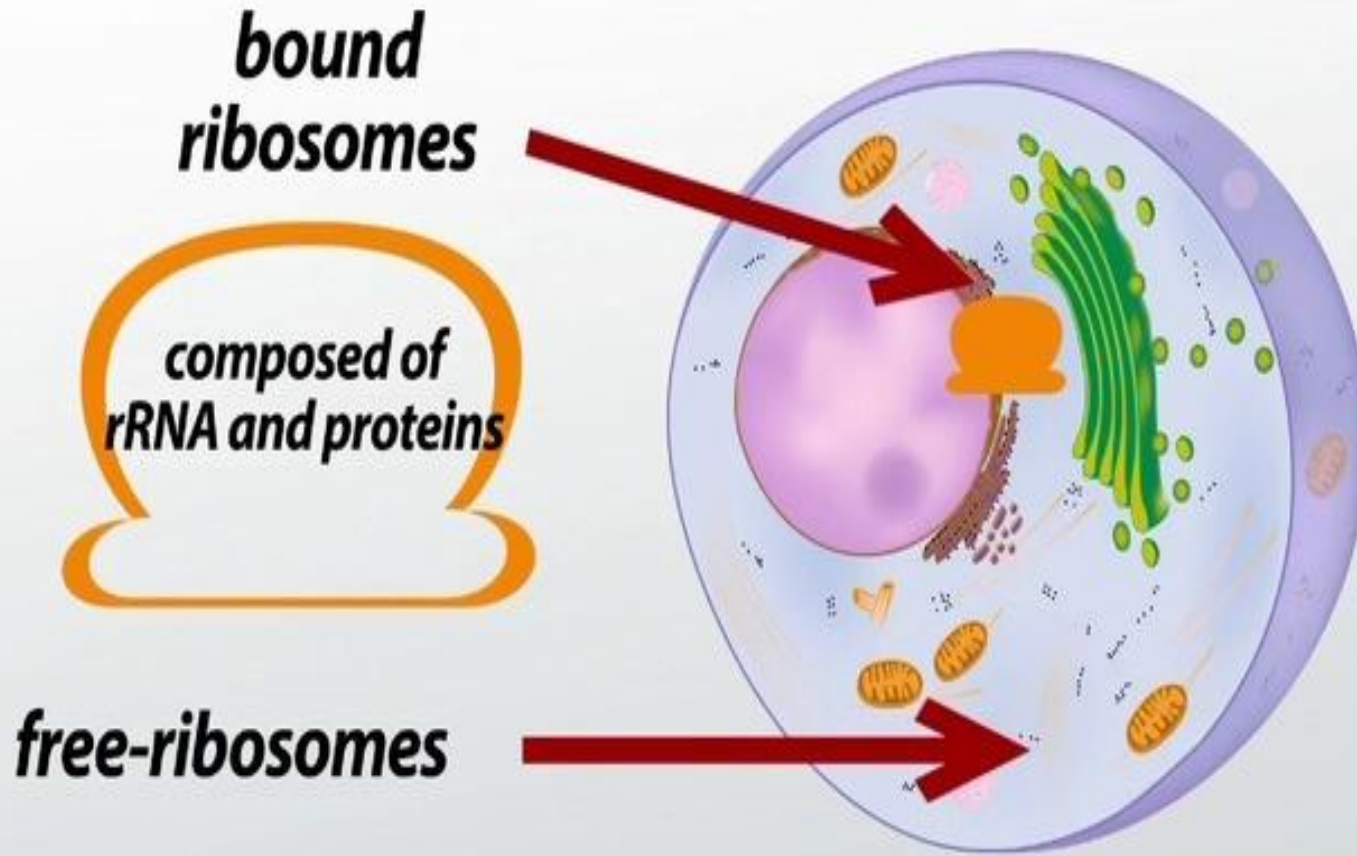
Transfer RNA

Messenger RNA

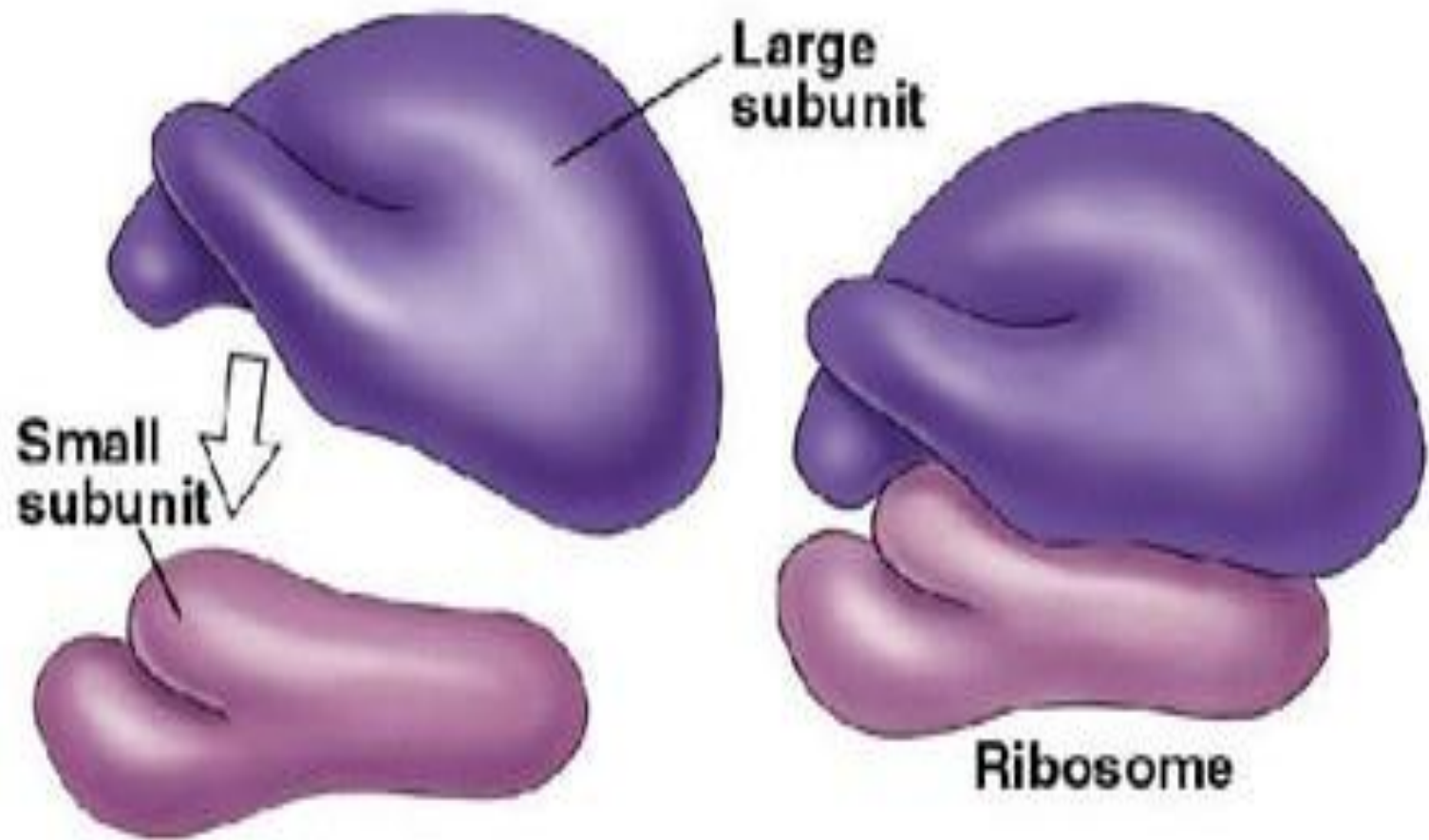
RIBOSOMAL RNA

- Most stable form of RNA
- Found in ribosomes
- Highest molecular weight
- Abundant of all the types
- 80% of the total RNA of the cell
- Represents 40-60% of the total weight of ribosomes

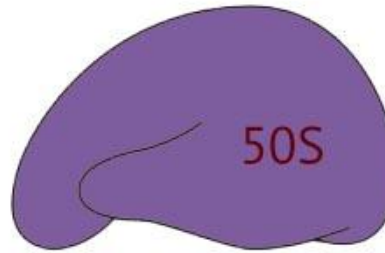
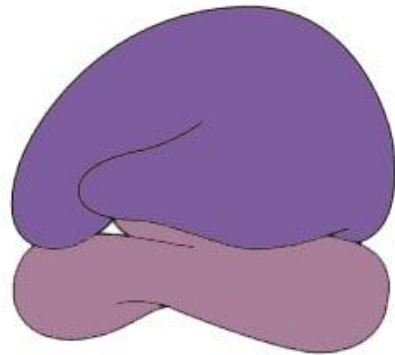
RIBOSOMES PERFORM PROTEIN SYNTHESIS



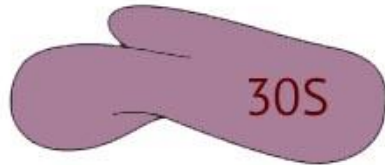
Ribosome



prokaryotic 70S ribosome

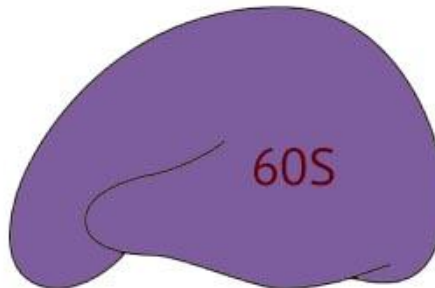


23S and
5S rRNAs
(34 proteins)

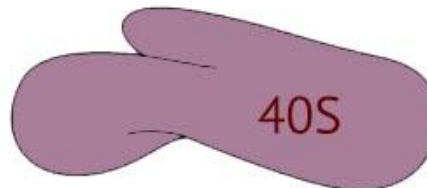


16S
(21 proteins)

eukaryotic 80S ribosome



28S, 5.8S,
and 5S rRNAs
(~45 proteins)



18S rRNA
(~30 proteins)

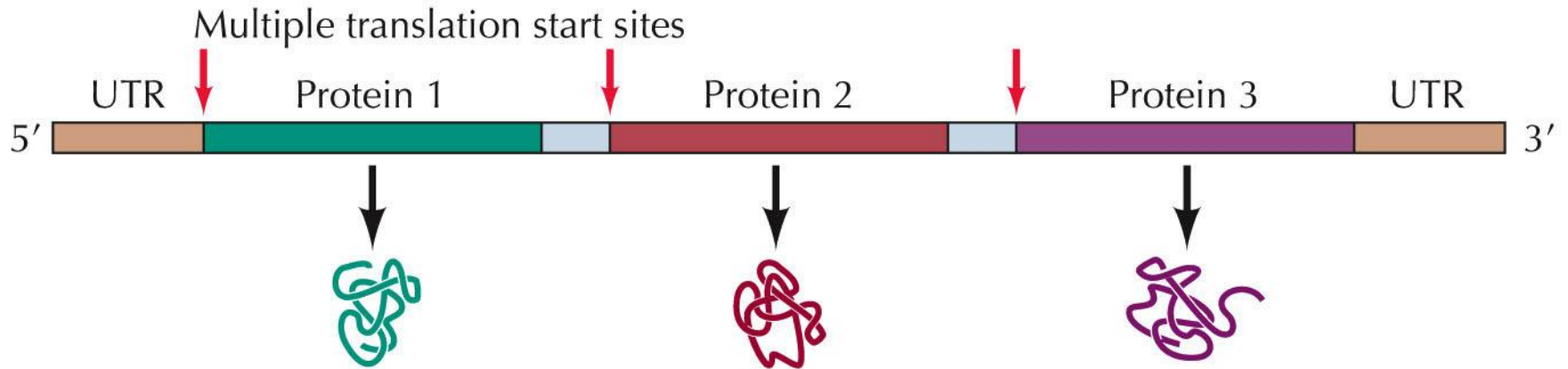
RIBOSOMAL RNA

- Highest GC content
- More than 50%
- Necessary for ribosomal assembly
- Provides specific sequence to which messenger RNA molecule can bind in order to be translated

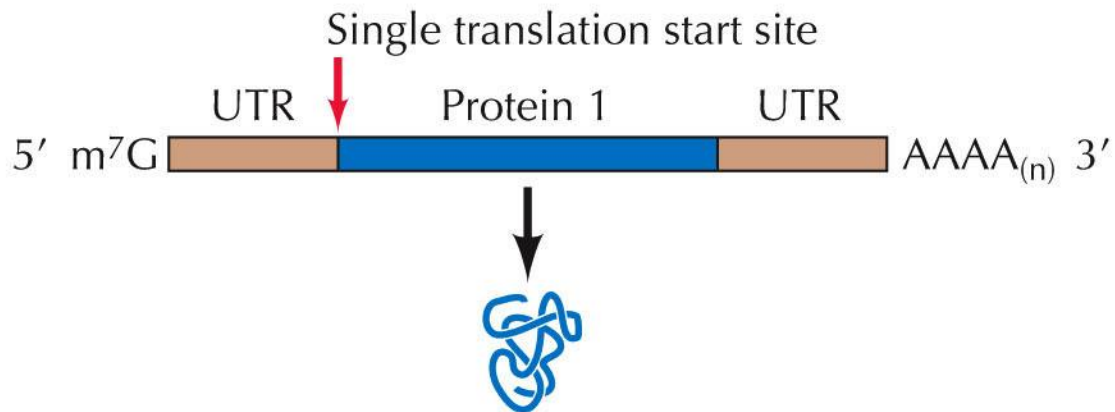
MESSENGER RNA

- Heterogeneous in size and stability
- Amounts to 5% of the total RNA
- Synthesized on the surface of DNA template
- Base sequence complementary to DNA
- Carries the genetic information (message) for the assembly of amino acids from DNA to ribosomes
- Synthesized by DNA dependent RNA polymerase

Prokaryotic mRNA



Eukaryotic mRNA



RNA capping and polyadenylation

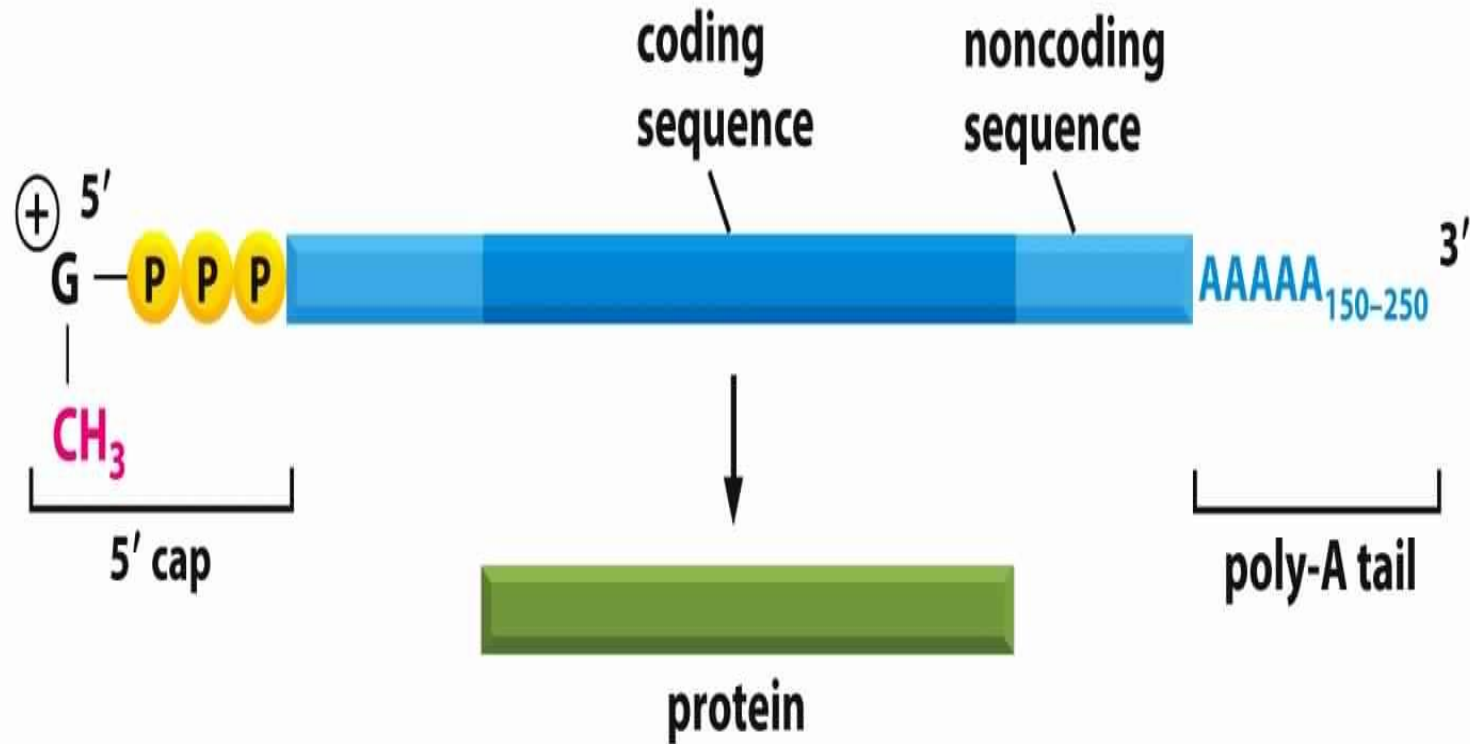


Figure 7-16a Essential Cell Biology 3/e (© Garland Science 2010)

TRANSFER RNA

- t RNA
- Soluble RNA
- Smallest polymeric form of RNA
- 15% of the total RNA of cell
- Acts as specific carriers of activated **amino acids** to specific sites on protein synthesizing templates

Structure of tRNA

- **Primary structure**- linear sequence of nucleotides
- **Secondary structure**-Clover leaf model
- **Tertiary structure**- 3-D structure of tRNA , L shape, Helix stacking

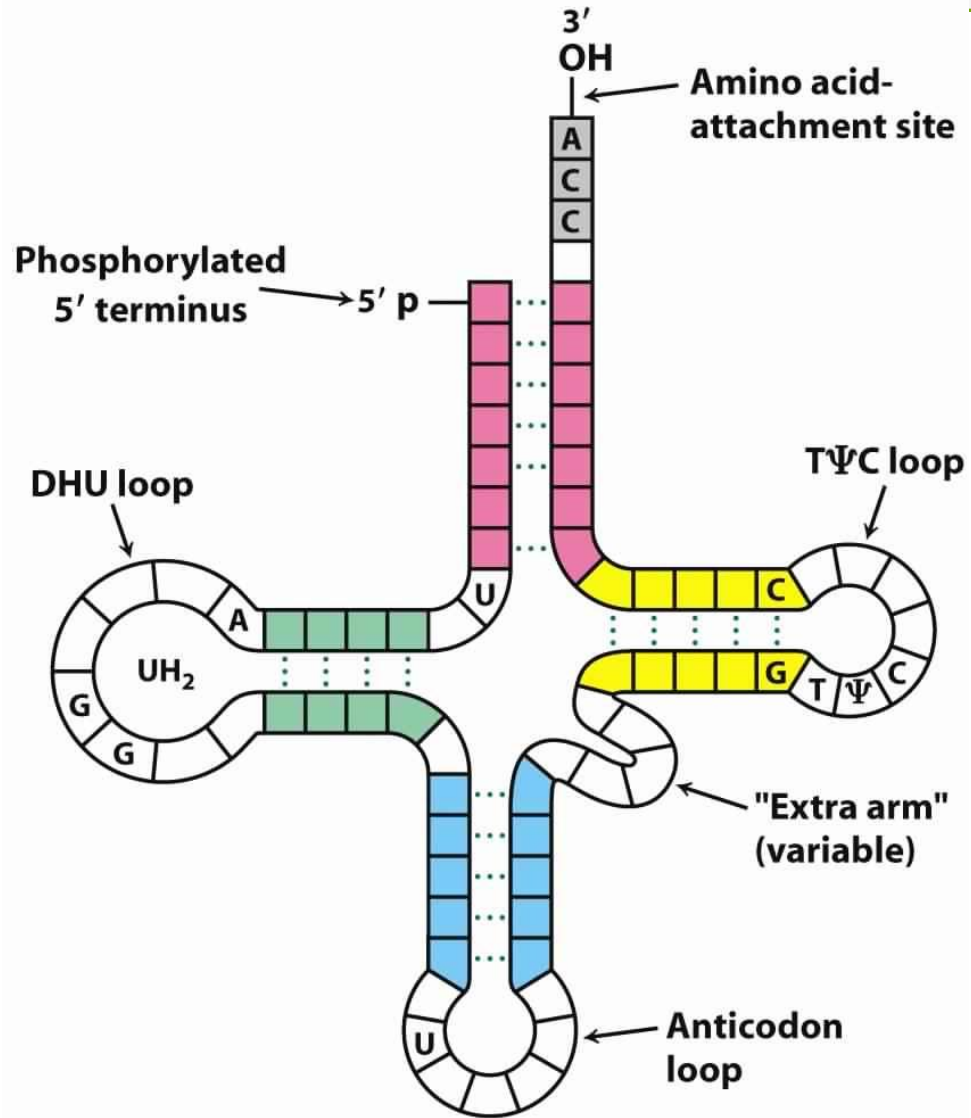


Figure 30.3

Biochemistry, Seventh Edition

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DIFFERENCE BETWEEN DNA & RNA

| | DNA | RNA |
|--------------------|-----------------------------------------------------|-----------------------------------------------------|
| Location | Chromatin of nucleus | Cytoplasm=90% Nucleolus=1-% |
| State | Never present in free state in cytoplasm | May be present in free state |
| Strands | Normally double stranded and rarely single stranded | Normally single stranded and rarely single stranded |
| | Both sense and antisense strands | Sequence is same as that of antisense strand |
| Sugar moiety | 2'deoxyribose | ribose |
| Nitrogenous bases | A, T, G, C (No uracil) | A, U, G, C (No thymine) |
| Base pairing | A=T & G=C | A=U & G=C |
| Base pairing | Entire length | 50% of the entire length (helical region) |
| Unusual base pairs | Few unusual base pairs | More unusual base pairs |

DIFFERENCE BETWEEN DNA & RNA

| | DNA | RNA |
|-------------------------------|-----------------------------------------------------|------------------------------------------------|
| Molecular weight | Large number of nucleotides , high molecular weight | Fewer nucleotides hence low molecular weight |
| Stability | Alkali stable | Alkali labile |
| Synthesis | Acts as template for its synthesis | Does not acts as template for its synthesis |
| Replication and Transcription | Undergoes Replication and transcription | Does not undergo replication and transcription |
| Genetic material | Usual genetic material | Only for some viruses |