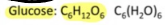
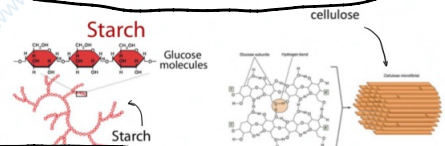


carbohydrates

The name «carbohydrates» means «hydrates of carbons» as many of them have the formula $C_n(H_2O)_m$.



They can form polymers and are used to store energy in living beings (starch, glycogen), as support structures in plants (cellulose), and are essential components of nucleic acids (DNA and RNA).



Carbohydrates can form polymers. Depending on their length, they can be classified as:

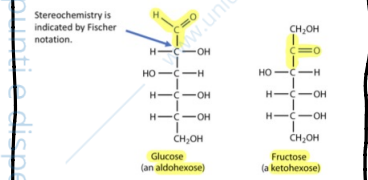
- Monosaccharides (1 unit)
- Bisaccharides (2)
- Oligosaccharides (few units) (2-10)
- Polysaccharides (many units)

Monosaccharides

Simple sugars
Cannot be further hydrolyzed into simple carbohydrates

Building blocks of oligo and polysaccharides

At the molecular level, they are polyhydroxyaldehydes or polyhydroxyketones.

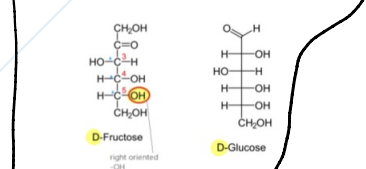


Stereochemistry is indicated by Fischer notation.
The chemical behaviour of carbohydrates is complicated by the presence of multiple chiral centers.

D and L monosaccharides

D-L notation describes the configuration of the chiral carbons of a monosaccharide. It uses as reference the penultimate carbon.

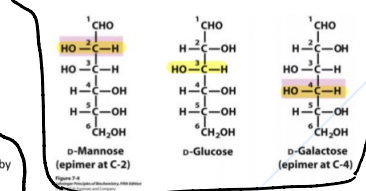
If written in the Fisher projection, with the oxidized part «up», D monosaccharides have the OH group of the penultimate carbon on the right, L have it on the left.



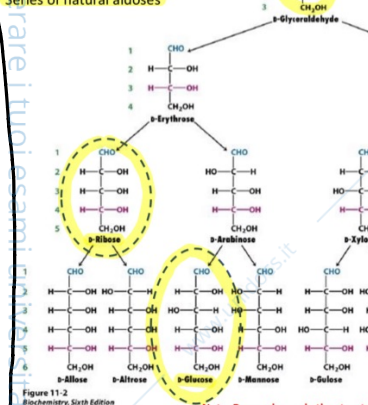
Most common monosaccharides are D.

Epimers

Two isomers differing in configuration at only one chiral center (i.e. two diastereoisomers).

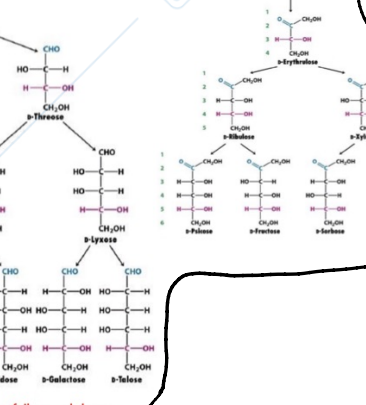


Monosaccharides: Series of natural aldoses



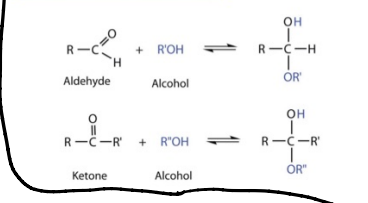
Note: Remember only the structures of ribose and glucose

Monosaccharides: Series of natural ketoses

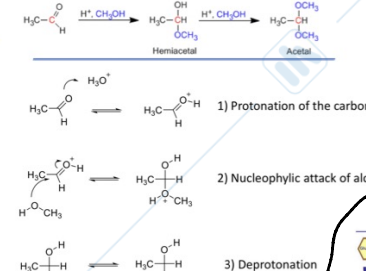


The cyclic structure of monosaccharides

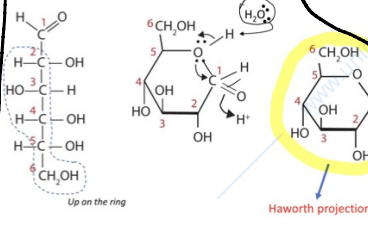
So far we have represented monosaccharides as linear molecules, but many of them also adopt cyclic structures. This conversion occurs because of the ability of aldehydes and ketones to react with alcohols.



Addition of 1 or 2 alcohols to a carbonyl group forms a hemiacetal, respectively, an acetal.

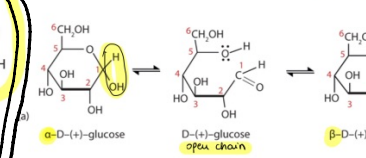


Formation of the cyclic structure of glucose



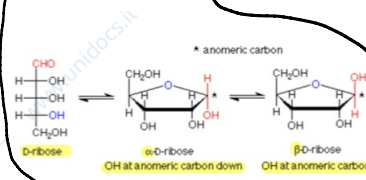
Formation of a hemiacetal. This reaction is catalyzed by acids. The reaction involves the OH on C5, to form a 6-membered ring.

The cyclization reaction produces two anomers: α and β DIASTEREOMERS

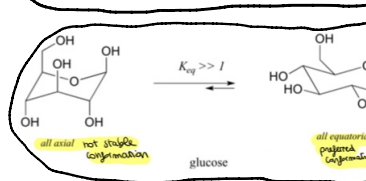
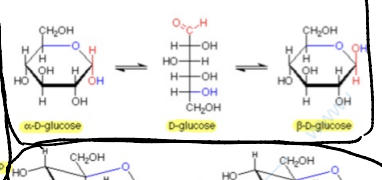


In water, at equilibrium, the mixture consists of about 36% α -D-glucose, 64% β -D-glucose, and less than 0.02% of the open-chain aldehyde form.

Ribose forms 5-membered rings

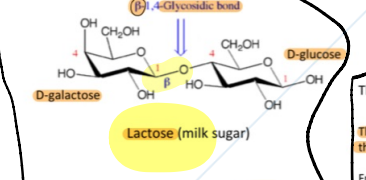


Chair conformation of 6-membered rings, α and β anomers



22-carbohydrates2

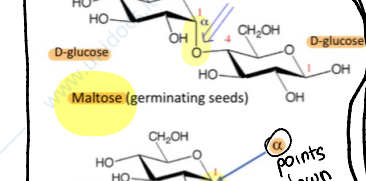
Disaccharides



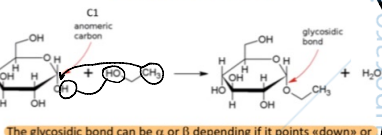
The glycosidic bond

The glycosidic bond is formed between the C1 of a monosaccharide (in the cyclic form) with the OH group of another molecule. From a chemical point of view, it is an acetal.

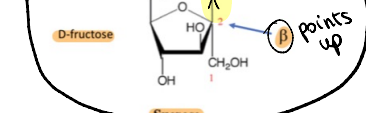
Maltose (germinating seeds)



Glycosidic bond between D-glucose and ethanol



Sucrose

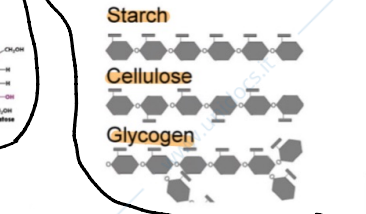


The glycosidic bond can be α or β depending if it points «down» or «up», respectively.

Polysaccharides

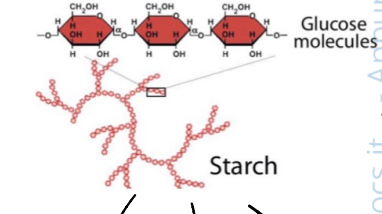
Large number of monosaccharide units bound together by glycosidic bonds.

Starch and glycogen are used as energy storage in plants and animals, respectively.

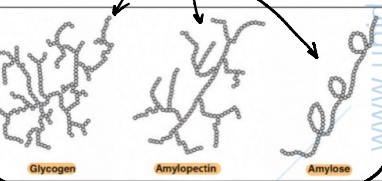
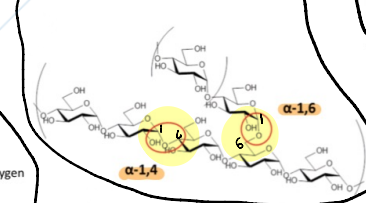


Starch and glycogen are formed by glucose units linked together mainly by α -1,4-glycosidic bonds.

Cellulose has structural functions in plants.



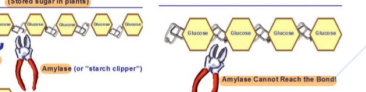
Ramifications are due to α -1,6 glycosidic bonds.



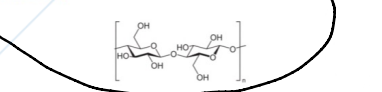
Starch. Amylose is the most abundant form of starch present in plants. Eating raw foods containing starches provides very little energy as the digestive system has a hard time breaking them down. Cooking breaks down the crystal structure of starches, making them much easier to break down in the human body.

Humans and animals store glucose energy from starches in the form of the very large molecule, glycogen. It has many branches that allow it to break down quickly when energy is needed by cells in the body. It is predominantly found in liver and muscle tissue in animals.

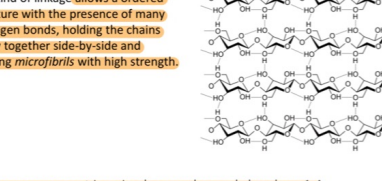
Starch (stored sugar in plants)



Cellulose

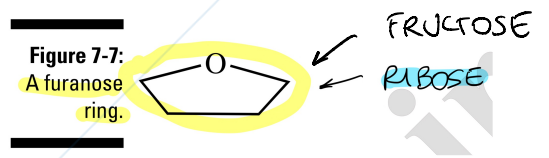
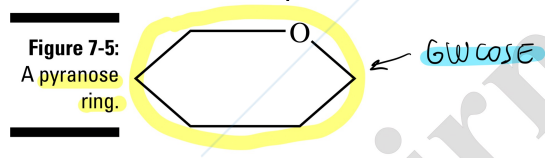
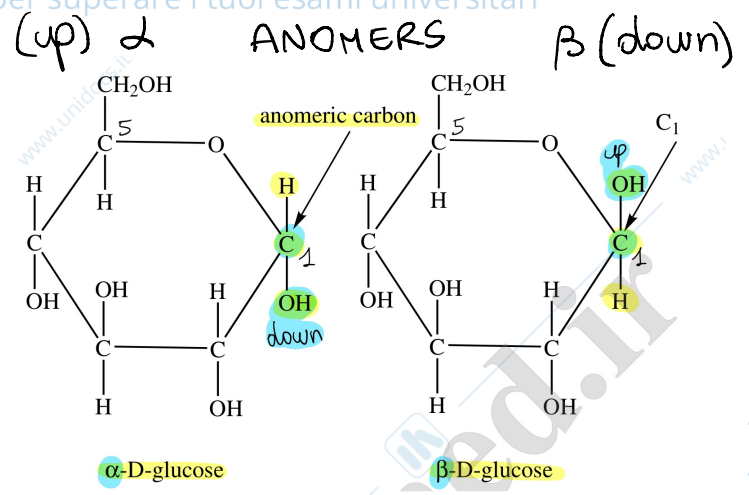
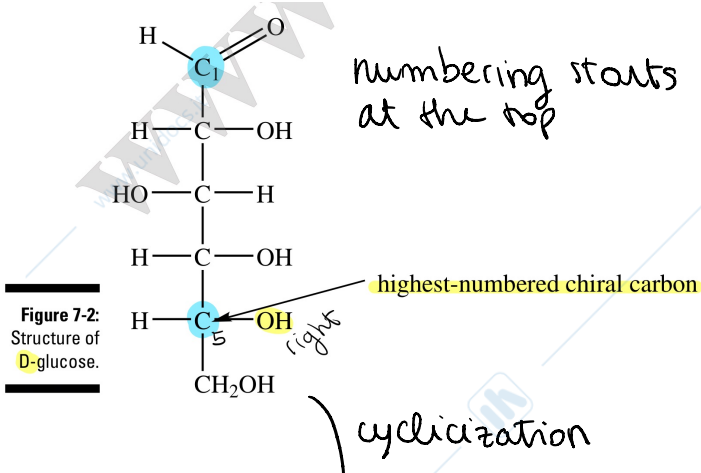


Cellulose is formed by glucose units linked together by β -1,4-glycosidic bonds

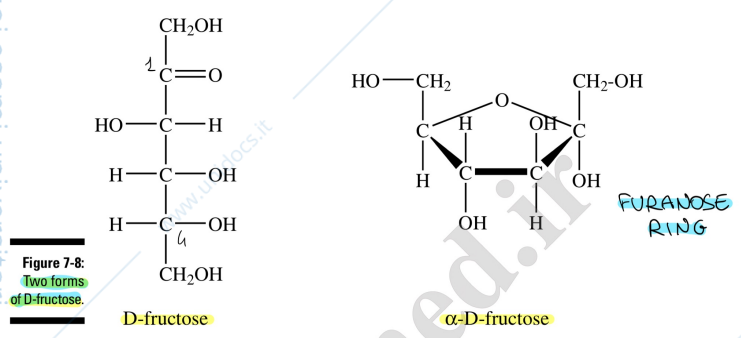
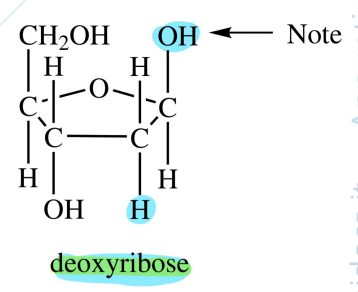
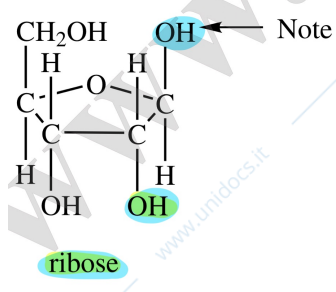
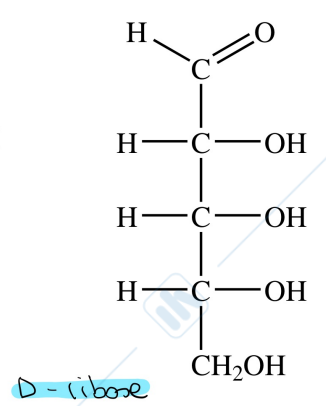
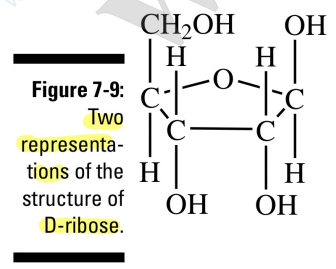
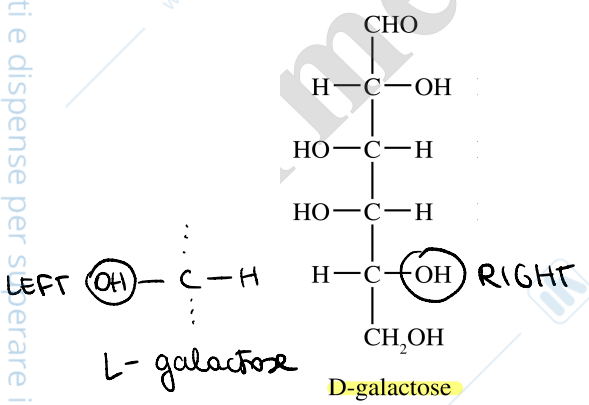


This kind of linkage allows an ordered structure with the presence of many hydrogen bonds, holding the chains firmly together side-by-side and forming microfibrils with high strength.

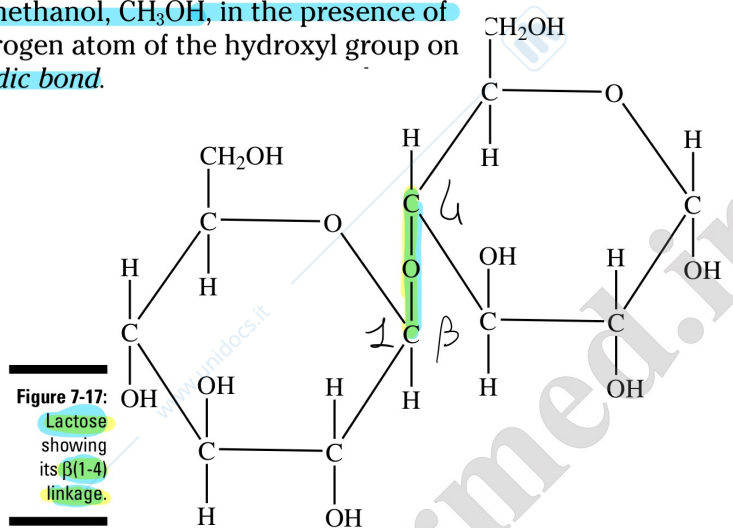
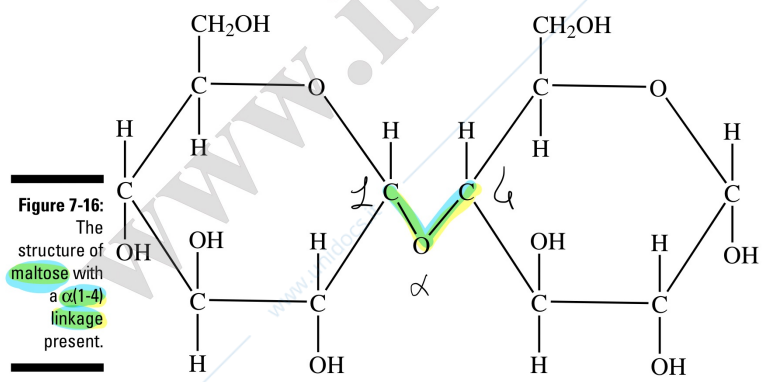
The enzyme present in animals α -amylase only breaks α -1,4 glycosidic bonds, so cellulose is not digested. Only herbivores host microbes that can digest cellulose.

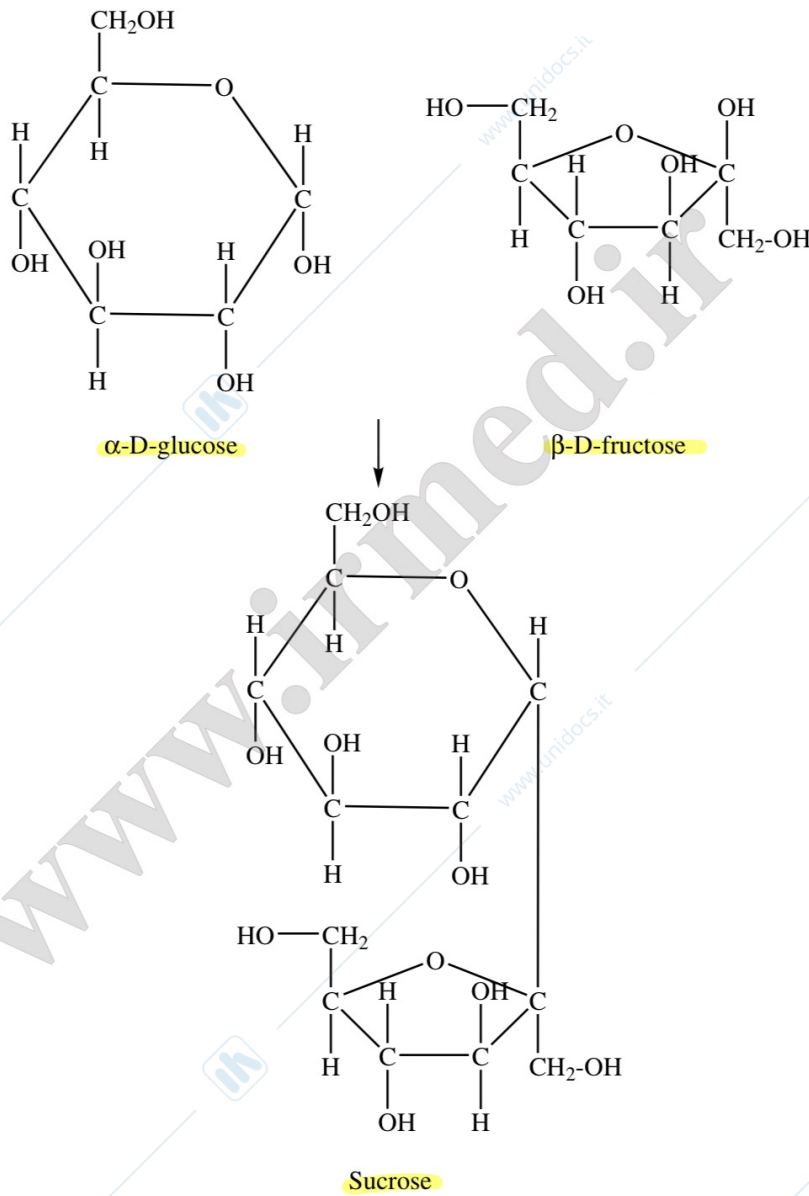


if the -OH on this carbon atom appears on the right, it's the D form of the monosaccharide; if it is on the left, it's the L form.



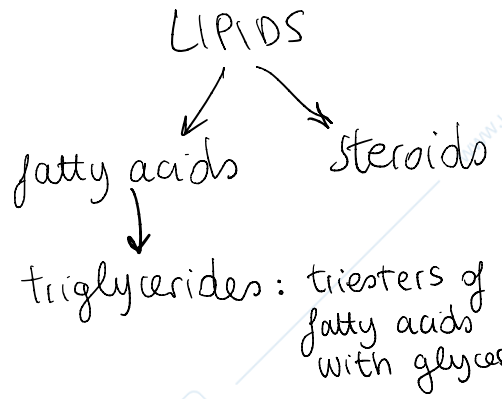
The reaction of a monosaccharide with methanol, CH₃OH, in the presence of hydrochloric acid, HCl, replaces the hydrogen atom of the hydroxyl group on C₁ with a methyl group, forming a glycosidic bond.



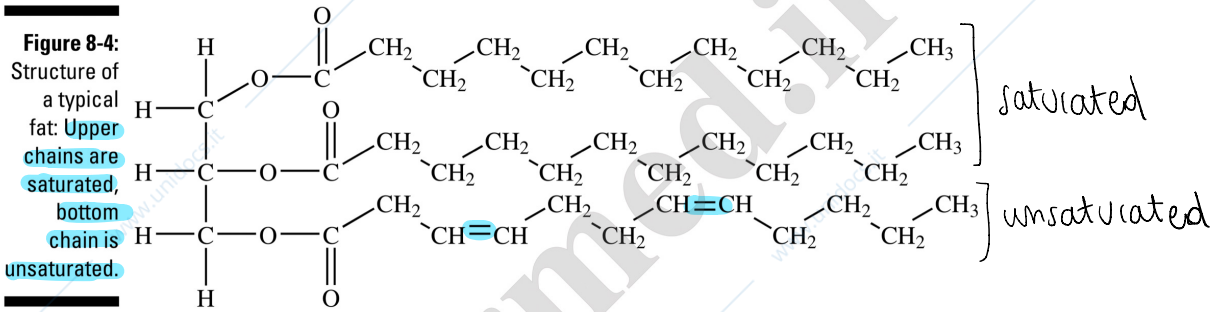
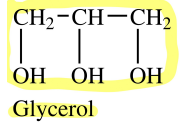


Starch is a polymer of α -D-glucose. There are three common types of starch: amylose, amylopectin, and glycogen. Amylose is the combination of $\alpha(1-4)$ glucose groups. Amylopectin, like amylose, has $\alpha(1-4)$ glucose linkages, but, in addition, it has $\alpha(1-6)$ branches.

Palmitic acid	$\text{CH}_3(\text{CH}_2)_{14}\text{COOH}$
Stearic acid	$\text{CH}_3(\text{CH}_2)_{16}\text{COOH}$
Oleic acid	$\text{CH}_3(\text{CH}_2)_7\text{CH}=\text{CH}(\text{CH}_2)_7\text{COOH}$
Linoleic acid	$\text{CH}_3(\text{CH}_2)_3(\text{CH}_2\text{CH}=\text{CH})_2(\text{CH}_2)_7\text{COOH}$
Linolenic acid	$\text{CH}_3(\text{CH}_2\text{CH}=\text{CH})_3(\text{CH}_2)_7\text{COOH}$

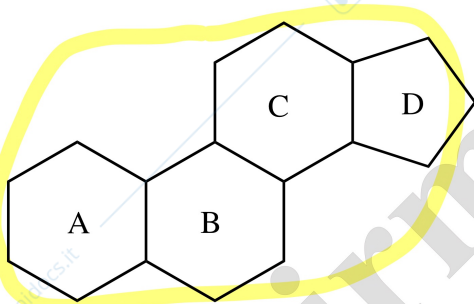


A **saturated fat** consists of fatty acids with no carbon-carbon double bonds. An **unsaturated fat** has a double bond while a **polyunsaturated fat** has multiple double bonds.



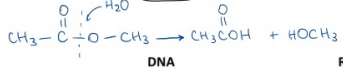
saponification reaction, hydrolysis of the ester groups in the presence of a base yields glycerol and the carboxylate ions of the three fatty acids.

Figure 8-11: Basic structure of a steroid.



Nucleic acids

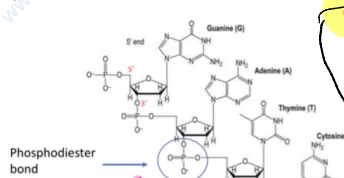
Controlled hydrolysis of nucleic acids yields three components.



DNA	RNA
Heterocyclic amines	Heterocyclic amines
2-deoxy-D-ribose	D-ribose
Phosphate ion (PO_4^{3-})	Phosphate ion (PO_4^{3-})

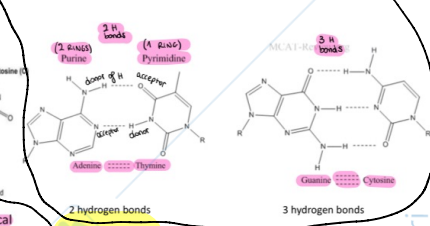
Structure of DNA

DNA is a **polynucleotide**, that is a biopolymer composed by several nucleotides. Nucleotides are linked by an ester bond between the phosphate of a nucleotide and the 3' OH group of another nucleotide. A sequence of polynucleotides is indicated by starting from the 5' end.

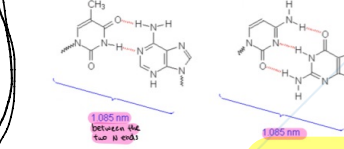


Base pairing

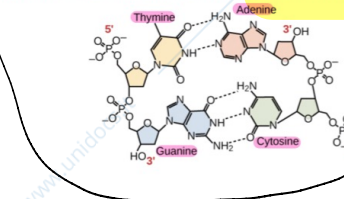
Adenine is able to pair with thymine, guanine with cytosine. As a general rule, a purine binds to a pyrimidine.



The dimensions of the A-T and G-C base pairs are identical



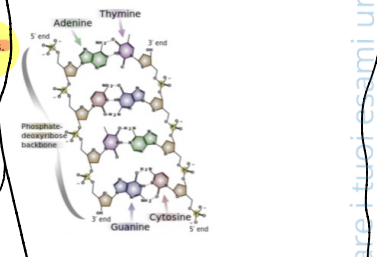
Two DNA single strands can bind if they have complementary bases.



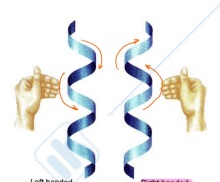
Double helix of DNA

A single polynucleotide strand is associated with another complementary strand forming a double strand.

The two strands are **antiparallel** (one is 5' → 3', the other is 3' → 5'), and are joined by hydrogen bonds between the nucleobases.



DNA helix is usually **right-handed**



Note that the handedness is retained when the helices are broken apart.

Nucleic acids contain information for organization and regulation of cellular functions.

DNA = deoxyribonucleic acid
RNA = ribonucleic acid

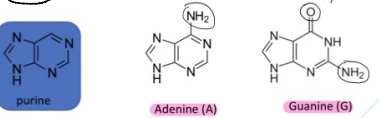
DNA is the repository of genetic information. RNA is the responsible for the transcription and for the synthesis of proteins.



Nucleobases

Pyrimidine bases

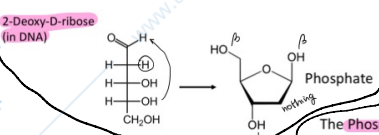
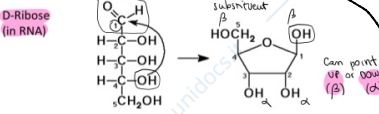
Purine bases



- Planar
- Weak bases.

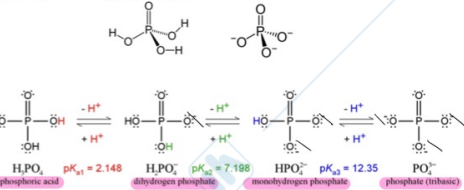
- Planar
- Weak bases.

At physiological pH, all nucleobases are neutral and are neither protonated nor deprotonated.



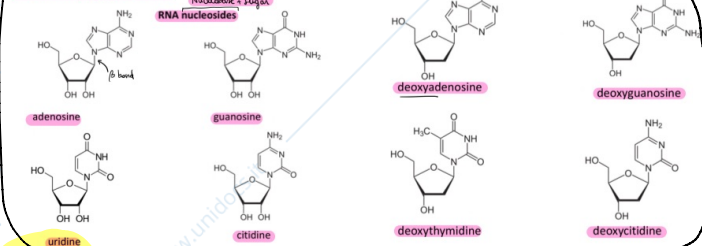
Numbering starts from the aldehyde group

The Phosphate group derives from phosphoric acid, a triprotic acid.



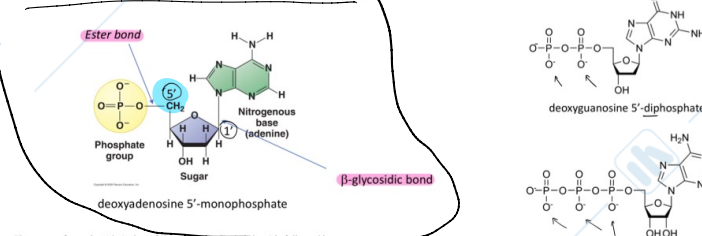
Nucleoside

Compound containing D-ribose (or 2-deoxyribose) bonded to a nucleobase by a β-glycosidic bond.



Nucleotide

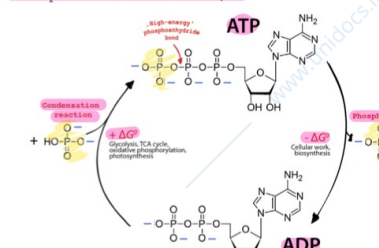
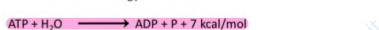
A nucleotide is a nucleoside bound to at least one phosphoric acid



The name of a nucleotide is that of the corresponding nucleoside followed by «monophosphate» and the position of the ester. The numbering of ribose is 1', 2', 3', 4', 5' to differentiate it from the numbering of the base.

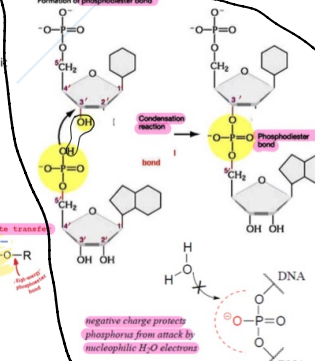
ATP - ADP system

ATP = adenosine triphosphate
ADP = adenosine diphosphate
Adenosine triphosphate (ATP) stores energy inside the cell and makes it available where energy is needed.



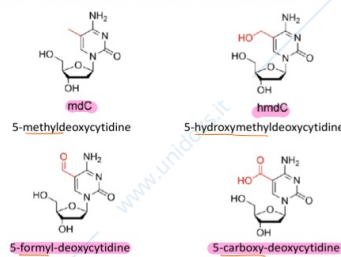
Polynucleotides

Formation of phosphodiester bond



Non-canonical bases in the genome

Probable regulatory elements of the transcription of genes.



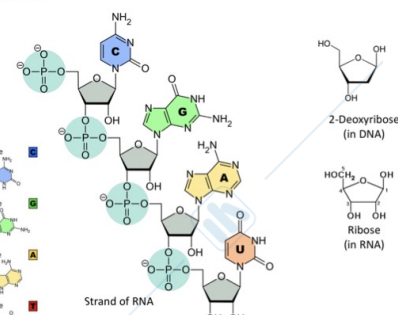
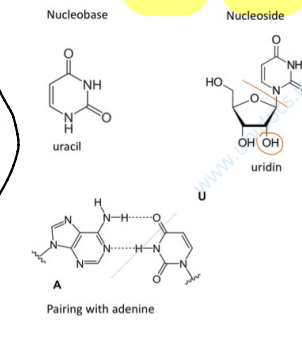
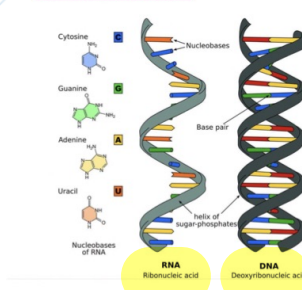
Ribonucleic acid (RNA)

It is a polynucleotide that transfer the information contained in DNA and performs the proteic synthesis.

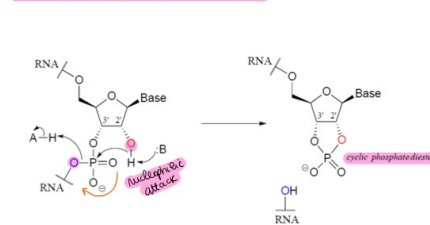
- Messenger RNA (mRNA) is the RNA that carries information from DNA to the ribosome (organelle that performs the proteic synthesis)
- Transfer RNA (tRNA) recognizes and transports a specific aminoacid
- Ribosomal RNA (rRNA) is the functional core of ribosomes.

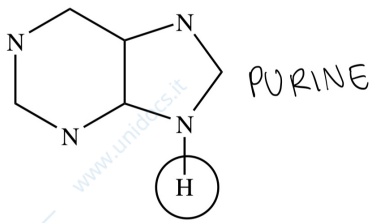
Differences between DNA and RNA

Ribose instead of 2'-deoxyribose
RNA is single stranded
Thymine is replaced with uracil

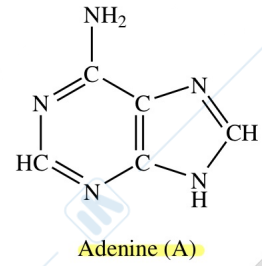
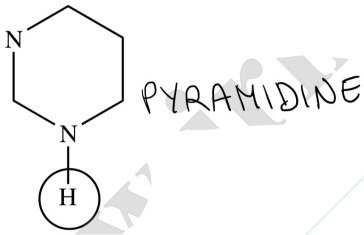


RNA nucleotides, unlike the deoxynucleotides of DNA, have a hydroxyl group at the neighboring 2' carbon. The 2' hydroxyl group is right next to the electrophilic phosphorus atom, poised in a good position to make a nucleophilic attack, breaking the RNA chain and forming a cyclic phosphate diester intermediate.

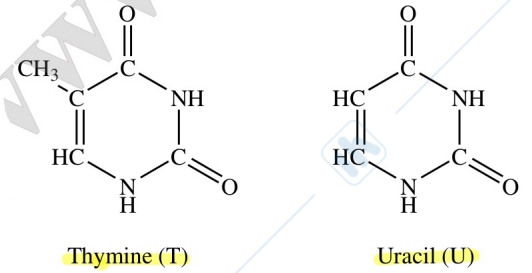
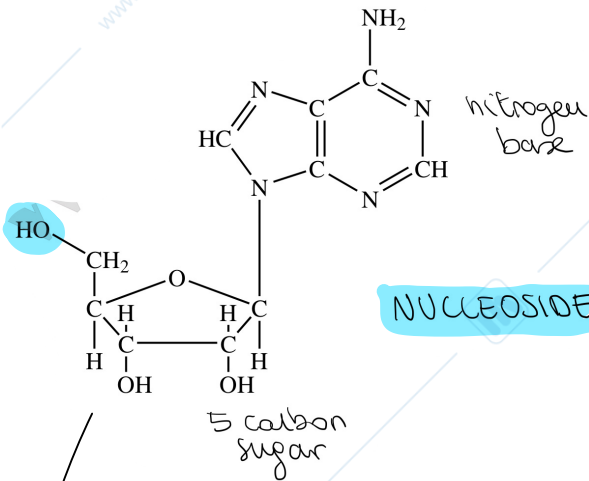
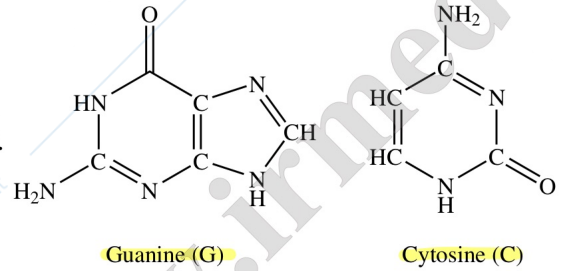




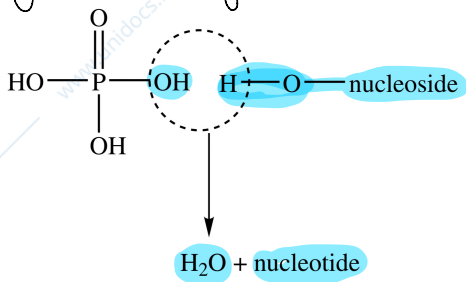
- ✓ The purines (adenine and guanine), composed of two fused rings incorporating two nitrogen atoms in each ring and
- ✓ The pyrimidines (cytosine, thymine, and uracil), composed of a single ring with two nitrogen atoms in the ring structure



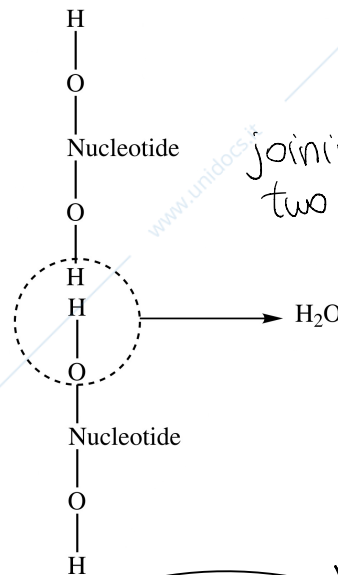
Adenine (A), guanine (G), and cytosine (C) occur in both DNA and RNA. Thymine (T) is only found in DNA, whereas uracil (U) only occurs in RNA.



formation of a nucleotide



joining of two nucleotides



NUCLEIC ACID

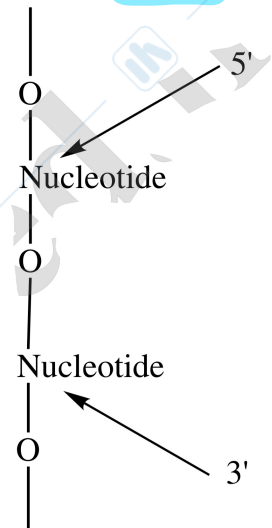
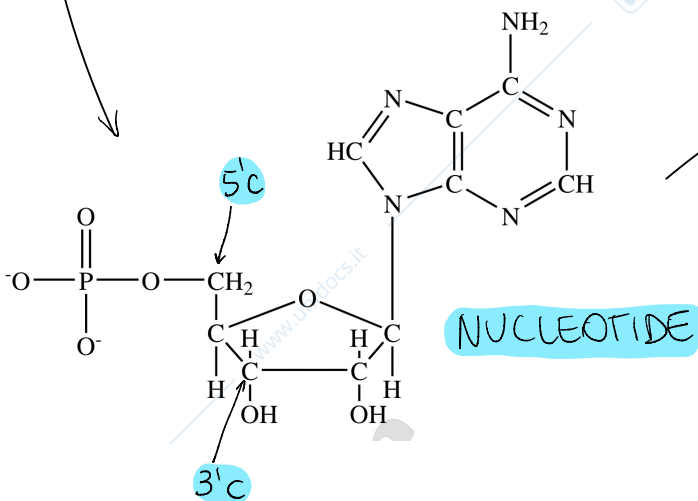


Figure 9-12:
Hydrogen bonds (dotted lines) form between **guanine** (right) and **cytosine** (left).

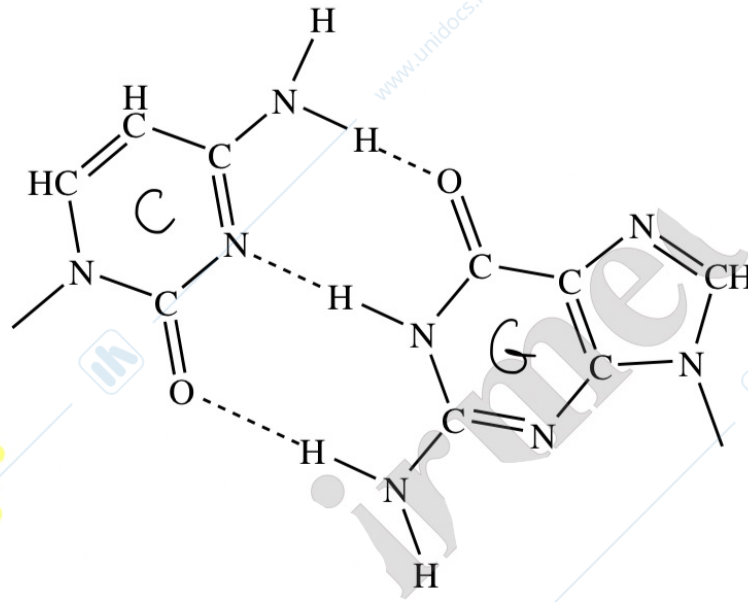
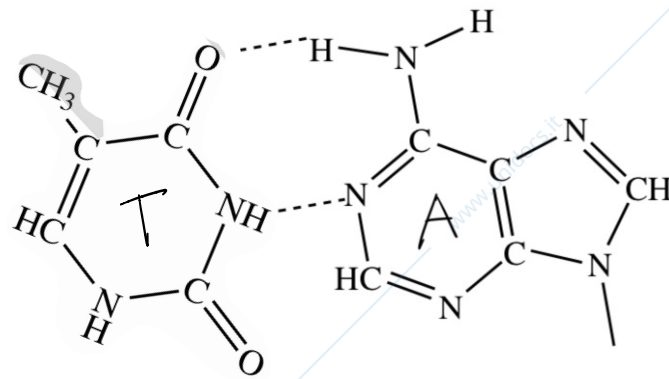
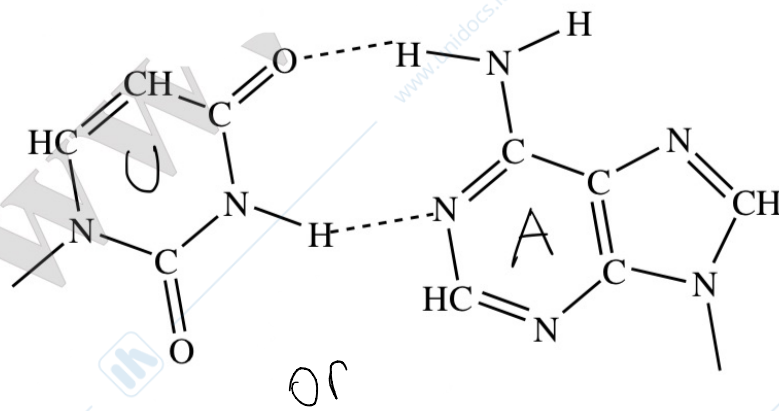


Figure 9-13:
Hydrogen bonds (dotted lines) form between **adenine** (right) and **uracil** (left).

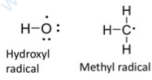


Free radicals

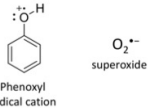
What are free radicals

Free radicals (or, more simply, radicals) are molecule or ions having unpaired electrons in the valence shell.

Neutral radicals (no charge)

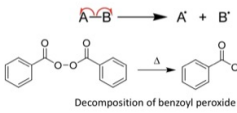


Radical ions (positive or negative charge)

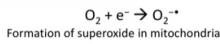


Radical formation

1) Homolytic cleavage of a bond, due to: heat and radiations



2) Capture or loss of one electron:



3) Bond cleavage coupled to an electron transfer:

Formation of HO• in the presence of iron II and hydroperoxides (Fenton reaction)



Reaction of radicals

Radicals are very reactive and have a short life in solution, because they lack the octet. Radicals undergo to two kinds of reactions:

- With themselves (radical-radical or decay reactions)
- With closed-shell molecules

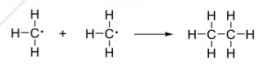


Radical-radical reactions

These reactions are extremely fast and cause the disappearance of radicals.

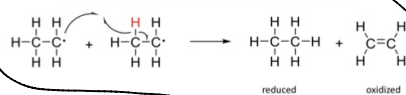
1) recombination

Two radicals bind together



2) disproportionation

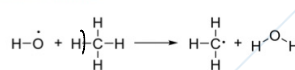
One radical is reduced while another one is oxidized.



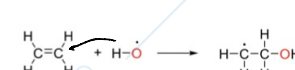
Reaction with closed-shell molecules

These reactions may be fast or slow, and transform one radical into another one.

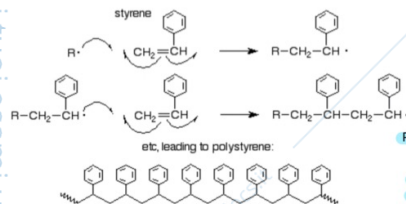
1) H atom abstraction



2) Addition to double bonds



Radical polymerization



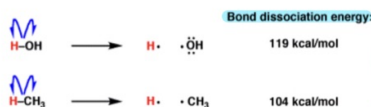
Persistent radicals

In very specific and limited cases, radicals can survive for long time, because recombination or disproportionation are not occurring. This happens in the case of radicals having bulky substituents.

Radical stability

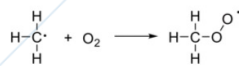
Radical stability describes how easily a radical can be formed starting from a closed-shell molecule. Also, this parameter is (roughly) inversely proportional to the tendency of a radical to react with closed-shell molecules.

Stability of neutral radicals is inversely proportional to the Bond Dissociation Energy (BDE), that is the energy required to break a bond (usually, the bond with H).



Peroxy radicals

Carbon-centered radicals quickly react with oxygen to form peroxy radicals.



With this reaction, O₂ is incorporated into organic molecules. This is the reason why radicals mediate oxidation reaction.

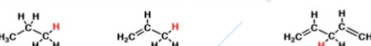
Radical stability increases in the order methyl < primary < secondary < tertiary



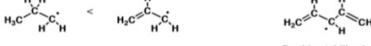
Strongest bond, Weakest bond



Radical stability increases if stabilized by resonance

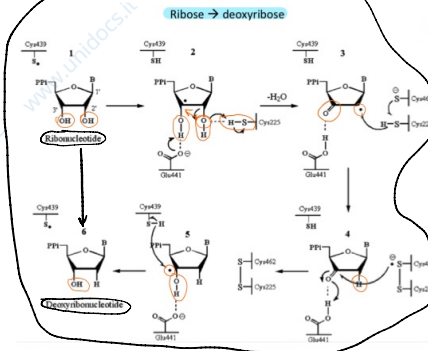


Strongest bond, Weakest bond



- Radical formation
- Reactions of radicals
 - recombination (polymerization)
 - H-atom transfer
- Stability and persistency
- Reactive oxygen species
- Lipid peroxidation
- Antioxidants

Radical enzymes: Ribonucleotide reductase



Radical damage to biomolecules

Oxidation of membranes (lipid peroxidation) is mediated by peroxy radicals (ROO•).

Damage to DNA is mediated by HO•.

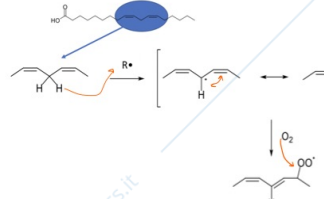
ROS: «Reactive Oxygen Species»

ROS is a term used in biochemistry that includes all species (radical or not) containing oxygen that are associated with radical damage.

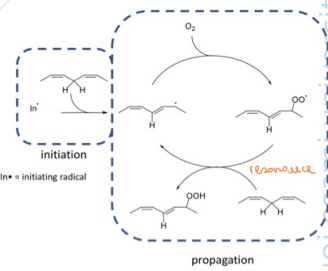
- radical: Hydroxyl radical: HO•
Superoxide radical: HOO• / OO•
Alkylperoxy radical: AlkOO•
- Non-radical: Hydrogen peroxide: H₂O₂

Lipid peroxidation

Cellular membranes contains polyunsaturated lipids. The bis-allylic moiety is very reactive toward radicals, because it can form a radical that is stabilized by resonance on both double bonds. Further reaction with oxygen yields peroxy radicals.



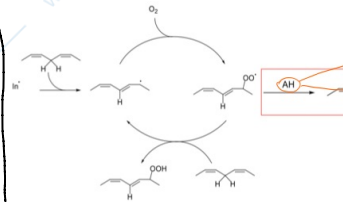
Lipid peroxidation radical cycle



Radical oxidation of lipids is a chain reaction. One initiation event destroys many lipid molecules.

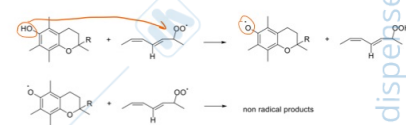
Chain-breaking antioxidants

Lipid peroxidation is blocked by molecules able to intercept peroxy radicals – the chain-breaking antioxidants (AH).

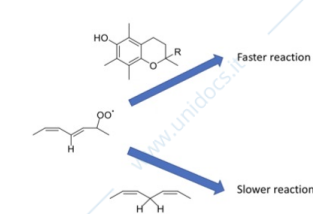


Alpha-tocopherol is a typical antioxidant

α-Tocopherol is the main component of Vitamin E. It is able to trap two peroxy radicals by the following reactions.



The antioxidant ability of α-tocopherol is due to the fact that it reacts with peroxy radical much faster than unsaturated lipids do.



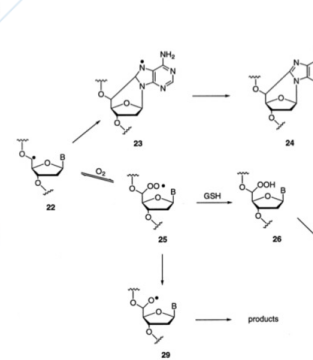
Radical damage to DNA

DNA is more resistant to radical attack than unsaturated lipids. Only the very unstable (i.e. very reactive) hydroxyl radical is able to attack it.

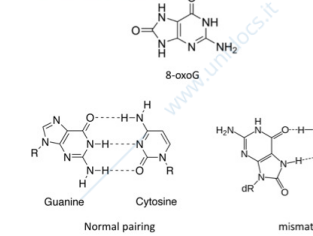


Scheme of the different possible oxidative attacks by OH• on DNA nucleobase (T) or backbone sugar moiety.

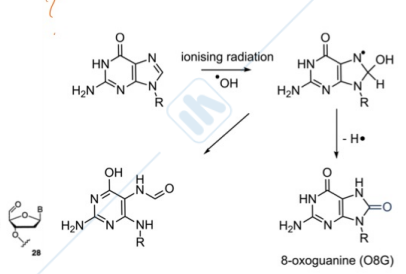
Damage to the sugar



The most common oxidative lesion is 8-oxoguanine. It causes pairing mismatch



Damage to the bases



Being so reactive, hydroxyl radicals (HO•) must be formed in proximity of DNA. These radicals are formed by the Fenton reaction between H₂O₂ produced by mitochondria and Fe²⁺ or Cu⁺ present in the nucleus, or by radiations.

Hydroxyl radicals are not efficiently blocked by antioxidants, because HO• reacts with DNA and with antioxidants at similar rates.

- The best defence against DNA radical damage is the prevention of HO• radicals formation:
- Limit H₂O₂ / ROOH build up inside cells.
 - Remove excess Fe²⁺ and Cu⁺ ions.
 - Avoid ionizing radiations.