



**2024  
SUMMARY**

**BIOLOGY 101**

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# Chapter 6:

★ Free Energy Change ( $\Delta G$ )  $\rightarrow$  The energy that can do work

difference  
Final - Initial

$$\Delta G = G_{\text{products}} - G_{\text{reactants}}$$

$$\Delta G = \Delta H - T\Delta S$$

Enthalpy  
Total energy

Entropy  
Temperature in Kelvin (K)

It is a measure of instability (tendency to become more stable)

★ Equilibrium  $\rightarrow$  Maximum stability state  $\rightarrow$  lowest G

Spontaneous Reaction

$\Rightarrow$  Exergonic Reaction

$\Rightarrow$  Catabolic Reaction

$\rightarrow$  break down molecules

$\Rightarrow$   $\Delta G = \ominus$  Negative

$\Rightarrow$  Release energy

$\Rightarrow$  Decrease in free energy ( $\downarrow G$ )

$\Rightarrow$  Increase the Stability ( $\uparrow$  stability)

$\Rightarrow$  Toward equilibrium

Non-Spontaneous Reaction

$\Rightarrow$  Endergonic Reaction

$\Rightarrow$  Anabolic Reaction

$\rightarrow$  Building molecules

$\Rightarrow$   $\Delta G = \oplus$  Positive

$\Rightarrow$  Absorbs, uses, consumes, stores energy

$\Rightarrow$  Increase in free energy ( $\uparrow G$ )

$\Rightarrow$  Decrease the Stability ( $\downarrow$  stability)

$\Rightarrow$  Away from equilibrium

Unstable  
 $\uparrow G$

The magnitude of  $\Delta G$  represents the amount of energy

$\Delta G = \oplus$   $\Delta G = \ominus$

The Reverse Reaction of an exergonic Reaction must be endergonic

Stable  
 $\downarrow G$

☆ The release of energy during an exergonic reaction is due to the **conversion** from high energy (unstable) state into a less energy (stable) state **NET** due to breaking bonds itself

☆ Energy stored in bonds  $\rightsquigarrow$  represents the **potential energy** released

☆ Living cells **Never** reaches **equilibrium**  $\rightsquigarrow$  because they are **open systems**  
 $\hookrightarrow$  If a cell reached equilibrium  $\rightsquigarrow$  dead

☆ Types of work:

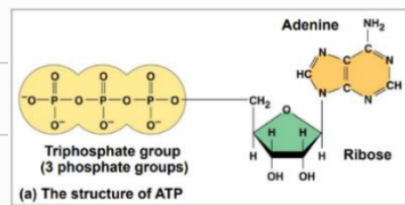
- **Chemical work**  $\rightsquigarrow$  Anabolic Reactions (synthesis of glutamine, building polymers)
- **Transport work**  $\rightsquigarrow$  Active transport (against gradient, Pumping)
- **Mechanical work**  $\rightsquigarrow$  Muscle contraction, beating of cilia, moving chromosomes

☆ **Energy coupling**  $\rightsquigarrow$  **Exergonic** + **Endergonic**  $\overset{\text{using}}{\rightsquigarrow}$   $\overset{\text{to drive}}{\rightsquigarrow}$  **Exergonic**  $\overset{\text{the Overall}}{=}$  **Exergonic**  
 $\rightsquigarrow$  It is mediated by **ATP**

RNA nucleotide  $\leftarrow$  Ribose

Adenine

3 Phosphate



ATP is used for energy coupling because:

Release a great amount of energy  $\rightsquigarrow$  due to its **instability**  $\rightsquigarrow$  due to **repulsion** between **negative charges** in the (P) tail

$\rightsquigarrow$  Release energy by breaking the bond of the **terminal (P)** forming **ADP + P<sub>i</sub>**

usually transferred to another

molecule (Phosphorylation) forming a **phosphorylated intermediate**  
**high energy**  $\leftarrow$  **Unstable**  $\leftarrow$  **Reactive**

☆ **ATP hydrolysis**  $\rightsquigarrow$   $\Delta G = -7.3$  kcal/mol (under **standard** conditions)  
 $\rightsquigarrow$   $\Delta G = -13$  kcal/mol (under **cellular** conditions)

☆ ATP is recycled by adding phosphate to ADP  $\rightsquigarrow$  **endergonic**  
 The energy for this reaction is acquired from **catabolic reactions** such as **break down glucose**

☆ Catalyst  $\rightsquigarrow$  chemical agent that speeds up reactions, without being consumed

$\Rightarrow$  Enzymes  $\rightsquigarrow$  Catalytic proteins  $\rightsquigarrow$  ends with -ase

Q) A small amount of enzymes catalyzing a huge number of reactions, Why?

Enzymes are not changed or consumed during reactions (released in its original form) so they can be used repeatedly

☆ Activation energy (EA)  $\rightsquigarrow$  Initial energy needed to start a reaction

$\Rightarrow$  Supplied as thermal energy  $\rightsquigarrow$  accelerating reactants & collide more until reaching transition state

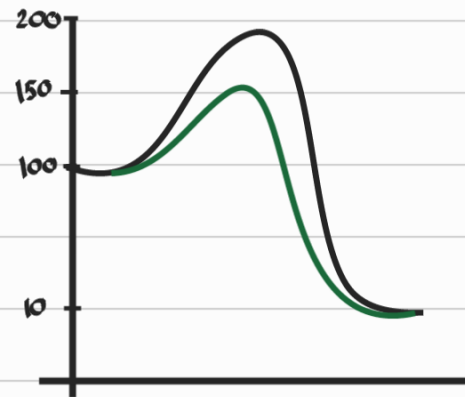
$\hookrightarrow$  Unstable (high energy) conditions of reactants

It is the energy between reactants & transition state

☆ How do enzymes work?

$\Rightarrow$  lowering EA barrier  $\rightsquigarrow$  reaching transition state easier

$\Rightarrow$  Enzymes do not affect  $\Delta G$



substrate  $\rightsquigarrow$  Reactants in enzyme catalyzed reaction

enzyme-substrate complex  $\rightsquigarrow$  enzyme binding a substrate

Active site  $\rightsquigarrow$  Region (pocket, groove) on the surface of the enzyme

$\hookrightarrow$  It binds to the substrate & catalyze the reaction

Induced fit  $\rightsquigarrow$  It is the tight binding after the initial contact between substrate & enzyme

☆ How does the active site lower EA barrier??

□ Orienting substrate correctly

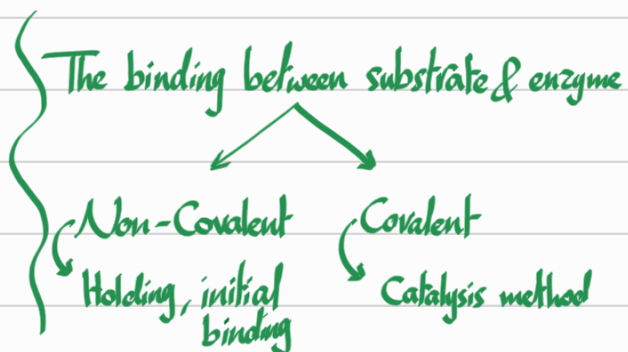
□ straining bonds

□ Provide favorable microenvironment

□ Covalent bonding with substrate

☆ Enzyme activity is affected by:

- 1- Environmental factors  $\rightsquigarrow$  Optimal conditions
- 2- Chemicals  $\rightsquigarrow$  most active shape





- ☆ Human enzyme has optimal temperature =  $35^{\circ}\text{C}$  -  $40^{\circ}\text{C}$ , optimal pH = 6 - 8
  - ⇒ optimal pH for pepsin (in the stomach) ⇒ very low about 2
  - ⇒ optimal pH for Trypsin (in the intestine) ⇒ about 8
- ☆ optimal temperature for thermophilic bacteria =  $70^{\circ}\text{C}$  and more

Note: as temperature increases ⇒ more collide ⇒ the rate increases until reaching optimal temperature ⇒ if temperature increased ⇒ the enzyme denatures and become inactive

- ☆ As the concentration of the substrate increases ⇒ the rate of the reaction increases until all enzyme molecules become engaged ⇒ saturated
- ☆ To increase the rate of the reaction when the enzyme is saturated we have to increase the amount of the enzyme

- ☆ Cofactors ⇒ Non-protein enzyme helpers
  - ↳ Inorganic (such as metals)
  - ↳ Organic ⇒ Coenzymes (such as vitamins)

- ☆ Cells regulate their metabolic pathways by:
  - 1) Switching on or of the genes that encodes a specific enzyme
  - 2) Regulating the activity of enzymes

### ☆ Enzymes Inhibitors

#### Competitive inhibitors

- ⇒ Binds the active site
- ⇒ Blocks the entry of substrate to the active site
- ⇒ Can be overcome by increasing substrate conc

#### Non-competitive inhibitors

- ⇒ Bind another site
- ⇒ Changes the shape making the active site less effective

If the binding is: weak ⇒ Reversible inhibition, covalent ⇒ irreversible inhibition

☆ Allosteric regulation → For enzymes composed from more than 1 subunit  
↳ the binding of a regulatory molecule to a site affects the function on other sites  
↳ activators (stimulate), inhibitors (inhibit)  
↳ A single regulatory molecule can affect all subunits

☆ Cooperativity

↳ It is a type of allosteric regulation that can amplify enzyme activity  
↳ one substrate bind to the active site of 1 subunit → increase the affinity for the substrate of all subunits  
↳ It is explained by hemoglobin (transport protein not an enzyme)

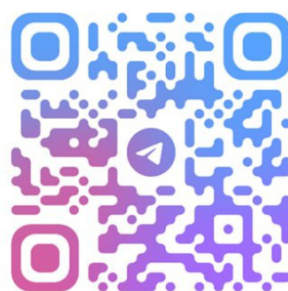
☆ Feedback inhibition

↳ The end product in a pathway inhibits an early enzyme  
↳ prevents wasting chemical resources & synthesis more products than needed

Notes:

- 1) Some enzymes are inserted in membranes
- 2) ATP acts as inhibitors for catabolic pathways  
ADP acts as activators for catabolic pathways
- 3) The variety between enzymes is due to mutations
- 4)  $\Delta G$  for the hydrolysis of glucose = -686
- 5) Dinoflagellates are marine organisms that convert chemical energy to light by bioluminescence

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# Chapter 10:

Potential energy is due to the arrangement of  $e^-$

Catabolic pathways  $\rightsquigarrow$  Break down complex molecules  $\rightsquigarrow$  Release energy

☆ Energy is released due to the transfer & rearrangement of  $e^-$   $\rightsquigarrow$  become simple, more stable with less energy

Some of this energy will be stored in ATP and the rest is dissipated (lost) as heat

☆ Redox Reactions  $\rightsquigarrow$  Reactions involve the transfer of  $e^-$

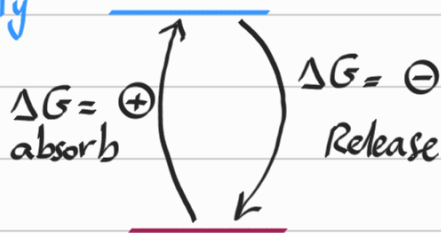
↳ Oxidation: lose electrons ( $e^-$  donor)  $\rightsquigarrow$  charge becomes more positive

↳ Reduction: gain electrons ( $e^-$  acceptor)  $\rightsquigarrow$  charge becomes less positive

☆ Oxidizing agent  $\rightsquigarrow$  get reduced (gain  $e^-$ )

☆ Reducing agent  $\rightsquigarrow$  get oxidized (lose  $e^-$ )

Unstable, low electronegativity  
High energy, Complex



Stable, High electronegativity  
low energy, Simple

Notes:

$\Rightarrow$  No oxidation without reduction

$\Rightarrow$  Some redox reactions don't involve the actual transfer of  $e^-$   
Such as  $O_2 + \text{Methane}$

☆ Oxygen has a very high electronegativity  $\rightsquigarrow$  strong pulling  $\rightsquigarrow$  ↑ Reduction  
So, the strongest oxidizing agent

↳ Any transfer of electrons toward oxygen  $\rightsquigarrow$  Release energy

☆ Organic molecules acting as fuels have abundance of H atoms

↳ because H has a very low electronegativity  $\rightsquigarrow$  high-energy electrons (hilltop electrons)

## ☆ Catabolic pathways:

- ⇒ Aerobic Respiration → Uses  $O_2$  & ETC → The most efficient
- ⇒ Anaerobic Respiration → Uses ETC without  $O_2$
- ⇒ Fermentation → **doesn't** use  $O_2$  and ETC

$NAD^+$  (Nicotinamide Adenine Dinucleotide): A coenzyme,  $e^-$  carrier, derivative of vitamin niacin

### ☆ Dehydrogenase:

↳ Removes a pair of H atoms ( $2e^-$ ,  $2H^+$ ) from the substrate

☆  $NAD^+$  accepts  $2e^-$  &  $1H^+$  forming NADH

☆ The other proton is released  $H^+$  ion into the solution

↳ acts with dehydrogenases  
Carry  $e^-$  so represent stored energy taped to make ATP

### Note:

$NAD^+$  → Oxidized form  
 $NADH$  → Reduced form

☆  $NADH$  transfers the  $2e^-$  to the ETC

☆ ETC transfers the  $2e^-$  to Oxygen

☆ Oxygen accepts  $2H^+$  forming  $H_2O$

↳ Electron transport chain:  
⇒ consists mainly of proteins  
⇒ Eukaryotes: inner mitochondrial membrane

Prokaryotes: Plasma membrane

⇒ Transferring electrons from the

High energy end  
↓ Electronegativity

Top  
↓

low energy end  
↑ Electronegativity

Bottom

## ☆ Aerobic Respiration Eukaryotes & Prokaryotes

3 stages:

1) Glycolysis → break down glucose into 2 pyruvate

2) TCA cycle → completes glucose break down

3) Oxidative phosphorylation → synthesizes most (90%) of ATP



# ☆ Glycolysis ☆

- Occurs in the Cytoplasm
- Occurs in the presence of  $O_2$  or not
- 10 steps divided into 2 phases:

Energy investment phase	Energy pay-off phase
spend 2 ATP	Repay (forming) 4 ATP
to break glucose into	Rearranging the atoms
<u>Two 3-C sugars</u>	forming 2 pyruvate
↳ G3P and DHAP	↳ 2 G3P
	isomerase

Notes:

⇒ **Oxidative phosphorylation**: adding  $P_i$  to ADP forming ATP, occurs **with** ETC

⇒ **Substrate level phosphorylation**: adding phosphate from the substrate to ADP forming ATP, occurs **without** ETC

Very Important Notes on glycolysis:

- 1) Steps that consume ATP = 1 & 3 / steps that form ATP = 7 & 10
- 2) Step number 6 forms 2 NADH / step number 9 forms 2  $H_2O$
- 3) **Phosphofructokinase**: The enzyme that catalyze the 3<sup>rd</sup> step → transfers phosphate to (fructose 6-phosphate) forming (fructose 1,6-bisphosphate)
- 4) **Enolase**: The enzyme that catalyze the 9<sup>th</sup> step → forming (=) in 2-phosphoglycerate converting it to Phosphoenolpyruvate (PEP) → a molecule with high energy
- 5) **G3P** = Glyceraldehyde 3-phosphate, **DHAP** = Dihydroxy Acetone Phosphate

The net products of Glycolysis per glucose molecule:

**2 ATP, 2 NADH, 2  $H_2O$ , 2 pyruvate molecules**

**No Release of  $CO_2$**

## ☆ Pyruvate oxidation

- Occurs in the mitochondria (eukaryotes) but in Cytosol (prokaryotes)
  - Pyruvate enters the mitochondria by **active transport**
- Steps: 1) Release  $CO_2$  2) Forming NADH 3) attaching CoA forming **acetyl CoA**
- **Coenzyme A (CoA)**: Sulfur containing compound, derivative of B vitamin, has high energy

The net products of Pyruvate Oxidation per:

Pyruvate molecule:  $1 \text{CO}_2$ ,  $1 \text{NADH}$ ,  $1 \text{acetyl CoA}$

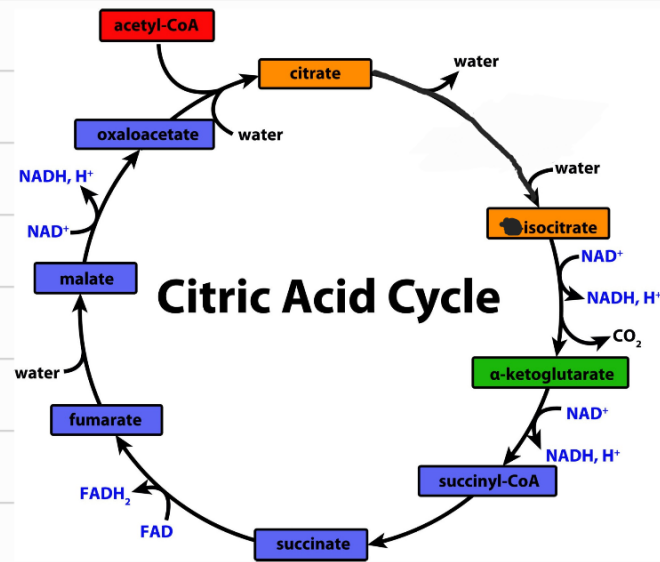
Glucose molecule:  $2 \text{CO}_2$ ,  $2 \text{NADH}$ ,  $2 \text{acetyl CoA}$

No ATP  
Synthesis

## ☆ TCA cycle ☆

- Called Citric acid cycle, kreb's cycle and Tricarboxylic acid (TCA) cycle

CIA Sent Soldiers  
For My Office



## Notes:

- 1) TCA cycle release  $2 \text{CO}_2$  in steps 3 and 4
  - 2) TCA cycle forms  $3 \text{NADH}$  in steps 3, 4 and 8
  - 3) TCA cycle forms  $1 \text{ATP}$  (or  $\text{GTP}$ ) in step 5
  - 4) TCA cycle forms  $1 \text{FADH}_2$  in step 6
  - 5) Citrate is converted to isocitrate by Removing  $\text{H}_2\text{O}$  then adding  $\text{H}_2\text{O}$  again
  - 6) Fumarate is converted to malate by adding  $\text{H}_2\text{O}$
- All the enzymes of TCA cycle are located in the mitochondrial matrix  
Except the enzyme of the 6<sup>th</sup> step (inner mitochondrial membrane)

The net products of TCA cycle per:

pyruvate molecule, per cycle:  $2 \text{CO}_2$ ,  $3 \text{NADH}$ ,  $1 \text{FADH}_2$ ,  $1 \text{ATP}$  (or  $\text{GTP}$ )

glucose molecule:  $4 \text{CO}_2$ ,  $6 \text{NADH}$ ,  $2 \text{FADH}_2$ ,  $2 \text{ATP}$  (or  $\text{GTP}$ )

# ☆ Oxidative Phosphorylation ☆

ETC

Prosthetic group: **Non-protein** components

↳ mainly proteins

⇒ Source of  $e^-$  for ETC → **NADH & FADH<sub>2</sub>** accepts  $2e^-$  and  $2H^+$

⇒ **NADH** transfers  $2e^-$  to **Complex I**

**Complex I (FMN → FeS) → CoQ → Complex III and IV (cytochromes) → O<sub>2</sub>**

⇒ **FADH<sub>2</sub>** transfers  $2e^-$  to **Complex II**

**Complex II (FeS) → CoQ → Complex III and IV (cytochromes) → O<sub>2</sub>**

**Flavoprotein:** The first molecule in ETC, Its prosthetic group is **Flavin Mono Nucleotide** <sup>FMN</sup>

**Fe-S:** Iron-sulfur protein, Its prosthetic groups are **iron & sulfur**

**Coenzyme Q, Ubiquinone (CoQ):** **hydrophobic, mobile**, the only member that isn't protein

**Cytochromes (Cyt):** proteins in Complexes III and IV, the last one (before O<sub>2</sub>) is **Cyt a<sub>3</sub>**

their prosthetic group is **heme**

↳ contains iron

☆ ETC is important to transfer electrons step by step & release energy in a more manageable way (Not in an explosive way)

**ATP synthase** → It is an enzyme that makes ATP (from ADP + P<sub>i</sub>)

↳ In the **inner mitochondrial membrane** (eukaryotes)

and **plasma membrane** (prokaryotes)

The energy released due to the transfer of  $e^-$  along ETC is used to

**pump proton (H<sup>+</sup>)** from the matrix to the **intermembranous space**

forming a proton gradient → then H<sup>+</sup> return back to the matrix through

ATP synthase → moving and **spinning** the rotator → catalyzing **ATP synthesis**

↳ indirect use of energy



**Chemiosmosis, Proton-motive Force**  $\rightsquigarrow$  It is the use of proton gradient to do a work (such as ATP synthesis)

### Notes:

- 1) **34%** of the energy in glucose is used to make ATP  $\rightsquigarrow$  the rest is lost as heat  
 $\Rightarrow$  **Uncoupling proteins**: proteins in the inner mitochondrial membrane  $\rightsquigarrow$  decrease the efficiency of ATP generation to **form heat** to maintain body temperature
- 2) The total ATP synthesized is about = **32 ATP**  
 $\Rightarrow$  **2 (glycolysis) + 2 (TCA) + 28 (ETC + Chemiosmosis) = 32**  
**Substrate level phosphorylation + Oxidative phosphorylation**
- 3) The transfer of electrons release energy enough to pump:  
 $\Rightarrow$  **NADH** (from mitochondria)  $\rightsquigarrow$  **10 H<sup>+</sup>**  $\rightsquigarrow$  forming **2.5 ATP**  
 $\Rightarrow$  **NADH** (from cytosol, glycolysis)  $\rightsquigarrow$  forming **1.5 ATP**  
less because of energy used to **actively transport** NADH to the mitochondria  
 $\Rightarrow$  **FADH<sub>2</sub>**  $\rightsquigarrow$  **6 H<sup>+</sup>**  $\rightsquigarrow$  forming **1.5 ATP**  
less because it transfer its e<sup>-</sup> to a **lower energy level** (complex II not I)

☆ The flow of \_\_\_\_\_ during aerobic respiration:

- 1) e<sup>-</sup>  $\rightsquigarrow$  **Glucose**  $\rightsquigarrow$  **NADH, FADH<sub>2</sub>**  $\rightsquigarrow$  **ETC**  $\rightsquigarrow$  **Oxygen**
- 2) energy  $\rightsquigarrow$  **Glucose**  $\rightsquigarrow$  **NADH, FADH<sub>2</sub>**  $\rightsquigarrow$  **ETC**  $\rightsquigarrow$  **Chemiosmosis**  $\rightsquigarrow$  **ATP**

☆ **Anaerobic Respiration**  $\rightsquigarrow$  in certain prokaryotes

- $\Rightarrow$  It uses **ETC** but **doesn't use Oxygen**  $\rightsquigarrow$  It use another electronegative molecule with **less electronegativity** (efficiency) than O<sub>2</sub> (such as SO<sup>-2</sup>)
- $\Rightarrow$  Marin bacteria uses **SO<sup>-2</sup>**  $\rightsquigarrow$  so the by product is **H<sub>2</sub>S** not H<sub>2</sub>O  
(H<sub>2</sub>S has **rotten egg odor**)



# ☆ Fermentation

- ⇒ In cellular respiration → make ATP by substrate level and oxidative phosphorylation, and  $\text{NAD}^+$  is recycled by ETC
- ⇒ Fermentation → make ATP only by substrate level phosphorylation and  $\text{NAD}^+$  is recycled by pyruvate or one of its derivatives
- ⇒ Fermentation = glycolysis +  $\text{NAD}^+$  recycling (regeneration)

## Alcohol Fermentation

- ☆ Pyruvate converted to ethanol by 2 steps:
  - 1) Release  $\text{CO}_2$  forming acetaldehyde
  - 2) Reducing acetaldehyde by NADH forming Ethanol +  $\text{NAD}^+$
- ☆ used by yeast (Fungus) & bacteria
- ☆ used in winemaking, brewing and baking

## Lactic acid Fermentation

- ☆ Pyruvate is converted directly to lactate
- ☆ No Release of  $\text{CO}_2$
- ☆ Used by fungi, bacteria & muscles
- ☆ Used in making yogurt and cheese
- ☆ Muscles use lactic acid fermentation when  $\text{O}_2$  is scarce (not enough) in strenuous exercise → causing lactate accumulation → which transported to the liver to regenerate pyruvate

- ☆ In fermentation → pyruvate or acetaldehyde are the last  $e^-$  acceptors
- ☆ In aerobic respiration →  $\text{O}_2$  is the last  $e^-$  acceptor
- ☆ In anaerobic respiration → Electronegative molecule is the last  $e^-$  acceptor

Organisms are classified into:

- 1) **Obligate anaerobe** → only Anaerobic Respiration and fermentation  
↳ Can't survive in the presence of  $\text{O}_2$  ( $\text{O}_2$  is toxic for them)
- 2) **Obligate aerobe** → only Aerobic Respiration (such as vertebrate's brain)
- 3) **Facultative anaerobe** → Both Fermentation + Respiration (such as yeast, some bacteria & muscle cells)

☆ Glycolysis & TCA cycle have many intersections with other catabolic and anabolic pathways

## Catabolic

- 1) Starch and glycogen  $\rightsquigarrow$  broken down into glucose  $\rightsquigarrow$  enters glycolysis
- 2) Proteins  $\rightsquigarrow$  broken down into amino acid  $\rightsquigarrow$  Removing their amine group by deamination  $\rightsquigarrow$  then enter TCA or glycolysis
- 3) Fats  $\rightsquigarrow$  Glycerol  $\rightsquigarrow$  converted to G3P  $\rightsquigarrow$  glycolysis intermediate  
 $\rightsquigarrow$  Fatty acids  $\rightsquigarrow$  broken down by beta oxidation to 2C fragments which are converted to acetyl CoA  $\rightsquigarrow$  enter TCA

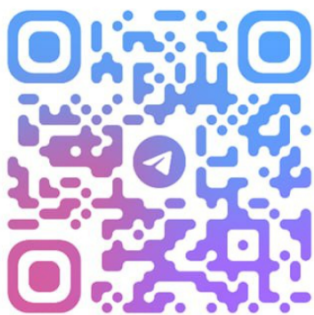
## Anabolic

- 1) Pyruvate can be used to make glucose
- 2) Acetyl CoA can be used to make fats
- 3) DHAP can be used to make fats
- 4) We can synthesis half of the 20 amino acid  
the rest are essential (only from diet)

## Regulation of Respiration:

phosphofructokinase (step 3 glycolysis)  $\rightsquigarrow$  the pacemaker of respiration  
Inhibited by ATP and citrate, activated by AMP and ADP

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