# **Instructor: Zhang Rong**

Department of Biochemistry & Molecular Biology
Sichuan University

# CHAPTER 6

# **BIOLOGICAL OXIDATION**

#### **Definition:**

The process that substance is oxidazed into carbon dioxide  $(CO_2)$  and water $(H_2O)$ , and releases energy in organism.

#### **Characteristics of biological oxidation:**

- 1. The reaction is generally occurred in the condition that PH approaches neutral and the temperature is  $37^{\circ}$  C.
- $2_{\sim}$  The produced energy is progressively released and can be storied .
  - 3. The reaction requires water.
  - 4. Use the decarboxylation way to produce CO2.
  - 5. Use the dehydrogenation way produce H<sub>2</sub>O.

#### **SECTION ONE**

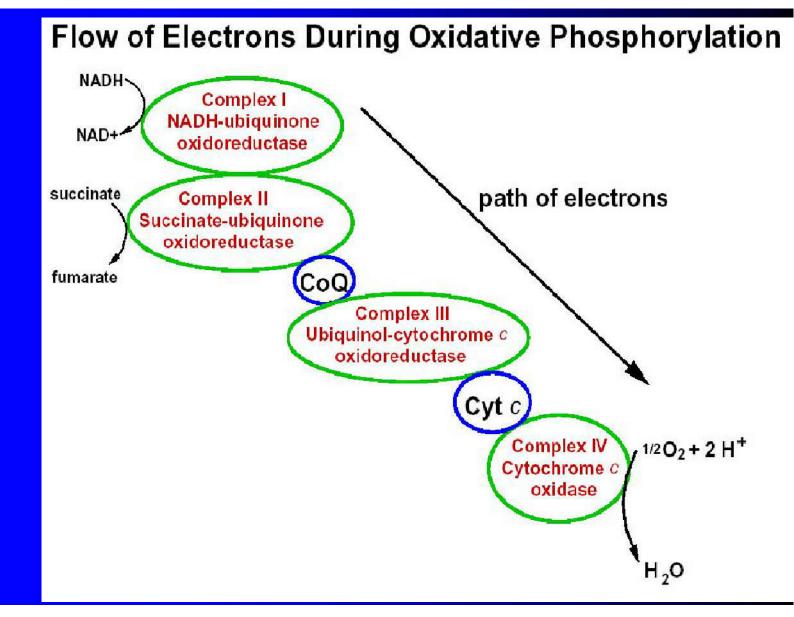
#### **OXIDATION SYSTERM OF ATP PRODUCED**

## 1, Respiratory chain

**Definition:** The chain reaction which is formed by the arrangement of a regular order of the enzymes and coenzymes on the inner membrane of mitochondria and is closely associated with the utilization of oxygen by cells is called respiratory chain.

### (1) Components of the respiratory chain

Four complex that still have the ability to transform electrons can be separated while using generally chemical methods



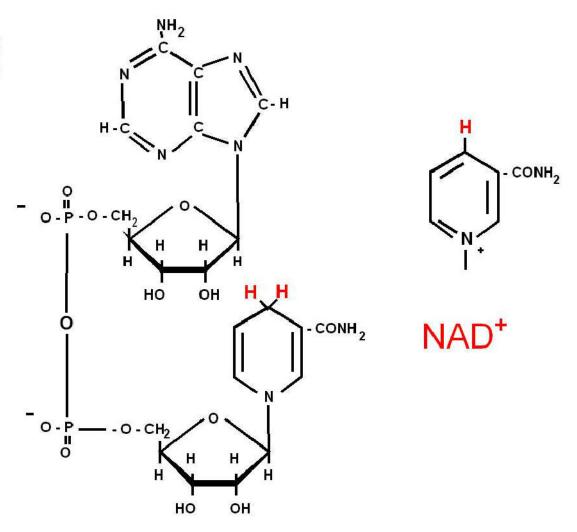
### Structure and function of electron transmitter:

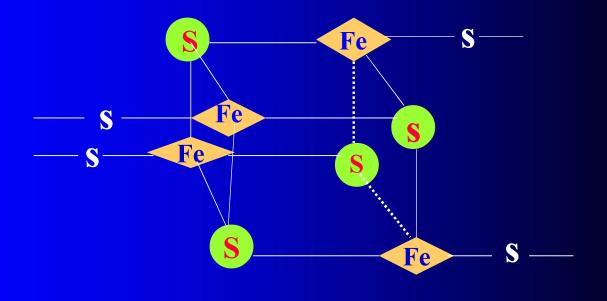
**□** Complex I: NADH-ubiquinone oxidoreductase Pass the reducing equivalents of NADH to Q

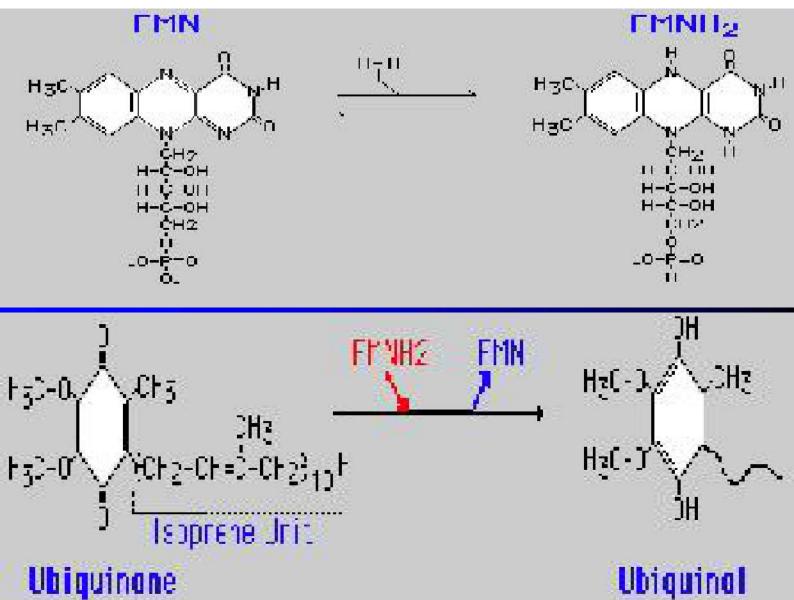
NADH FMN Q
(FeS)

- (1) Flavoprotein: The prosthetic group is FMN
- (2) Iron-sulfur protein (FeS)

# **NADH**







Complex II: succinate-ubiquinone oxidoreductase

Pass the electrons from succinate to ubiquinone

succinate FAD Q (FeS)

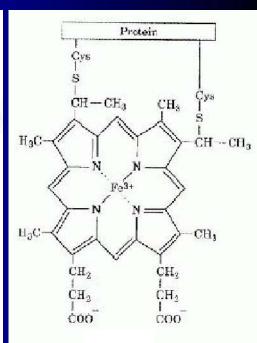
- (1) Flavoprotein: Prosthetic group is FAD
- (2) Iron-sulfur protein (FeS)
- (3) Cytochrome<sub>560</sub>

# Riboflavin (B<sub>2</sub>)

- **Complex** III: Pass the electrons from Q to  $CytC_1$ .
  - (1) Iron-sulfur protein (FeS)
  - (2) Cytochrome: Cytb<sub>562</sub>, Cytb<sub>566</sub>, Cytc<sub>1</sub>

Q Cyt b<sub>562</sub> Cyt b<sub>566</sub> Cyt c<sub>1</sub>

$$\begin{array}{c} \text{CH}_3 \\ \text{CH}_2 + \text{CH}_2 - \text{CH} = \text{C} - \text{CH}_2 +_{\overline{q}} \text{H} \\ \text{HO} - \text{CH} & \text{CH}_3 \\ \text{CH}_3 \\ \text{CH}_4 & \text{CH}_2 \\ \text{CH}_2 & \text{CH}_2 \\ \text{CH}_2 & \text{CH}_2 \\ \text{CH}_2 & \text{CH}_2 \\ \text{COO} & \text{COO} \\ \end{array}$$



# Complex IV: Cytochrome c oxidase Pass the electrons from Cytc to oxygen.

Cyt c

Cyt a

Cyt a<sub>3</sub>

 $1/2O_2$ 

This complex contain Cyta and Cyta3

Cytochrome C is the only soluble cytochrome and, togethe with uquinone (Q), it seems to be a more mobile component of the respiratory chain.

# (2) The sequence of respiratorychain:

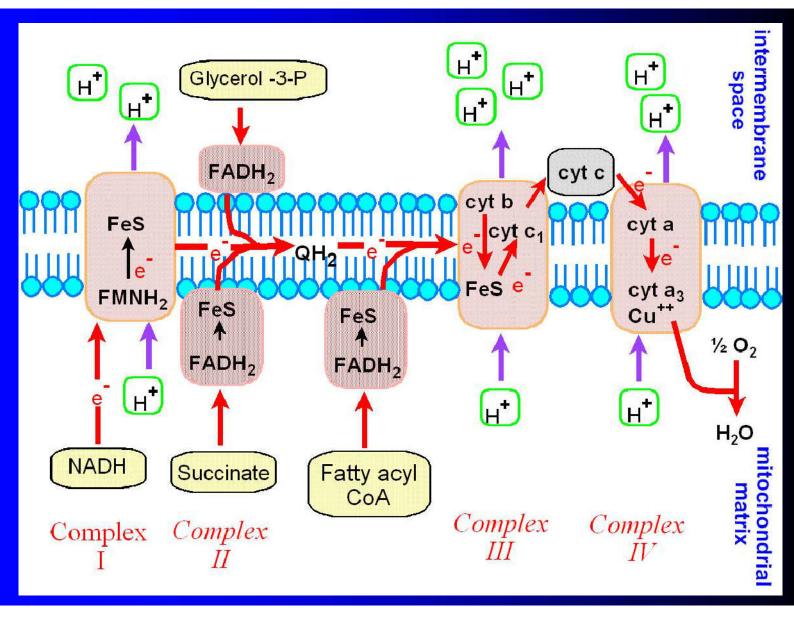
A. Determine the redox potential of all the electrons transmitters of the respiratory chain  $(\Delta E^{0'})$ .

# Some redox potentials of special interest in mammalian oxidation systems.

Oxidation-reduction couple	$\triangle \mathbf{E}^{0/}(\mathbf{V})$
$2H^+ + 2e \longrightarrow H_2$	-0.42
$NAD^+ + 2H^+ + 2e \longrightarrow NADH + H^+$	-0.32
$FMN + 2H^+ + 2e \longrightarrow FMNH_2$	-0.12
$FAD + 2H + 2e \longrightarrow FADH_2$	-0.06
$CoQ + 2H^+ + 2e \longrightarrow CoQ$	0.04
Cytb(Fe <sup>3+</sup> ) + e $\longrightarrow$ Cytb(Fe <sup>2+</sup> )	0.07
$CytC_1(Fe^{3+}) + e \longrightarrow Cyt C_1(Fe^{2+})$	0.22
$CytC(Fe^{3+}) + e \longrightarrow CytC(Fe^{2+})$	0.25
Cyta (Fe <sup>3+</sup> ) + e $\longrightarrow$ Cyta (Fe <sup>2+</sup> )	0.29
$Cyta_3(Fe^{3+}) + e \longrightarrow Cyta_3(Fe^{2+})$	0.39
$1/2O_2 + 2H^+ + 2e \longrightarrow H_2O$	0.82

- B. Add some respiratory chain inhibitors which the affect locus are afferent.
- C. Analyse with the specially absorption spectrums properties of the electrons transmitter.
- D. Rebuild the electrons transmitter in external
- The two respiratory chain which their arrangements are known till now:

succinate		FAD ()					
NADH	FMN	CoQ	Cytb	$\mathbf{c_1}$	c	aa <sub>3</sub>	O



# 2. Oxidation phosphorylation

Definition: Redox reaction of substrates producing reducing equivalents(H or electrons), There are funneled into the respiratory chain to their final reaction with oxygen to form water, and released free energy that used to synthesize ATP from ADP and Pi, in this way, oxidation is closely coupled to phosphorylation.

- (1) The location of for oxidative phosphorylation
  - ①, P/O ratio:

Definition: When substrates are oxidated producing equivalents (H or electrons), these are funneled into the respiratory chain ,their final reaction with oxygen to form water. The mol of the Pi that is consumed when one gram of mol oxygen  $(O_2)$  is consumed is called P/O ratio.

### **Substrates**

oxygen

**Equivalents** FAD

NADH FMN CoQ Cytb c<sub>1</sub> c aa<sub>3</sub> O<sub>2</sub>

 $H_2O$ 

3ADP + Pi 3ATP P/O Ratio = 3

## P/O ratio is measured in extra-somatic mitochondria:

Substrate	Electron transfer channel					P/O ratio	ATP quality
ß-羟丁酸	NADH	FMN	CoQ	Cyt	$\mathbf{O_2}$	2.4-2.8	3
琥珀酸	FAD	CoQ	Cyt	$O_2$		1.7	2
抗坏血酸	Cytc	Cytaa <sub>3</sub>	$O_2$			0.88	1
细胞色素C	Cyt	taa <sub>3</sub>	$O_2$			0.61-0.68	1

 $\overline{\text{FADH}_2}$ 

NADH FMN CoQ Cytb Cytc<sub>1</sub> Cytc Cytaa3 O2

# ②、Inferring from the free energy released in different phases:

It is clear that, an approximate 30.5KJ/mol free energy is needed to from one gram molecular of ATP, which is equal to 0.2v of  $E^{\text{O}'}$  respiratory chain. Therefore we can infer the location the ATP produced from the numerical value of the potential difference between two neighbor carriers on the respiratory chain.

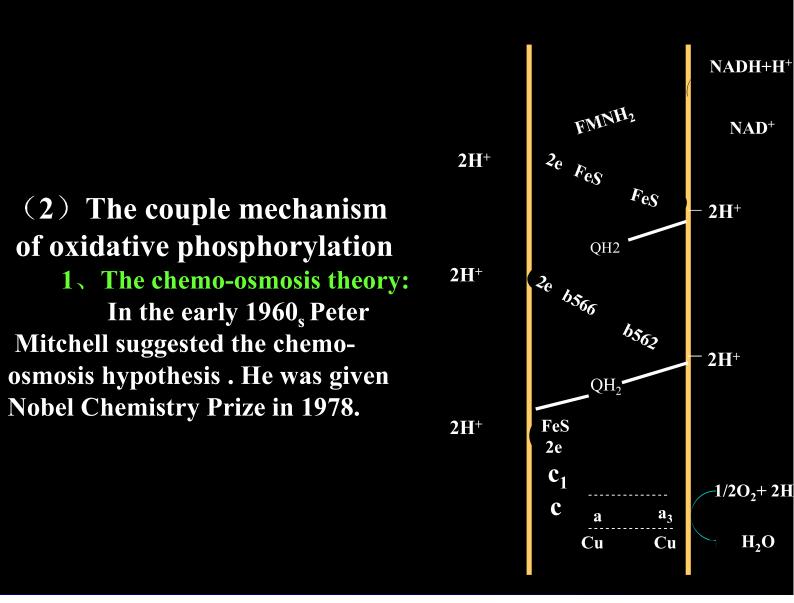
Then  $\Delta E^{o/}$  is used to calculate  $\Delta G^{o/}$ :

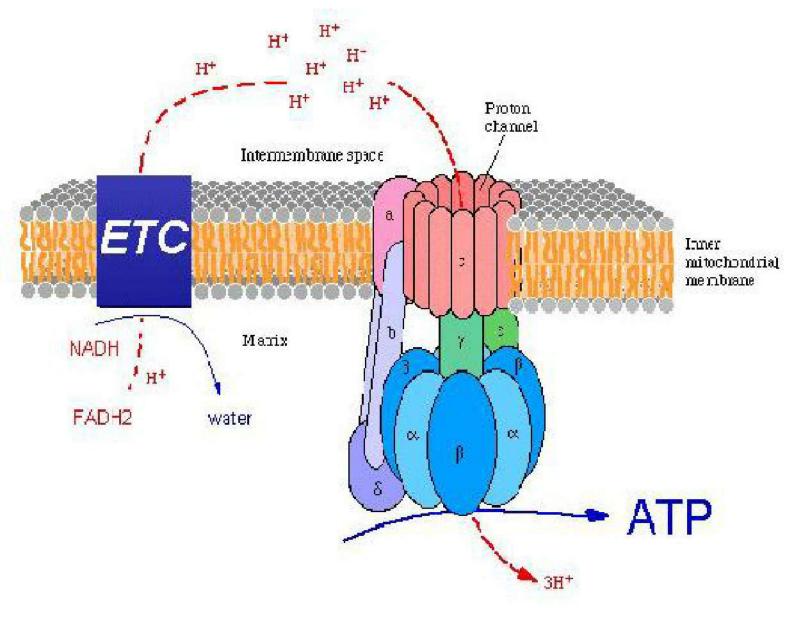
$$\triangle G^{O/} = - nF \triangle E^{O/}$$
  
F=96.5KJ/mol. V

 $FADH_2$ 

#### 0.1v=20KJ

NADH	FMN	CoQ		Cytb	Cytc	Cytaa <sub>3</sub>	$O_2$
EO/	0.36v		EO/	0.21v		$\mathbf{E}^{\mathbf{O}/}$	<b>0.53</b> v
$G^{0/}$	69.5KJ		$G^{O/}$	40.5KJ		$G^{O/}$	102.3KJ
ADP Pi			ADP Pi			ADP Pi	
A	ГР		A	ТР		AT	





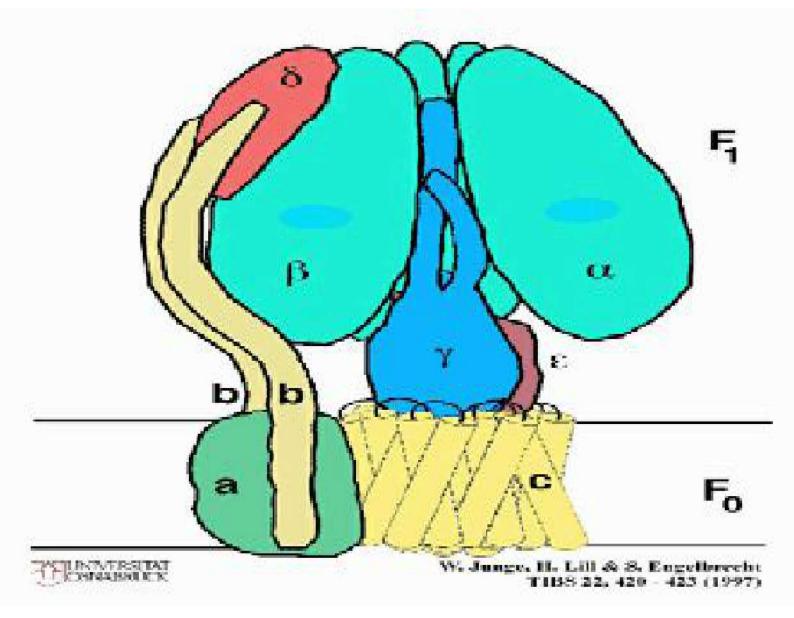
### 2. ATP synthetase:

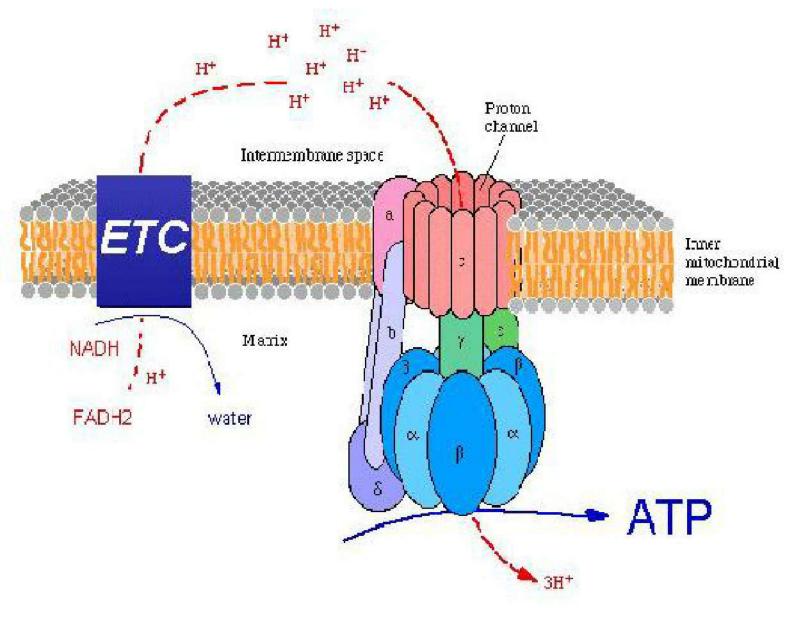
#### (1) structure of ATP synthetase:

ATP synthetase is a large membrane-protein complex . The ATP-synthesizing enzyme complex of the inner mitochondrial membrane has two major components,  $F_1$  and  $F_0$ 

#### $\triangleright$ $F_1$ : $F_1$ dissolves in water.

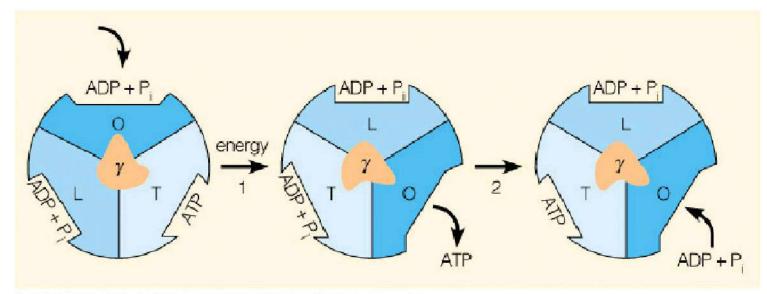
A tight binding site for ATP, apparently identical to the catalytic site, is located on each  $\beta$  subunit, or perhaps between each  $\beta$  and its associated  $\alpha$  subunit.





 $ightharpoonup F_0$ :  $F_0$  is liposoluble the  $F_0$  complex consists of a subunit, b subunit, c subunit and oligomycin-sensitivity-conferring protein(OSCP). Its structure is  $ab_2c_n$ . The primary role of  $F_0$  is transmitting the energy produced by the proton gradient to  $F_1$ .

### (2) The detailed mechanism of the synthesis of ATP



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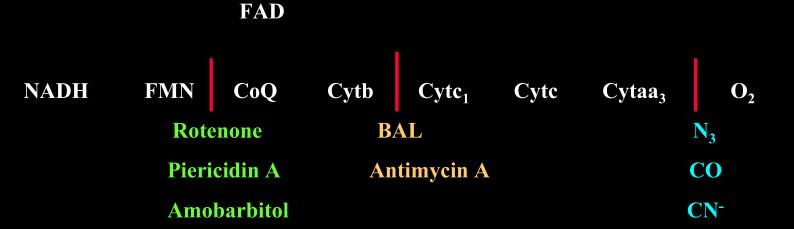
# 3. The factors influenced of oxidation and phosphorylation

- (1), ADP+Pi/ATP ratio is the primary factor to regulate the oxidation phosphorylation
- (2). Oxidative phosphorylation is also regulated by hormone: The thyroxine is able to facilitate the reaction of hydrolysis to resolve ATP into ADP and Pi.
- (3). The inhibitors of the oxidation phosphorylation Inhibitor may be divided into the inhibitors of the respiratory chain proper the inhibitors of the oxidative phosphorylation, and the uncouples of the oxidative phosphorylation

#### **1** The inhibitors of the oxidative phosphorylation:

Medicines: rotenone, piericidin A, amobarbital, antimycin A, dimercaptopropanol (BAL).

Poisons: CO, CN $^-$ ,  $N_3^+$ 



#### 2. Uncouples:

Dinitrophenol, Pentachlorophenol, dinitrocresol

The action of the uncouples is to dissociate oxidation in the respiratory chain from phosphorylation. P/O ratio decreases the quality of ATP synthesized is decreased, and speed up the oxidative of substrates.

#### 3 Oligomycin:

The oligomycin completely blocks the electrons from flowing back to  $\mathbf{F}_1$  from  $\mathbf{F}_0$ . This inhibitors prevent ATP from being synthesized from ADP and Pi . It lead to the uncouples of the oxidative phosphorylation.

(4) The mutation of the mitochondrial DNA (mt DNA)

# 4. High-energy phosphates — ATP

# (1). High-energy phosphates bond

If the free energy that released when the phosphate esterbond is hydrolyzing is more than 21KJ/mol, we call it the high energy phosphate bond . Lipmann introduced the symbol "~P" to represent it.

The compounds contained the high energy phosphate bond are called high energy phosphate compounds.

(2). The common some high-energy compounds in organism:

### (3), ATP cycle:

Oxidative
phosphate

Creatine

Muscle contraction
Materials transport

~P

Substrate level
phosphorylation

Creatine
phosphate
ADP

Transmit messages

### **The role of ATP cycle:**

- **Exchanging energy:** ATP acts as the "energy currency" of the cell, transferring free energy derived from substances of higher energy potential to those of lower energy potential.
- **Connect the phosphorylation and dephosphorylation in the progress of metabolism.**

## (4) The transformation of the high-energy phosphate bond

① Adenylate kinase interconversts adenine nucleosides:
The enzyme adenylate kinase is prevent in most cells. It catalyzed the interconversion of ATP and AMP on the hand and ADP on the other:

Adenosine kinase

ATP + AMP

ADP + ADP

# 2 Other nucleoside tri-phosphates participate in the transfer of high-energy phosphate.

By means of the enzyme nucleoside di-phosphate kinase, nucleoside tri-phosphates similar to ATP but containing an alternative base to adenine can be synthesized from their di-phosphates, eg:

nucleoside diphosphate kinase

ATP + UDP ADP + UTP

nucleoside diphosphate kinase

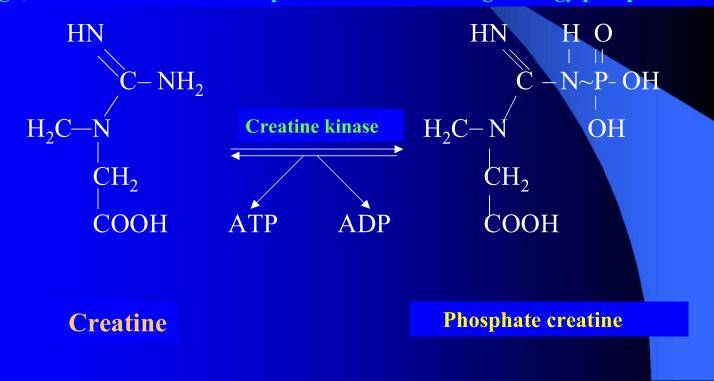
ATP + GDP ADP + GTP

nucleoside diphosphate kinase

 $\overline{ATP + CDP}$   $\overline{ADP + CTP}$ 

#### 3. Creatine phosphate shuttle:

The phosphagens, act as storage forms of high-energy phosphate, occurring in vertebrate skeletal muscle, heart and brain, under physiologic conditions phosphagens permit ATP concentrations to be maintained in muscle when ATP is rapidly being utilized as a source of energy for muscular contraction. On the other hand ,when ATP is plentiful and the ATP/ADP ratio is high, their concentration can up to act as a store of high-energy phosphate.



## 5. Substance transporter systems of the mitochondrial inner membrane

(1) Shuttle systems are required for mitochondrial oxidation of cytosolic NADH:

The dehydrogenase of the inner mitochondrial membrane of animal cells can accept electrons only from NADH in the matrix, given that the inner membrane is not permeable to cytosolic NADH, how can the NADH generated by glycolysis outside mitochondrial be re-oxidized to NAD $^+$  by  $O_2$  via the respiratory chain?

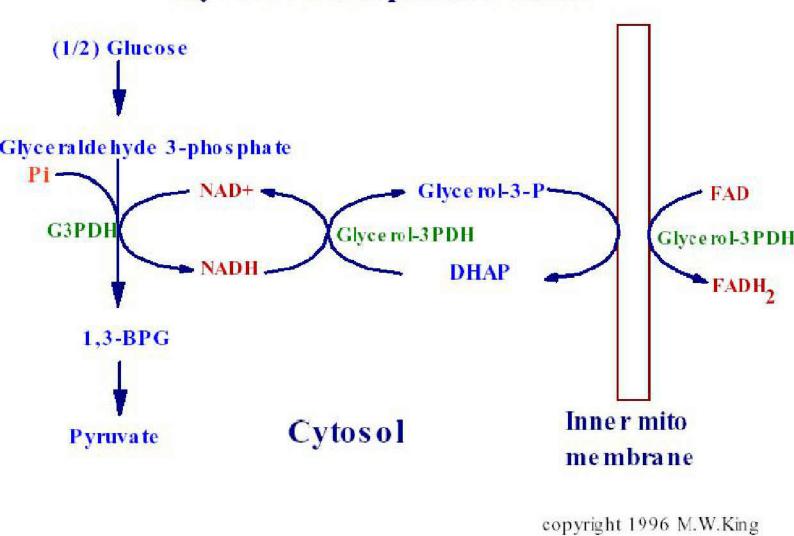
Special shuttle systems carry reducing equivalents from cytosolic NADH into mitochondrial by an indirect route.

The most active NADH shuttle, is the malate-aspatate shuttle and the glycerol-3-phosphate shuttle.

#### 1. The glycerol-3-phosphate shuttle:

This shuttle in skeletal muscle and brain, in that it delivers the reducing equivalents from NADH into complex  $\Pi$  (into succinate oxidation respiratory chain) not complex  $\Pi$ , providing only enough energy to synthesize two ATP(2 ATP)molecules per pair of electrons.

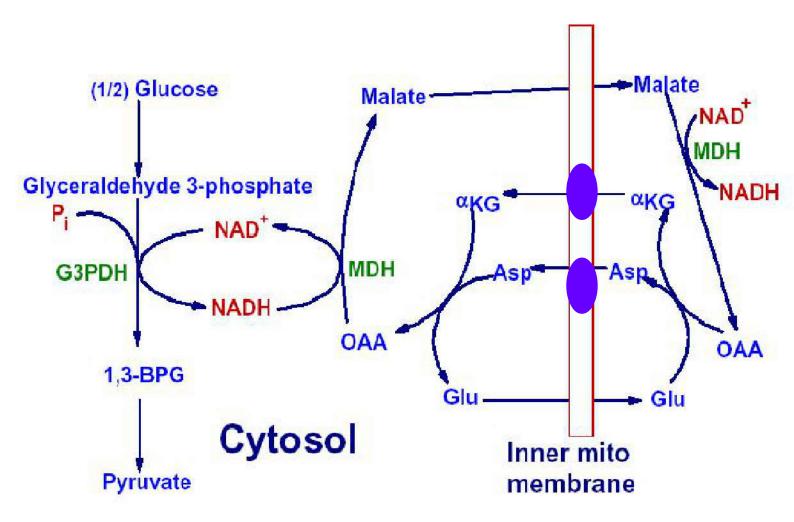
#### Glycerol Phosphate Shuttle



#### 2. The malate – asparate shuttle:

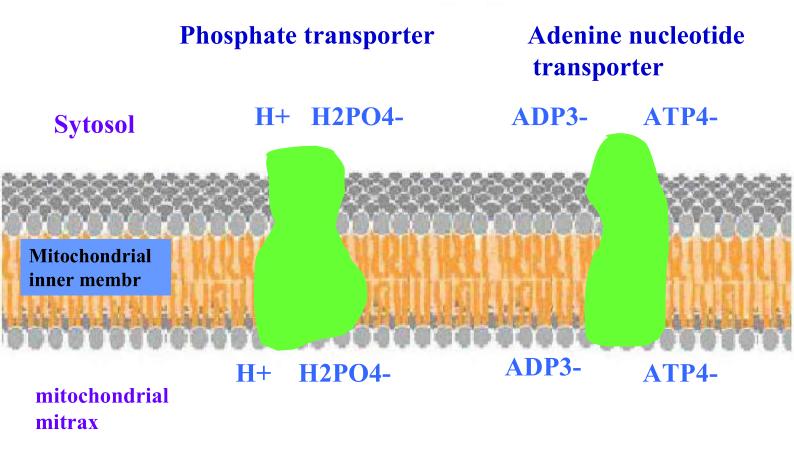
This shuttle in liver, kidney and heart mitochondria, in that it delivers the reducing equivalents from NADH into complex  $\, {\rm I} \,$  (into NADH oxidation respiratory chain), three molecules of ATP are generated as this pairs of electrons passes to  $\, {\rm O}_2 \, . \,$ 

### **Malate-Aspartate Shuttle**

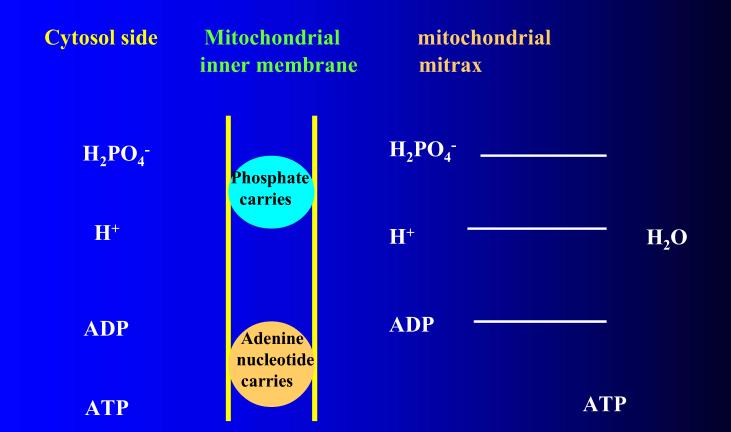


# (2) Adenine nucleotide transporter (ATP-ADP carrier)

The adenine nucleotide transporter allows the exchange of ATP and ADP but not AMP. It is vital in allowing ATP exit from mitochondria to the sites of extramitochondria utilization and allowing the return of ADP for ATP production within the mitochondrial.



**H2O** 



(3) The protein transports across

the mitochondrial membrane into the mitochondria by receptors that existed in membrane.

## 总结

生物氧化的概念、特点

生成ATP的氧化体系

- 一、<u>呼吸链</u> 定义、组成、排列顺序、两条主要的呼吸链
- 二、<u>氧化磷酸化</u> 定义、偶联部位、偶联机制
- 三、影响氧化磷酸化的因素 ADP、激素、抑制剂、解偶联剂、线粒体DNA突变

四、高能磷酸化合物—— ATP 高能磷酸键、常见的高能化合物、ATP循环、高能磷酸键的 转移。

五、通过线粒体内膜的物质转运 <u>胞液中NADH的氧化: a-磷酸甘油穿梭</u> <u>苹果酸-天冬氨酸穿梭</u>

腺苷酸载体

线粒体蛋白的跨膜转运

### 生物氧化复习题

#### 一、名词解释

呼吸链 磷氧比值 生物氧化 ATP循环 高能化合物 氧化磷酸化

#### 二、问答题

- 1、简述呼吸链的组成成分及主要功能?
- 2、试述ATP在能量转换的核心作用?
- 3、简述线粒体外产生的NADH如何进行氧化磷酸化?