

OUTLINE:

Introduction and review
Transport
Glycogenolysis
Glycolysis
Other sugars
Pasteur- Anaerobic vs Aerobic
Fermentations
Pyruvate
Krebs' Cycle
Oxidative Phosphorylation
Electron transport
Chemiosmotic theory: Phosphorylation

Exam-1 material

Exam-2 material



THE EVACUATION OF BOSTON

Fat Catabolism

diet
storage
Fatty acid Catabolism
FOUR stages in the catabolism of lipids:
Mobilization from tissues (mostly adipose):
hormone regulated
specific lipases
glycerol
Activation of fatty acids
Fatty-acyl CoA Synthetase
Transport
carnitine
Oxidation
Rationale
Saturated FA
β-oxidation
4 steps:
dehydrogenation
hydration
oxidation
thiolase
energetics
Unsaturated FA
energetics
Odd-chain FA
Ketone Bodies
Other organelles

Exam-3 material

Protein Degradation (Catabolism)

Digestion
Inside of cells
Protein turnover
Ubiquitin
Proteasome
Amino-Acid Degradation
Dealing with the nitrogen
Ammonia
free
transamination
Carbamoyl-phosphate synthetase
Urea Cycle
5 Steps
Ornithine transcarbamylase
Arginino-succinate synthetase
Arginino-succinase
Arginase
Energetics
Urea Bi-cycle
Dealing with the carbon

Seven Families

1. ADENQ (Trans-/de- aminase Family)
2. RPH (Glu Family)
Oxidase
One-carbon (1-C) metabolism
THF
SAM
3. GSC (Pyruvate Family)
PLP uses
4. MT - 1-C metabolism (α-KetoB Fam)
5. FY - oxidases (Aromatic Family)
6. KW (α-Ketoadipate Family)
7. VIL (Branched-chain AA Family)

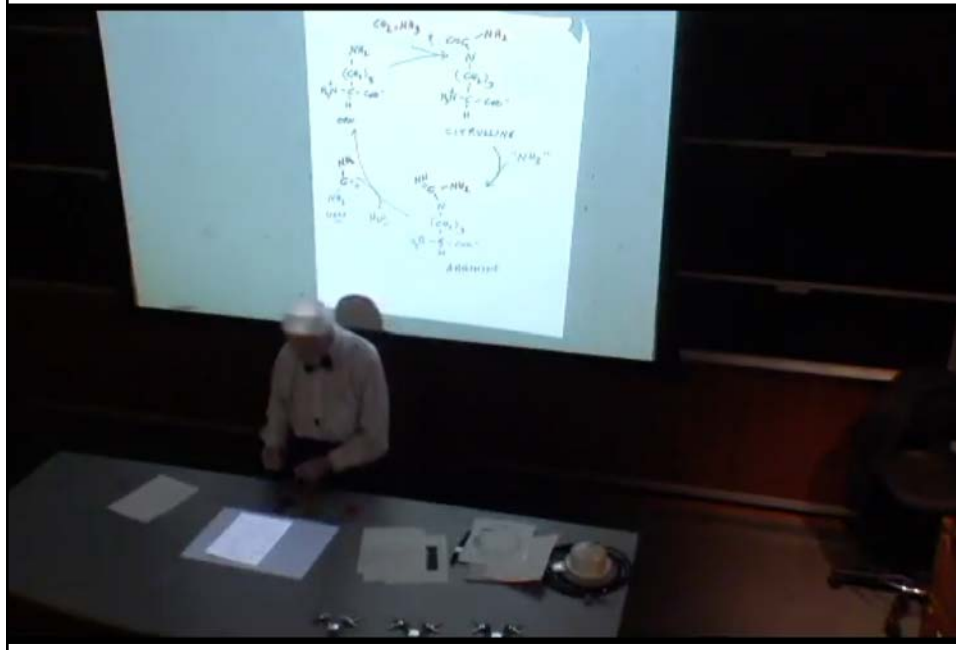
Nucleic-Acid Degradation

nucleotide to bases
salvage vs oxidation
purine
pyrimidine



Know mechanism →

Amino Acid Catabolism: Urea Cycle



Amino Acid Degradation: the carbon “skeletons”

A. Concepts

1. Convergent
2. ketogenic/glucogenic
3. Reactions seen before

The SEVEN (7) Families

B. Transaminase (A,D,E) / Deaminase (N,Q) Family (5)

C. Related to biosynthesis (8)

1. Glu Family (R,P,H)

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- a. PLP reactions

3. α -Ketobutarate Family (M,T)

- a. 1-C metabolism

In total there are 29 Nitrogen atoms in the 20 amino acids:

9 are given off as ammonia

18 are taken off in transaminase reactions

2 leave as urea

D. Dedicated (7)

1. Aromatic Family (F,Y)

- a. oxidases/oxygenases

2. α -Ketoadipic Family (K,W)

3. Branched-chain Family (V,I,L)

There are ~75 reactions in total

E. Convergence with Fatty Acids: propionyl-CoA

Amino Acid Degradation

- Intermediates of the **central metabolic pathway**
- Some amino acids result in more than one intermediate.
- Ketogenic amino acids can be converted to ketone bodies.

Seven to **Acetyl-CoA** Leu,⁷ **Ile**,⁷ **Thr**, Lys,⁶ **Trp**,⁶ **Phe**,⁵ **Tyr**⁵

- Glucogenic amino acids can be converted to glucose.

Six to **pyruvate**^{3,1} Ala, Cys, Gly, Ser, **Thr**, **Trp**

Five to **α -ketoglutarate**^{2,1} Arg, Glu, Gln, His, Pro

Four to **succinyl-CoA**^{7,4} **Ile**, Met, **Thr**, Val

Two to **fumarate**⁵ **Phe**, **Tyr**

Two to **oxaloacetate**¹ Asp, Asn

¹Trans-/de-aminase family

²Glu family

³Pyruvate family

⁴Alpha-keto butyric family

⁵Aromatic family

⁶Alpha-keto adipic family

⁷Branched-chain family

NOTES:

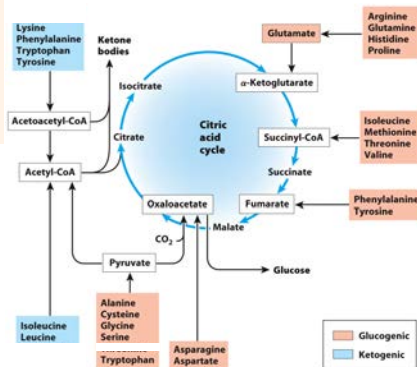
blue bkgd=ketogenic

salmon bkgd=glucogenic

Red=both keto- and gluco-genic

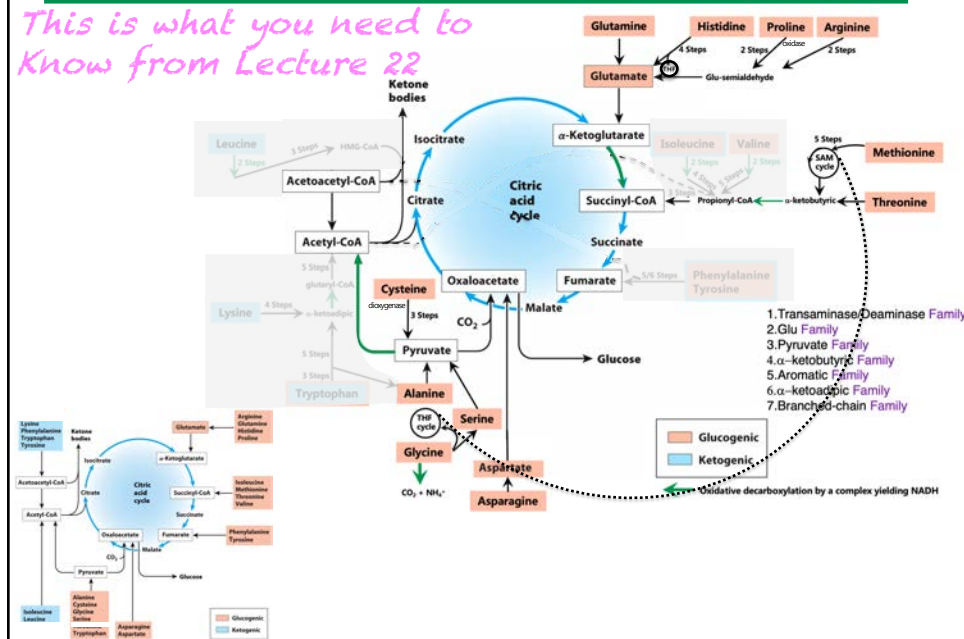
Italics=different non-human pathways

Bold=pathway discussed in class



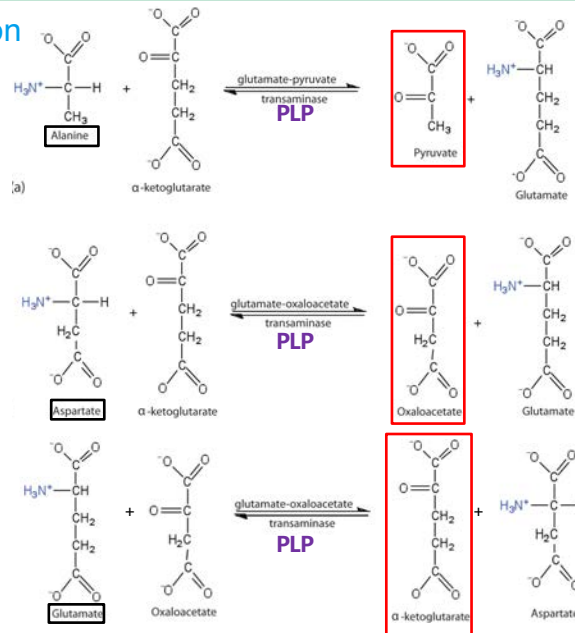
Amino Acid Degradation

This is what you need to know from Lecture 22



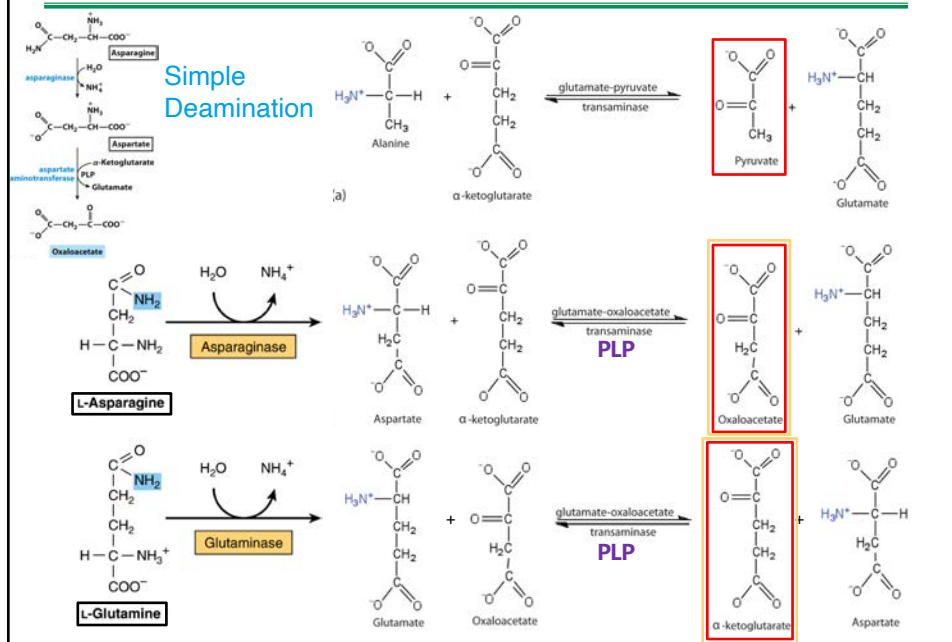
Amino Acid Degradation A,D,E

Simple Transamination



Amino Acid Degradation

N,Q



Amino Acid Degradation

A,D,E,N,Q

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Two to **oxaloacetate**¹

✓¹Trans-/de-aminase family

²Glu family

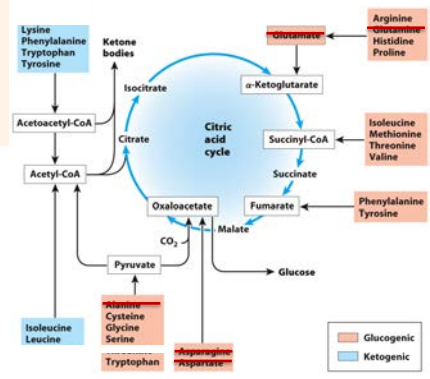
³Pyruvate family

⁴Alpha-keto butyric family

⁵Aromatic family

⁶Alpha-keto adipic family

⁷Branched-chain family



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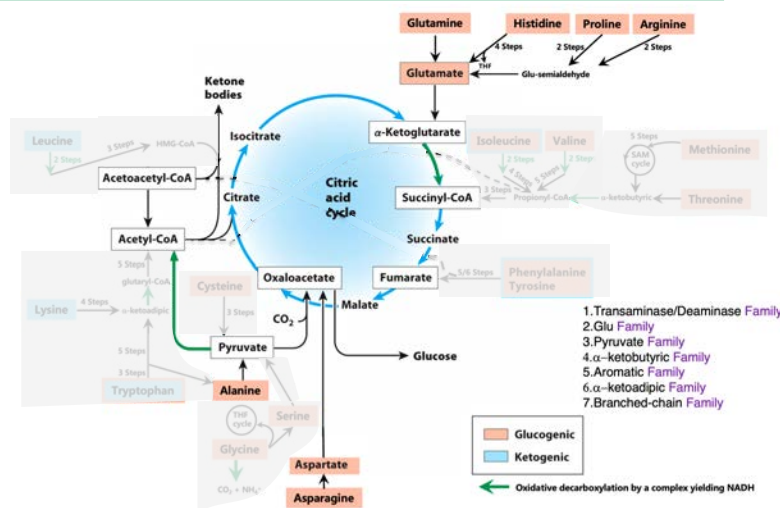
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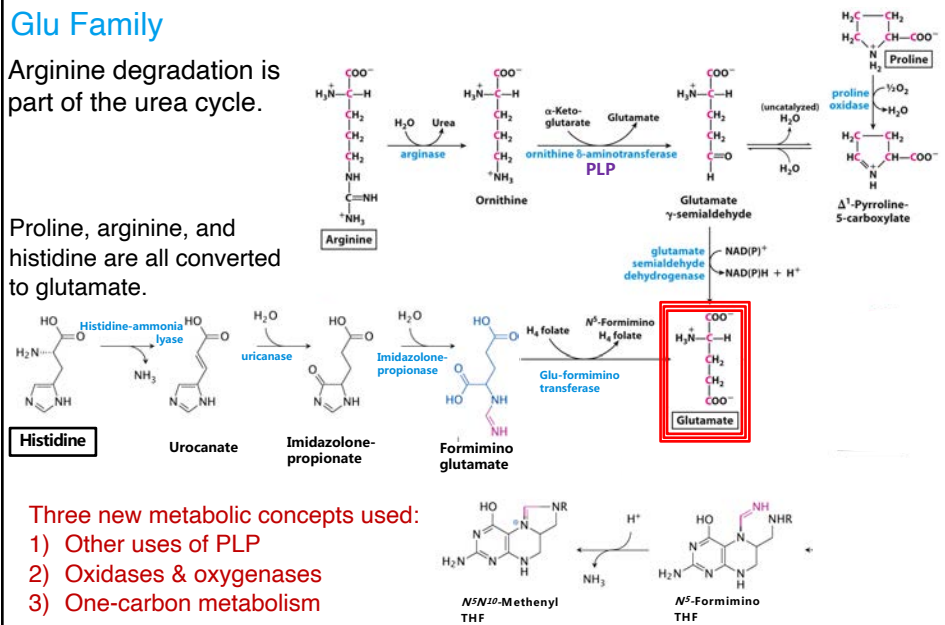
Before Glu

Amino Acid Degradation



R,P,H

Proline, arginine, and histidine are all converted to glutamate.

[illegible]

Amino Acid Degradation R,P,H

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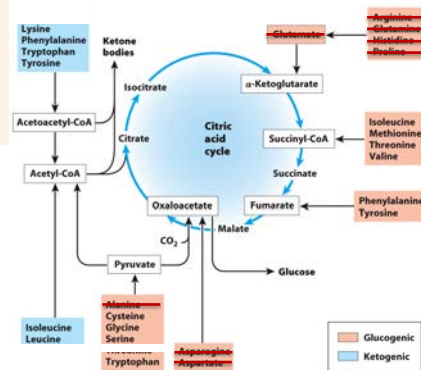
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Two to fumarate⁵ Phe, Tyr

Two to oxaloacetate¹

- ✓¹Trans-/de-aminase family
- ✓²Glu family
- ³Pyruvate family
- ⁴Alpha-keto butyric family
- ⁵Aromatic family
- ⁶Alpha-keto adipic family
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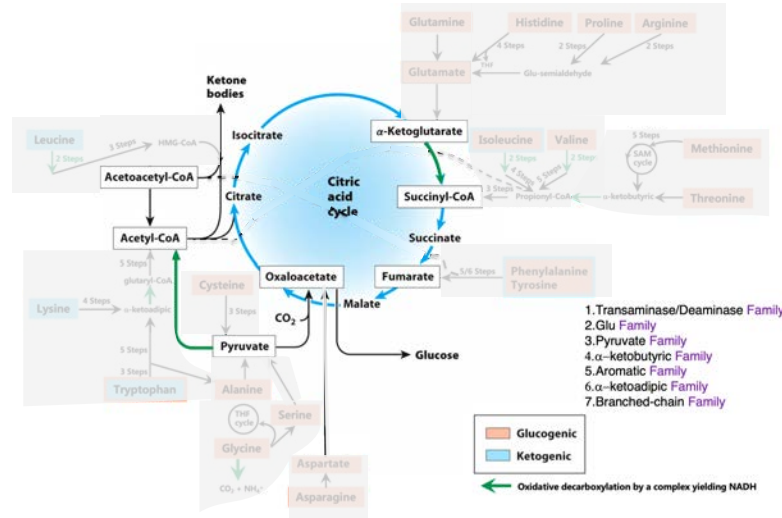
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Amino Acid Degradation



Amino Acid Degradation

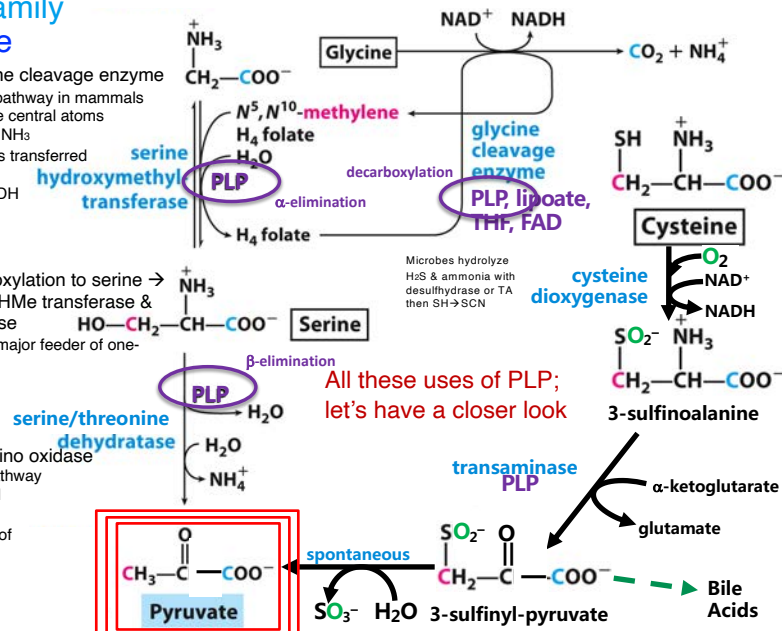
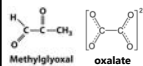
G,S,C

Pyruvate Family Glycine

- Pathway #1: glycine cleavage enzyme
 - apparently major pathway in mammals
 - separation of three central atoms
 - releases CO_2 and NH_3
 - methylene group is transferred to THF
 - mechanism like PDH complex

- Pathway #2: hydroxylation to serine → pyruvate via SerOHme transferase & Ser/Thr dehydratase
 - Reverse reaction major feeder of one-carbon to THF

- Pathway #3: D-amino oxidase
 - relatively minor pathway
 - ultimately oxidized to oxalate
 - major component of kidney stones

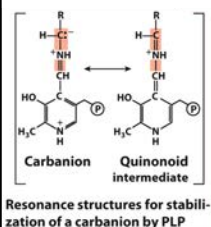
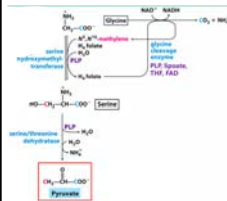


Amino Acid Degradation G,S,C

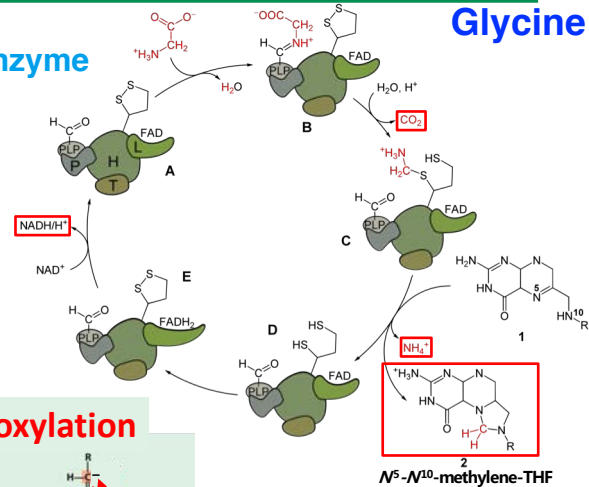
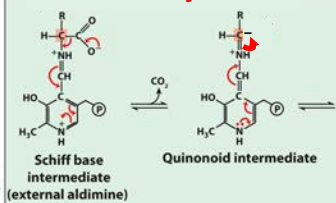
Pyruvate Family: Glycine Cleavage Enzyme

Glycine

See [Achieve](#) animated Figures



Decarboxylation



This is similar to how THF was used to pick off the formimino group in His catabolism

In Photosynthetic Parts of Plants, this is one of the most Abundant proteins and its called Glycine Decarboxylase

http://media.saplinglearning.com/priv/he/lehninger/acm/1820c_glycine_cleavage.html

Mechanism of Glycine Decarboxylase

- In plants, this same ortholog of Gly-cleavage enzyme is called Gly decarboxylase
- Flux through rubisco reaction and glycolate pathway can be high in bright sun, so plants need a lot of glycine decarboxylase.
 - makes up 50% of protein in some plant leaves
 - but non-photosynthetic parts of plants (tubers, etc.) have little glycine decarboxylase

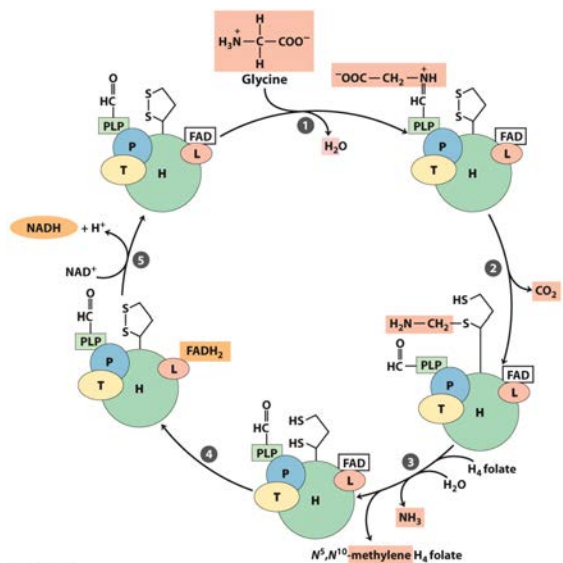
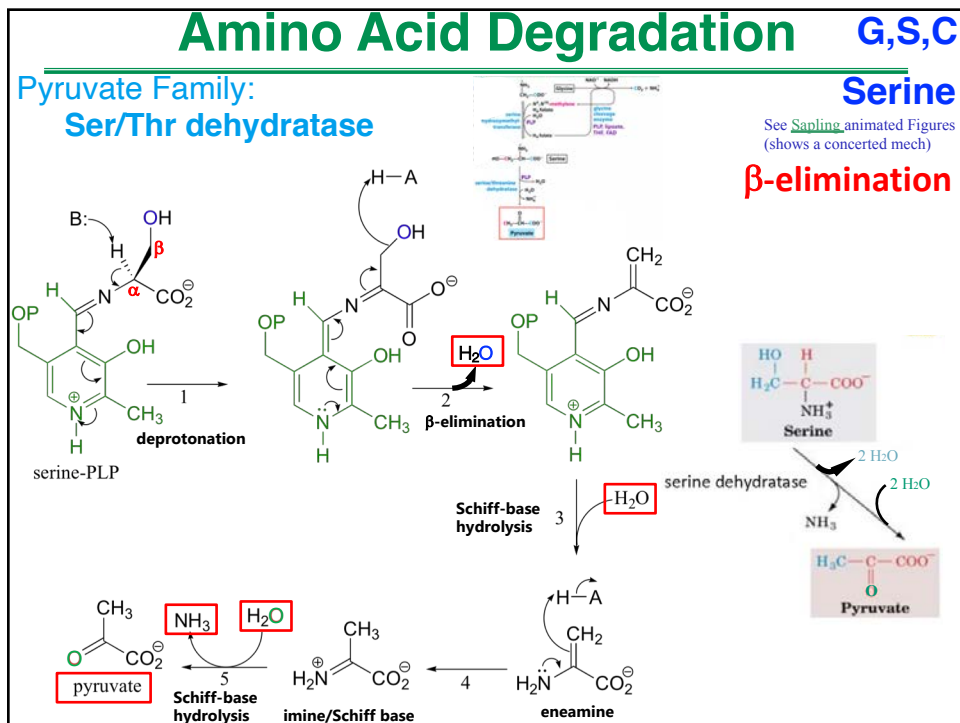
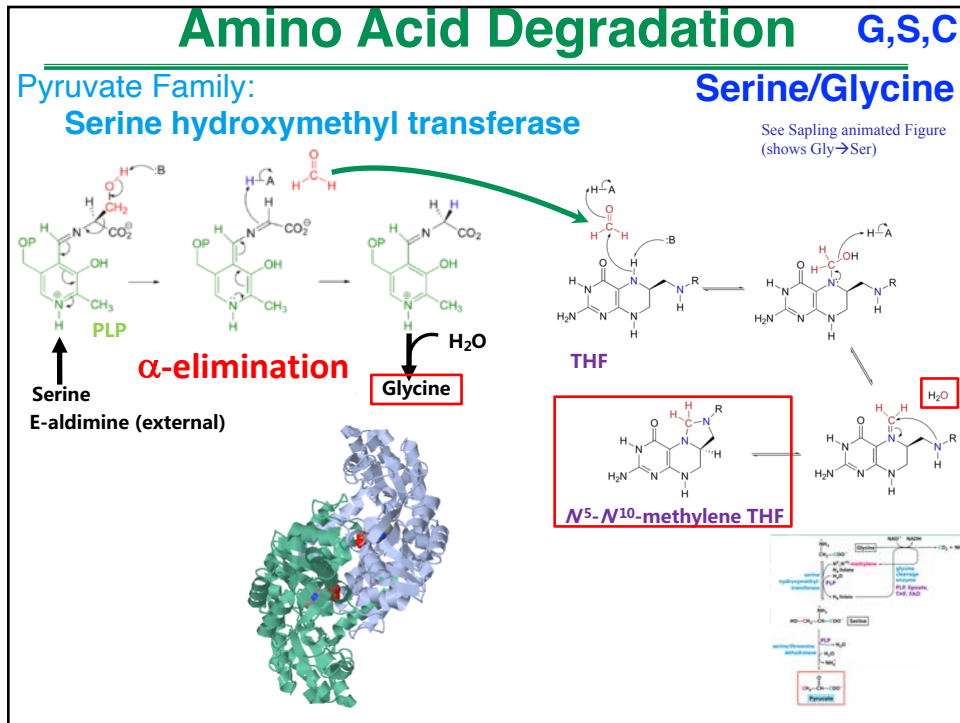


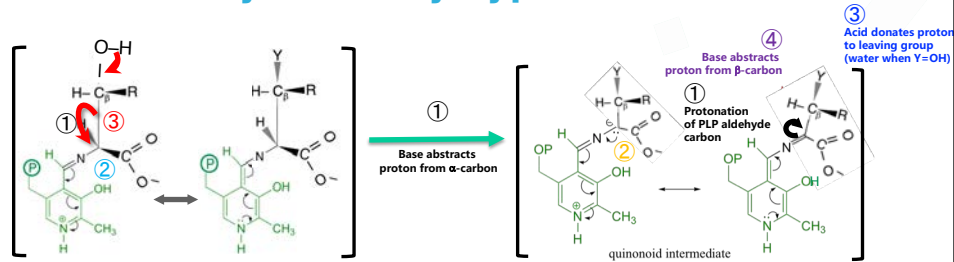
Figure 20-49

Glycine Decarboxylase Is Abundant in Photosynthetic Parts of Plants



Amino Acid Degradation

PLP Catalyzes Many Types of Reactions

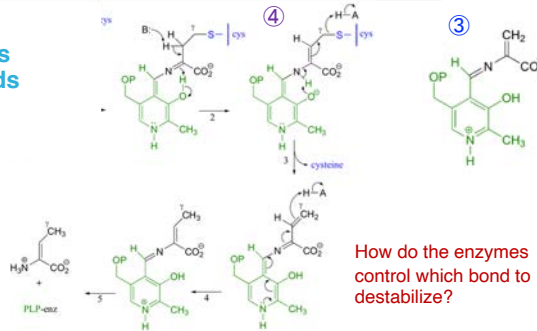


C α -bond:

- ① Transamination of Amino Acids
- ② Decarboxylation of Amino Acids
- ③ α -elimination of Amino Acids

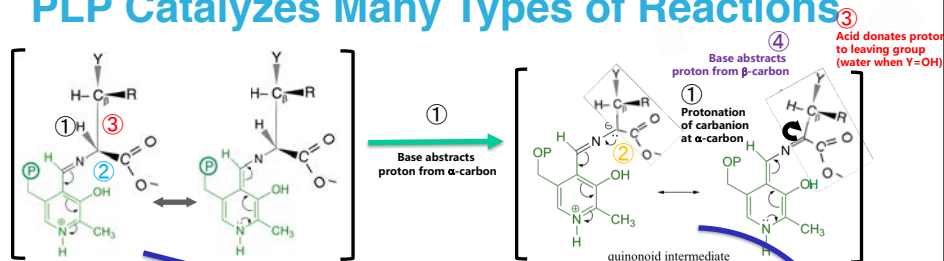
After C α -H abstracted ①:

- ② Racemization of Amino Acids
- ③ β -elimination of Amino Acids
- ④ γ -elimination of Amino Acids



Amino Acid Degradation

PLP Catalyzes Many Types of Reactions

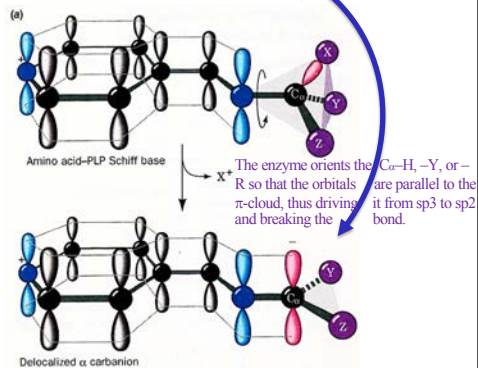


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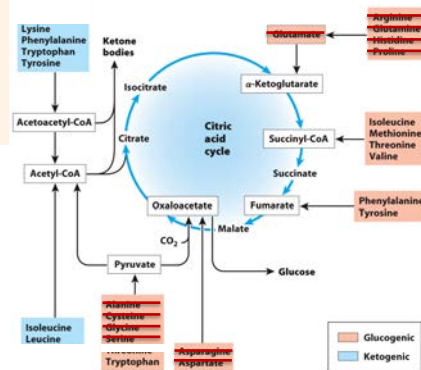
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