

BI/CH 422/622

OUTLINE:

Introduction and review
Transport
Glycogenolysis
Glycolysis
Other sugars
Pasteur: Anaerobic vs Aerobic

Fermentations

Exam-1 material

Pyruvate

Exam-2 material

Krebs' Cycle

Oxidative Phosphorylation

Electron transport

Chemiosmotic theory: Phosphorylation

Fat Catabolism

Exam-3 material

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

Protein Degradation (Catabolism)

Digestion

Inside of cells

Protein turnover

Ubiquitin

Proteasome

Amino-Acid Degradation

Dealing with the nitrogen

Ammonia

free

transamination ← **know mechanism**

Carbamoyl-phosphate synthetase

Urea

Cycle

4 Steps

Ornithine transcarbamylase

Arginino-succinate synthetase

Arginino-succinase

Arginase

Urea

Bi-cycle

Dealing with the carbon

Seven Families

1. **ADENQ** (Transaminase/deaminase 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 -Ketobutyrate Family)

5. **FY** - oxidases (Aromatic Family)

6. **KW** (α -Ketoadipate Family)

7. **VIL** (Branched-chain AA Family (BCAA))

Convergence with Fatty acid-odd chain

Nucleic Acid and Nucleotide Degradation

Nucleic Acids

Nucleotides

Salvage pathway

Degradation of purines

Degradation of pyrimidines

Connection of sugar metabolism

Fates of the 29 nitrogen
atoms in 20 AA:

9 ammonia

17 transamination

1 reduc. deamin.

2 urea

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 (Q,N) Family

C. Related to biosynthesis (R,P,H; C,G,S; M,T)

1. Glu Family

- a. Introduce oxidases/oxygenases
- b. Introduce one-carbon metabolism (1C)

2. Pyruvate Family

- a. PLP reactions

3. α -Ketobutyric Family (M,T)

- a. 1-C metabolism

D. Dedicated

1. Aromatic Family (F,Y)

- a. oxidases/oxygenases

2. α -Ketoadipic Family (K,W)

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

E. Convergence with Fatty Acids: propionyl-CoA

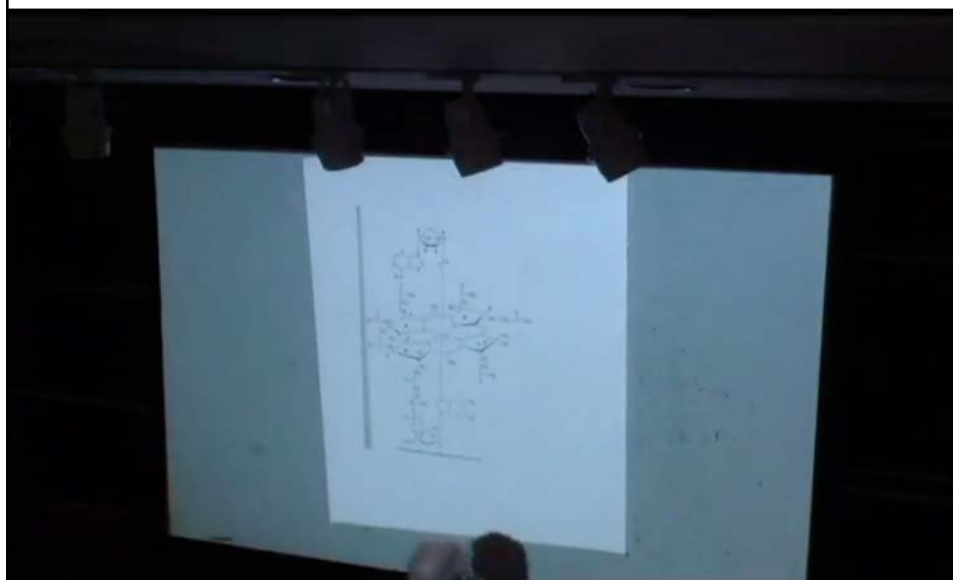
Before VIL

Only a handful of enzymes require Coenzyme B₁₂ (cobalamin)



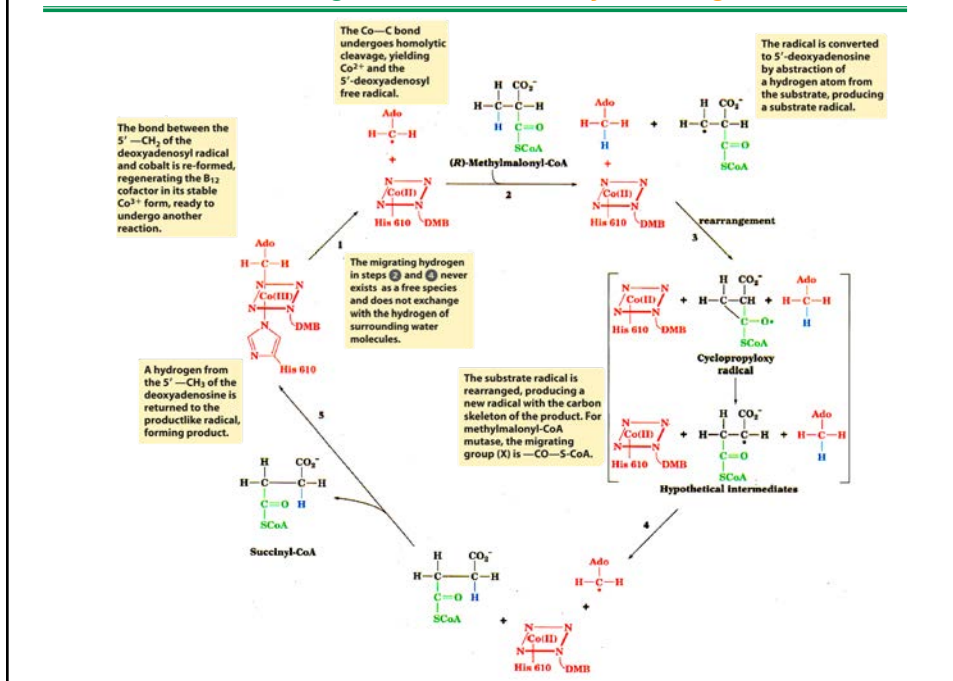
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The Vitamin B₁₂ Story

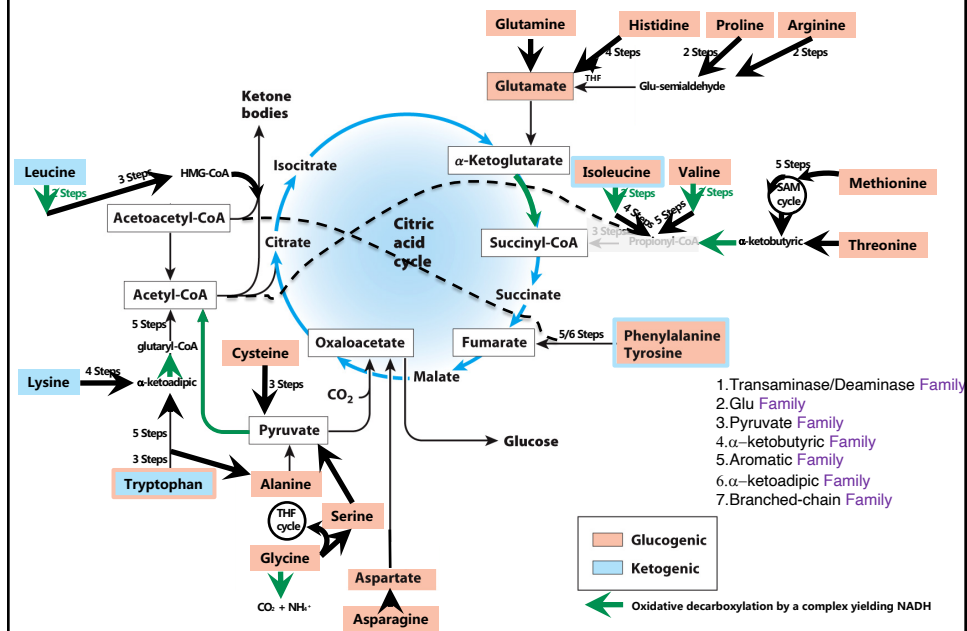


Dr. Kornberg: Lecture 16 02.27.17 (28:19-29:33) VitB-12 1.3 min

Amino Acid Degradation Meets Fatty Acid Degradation



Amino Acid Degradation: Overview



Amino Acid Degradation: Overview

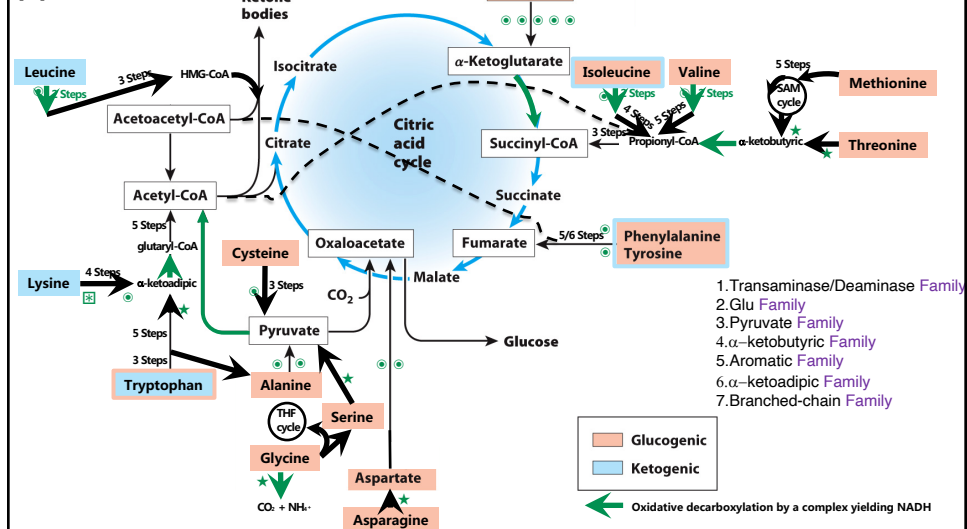
Fates of the 29 nitrogen atoms in the 20 amino acids:

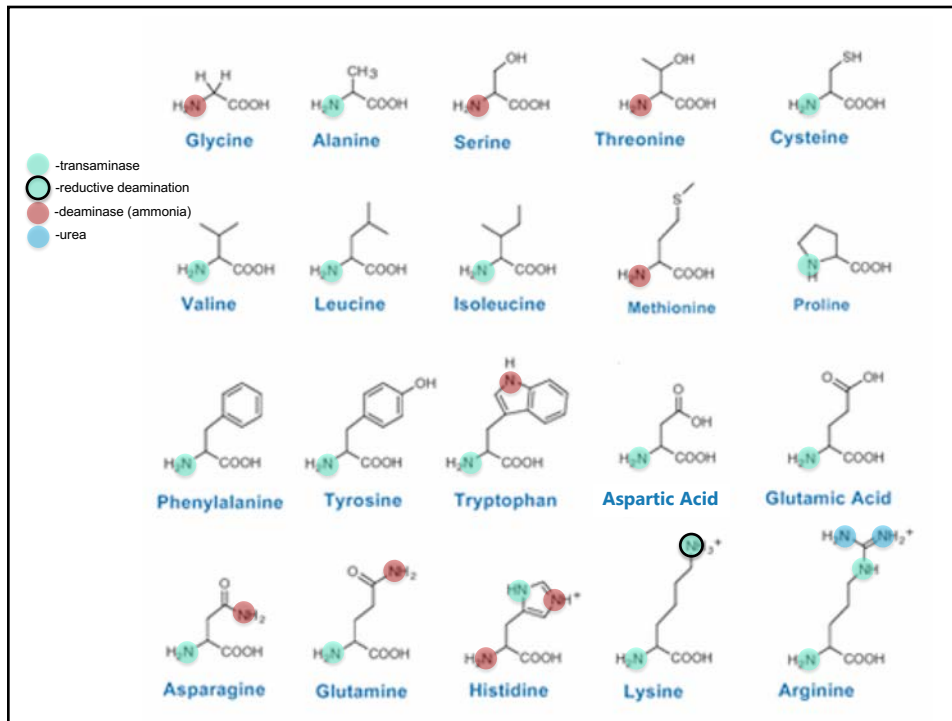
[★] 9 ammonia

[⊙] 17 transaminase (mostly Glu)

[⊞] 1 reductive deamination (also Glu)

[✱] 2 urea





Amino Acid Degradation: Overview

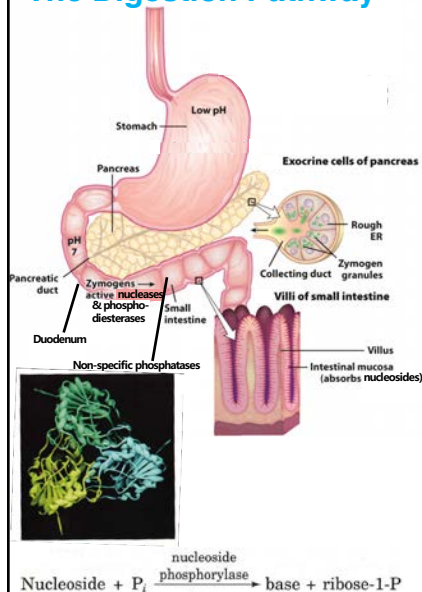
We learned that:

- amino acids from protein are an important **energy source** in carnivorous animals
- the first step of AA catabolism is transfer of the NH_3 via PLP-dependent aminotransferase usually to **α -ketoglutarate** to yield L-glutamate
- in most mammals, toxic ammonia is quickly recaptured into carbamoyl phosphate and passed into the **urea cycle**
- amino acids are degraded to pyruvate, acetyl-CoA, α -ketoglutarate, succinyl-CoA, and/or oxaloacetate
- amino acids yielding acetyl-CoA are ketogenic
- amino acids yielding other end products are glucogenic
- genetic defects in amino-acid degradation pathways result in a number of human diseases
- amino acid catabolism is dependent on a variety of cofactors, including THF, ado-Met (SAM), Cbl, biotin, and **PLP**

Nucleic-Acid Degradation

Nucleic-Acid Degradation

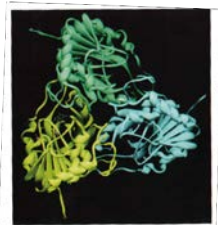
The Digestion Pathway



- Ingestion of food always includes nucleic acids.
- As you know from BI 421, the low pH of the stomach does not affect the polymer.
- In the duodenum, zymogens are converted to nucleases and the nucleotides are converted to nucleosides by non-specific phosphatases or nucleotidases.
- Only the non-ionic nucleosides are taken up in the villi of the small intestine.
- **In the cell**, the first step is the release of the ribose sugar, most effectively done by a non-specific **nucleoside phosphorylase** to give ribose 1-phosphate (Rib1P) and the free bases.
- **Most ingested nucleic acids are degraded to Rib1P, purines, and pyrimidines.**

Nucleotide Degradation: Overview

Fate of Nucleic Acids:



Purine Nucleoside Phosphorylase

Nucleotides

Nucleosides

Bases

Purine & Pyrimidine Salvage Pathway

The Salvage Pathways are in competition with the *de novo* biosynthetic pathways, and are both **ANABOLISM**

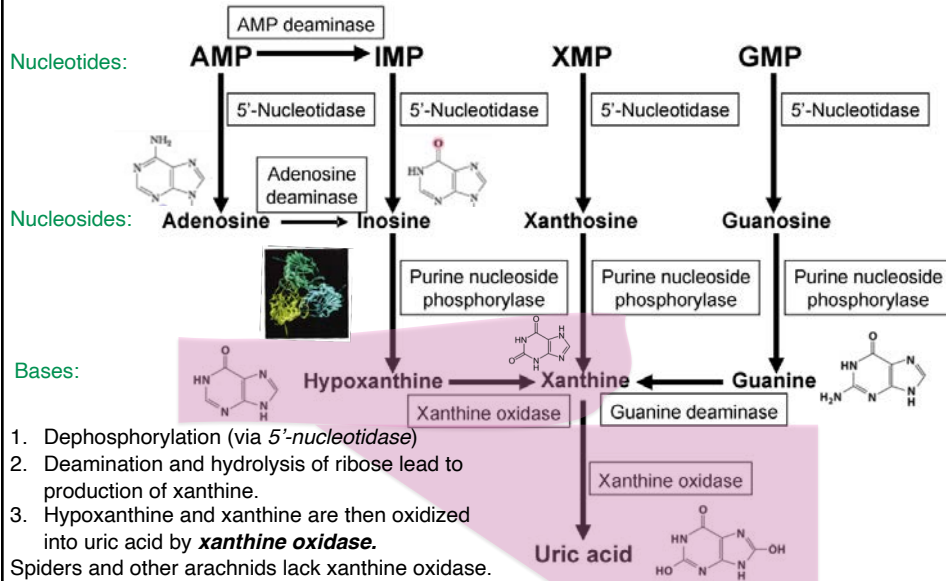
Once broken down to the nitrogenous bases they are either:

1. Salvaged for recycling into new nucleic acids (most cells; from internal, not ingested, nucleic acids).
2. Oxidized (primarily in the intestine and liver) by first converting to nucleosides, then to
 - Uric Acid (purines)
 - Acetyl-CoA & succinyl-CoA (pyrimidines)
 - Both yield ammonia

Oxidation

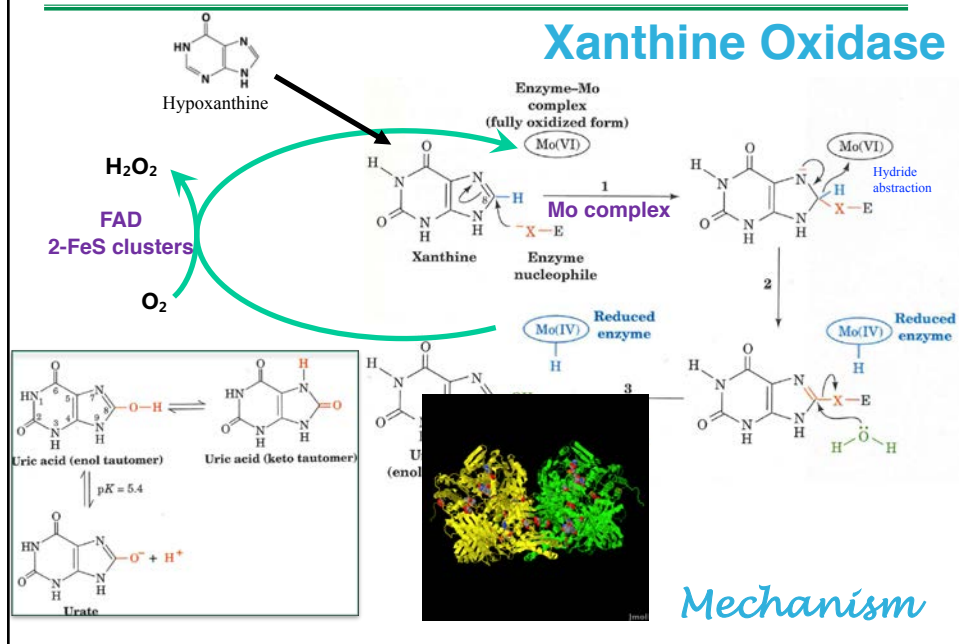
Nucleotide Degradation

Catabolism of Purines

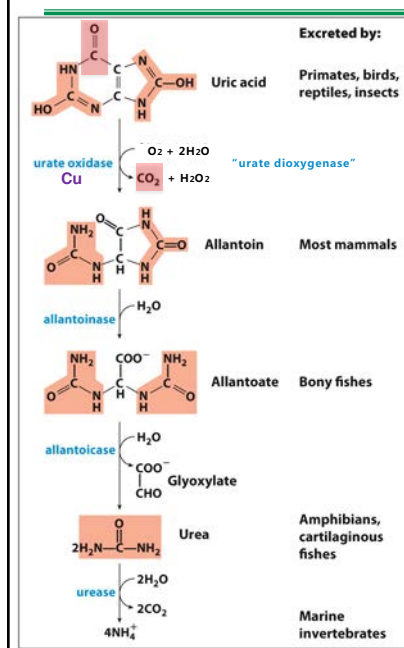


Nucleotide Degradation

Xanthine Oxidase



Nucleotide Degradation



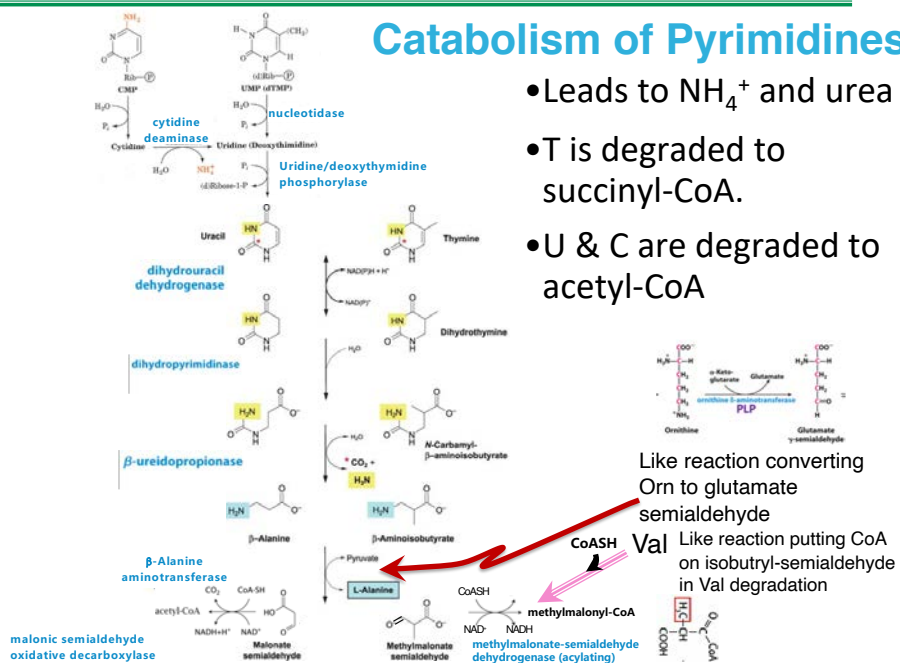
- Degree of further oxidation of uric acid is organism dependent.
- Birds and insects don't excrete amino-acid nitrogen as urea, but as uric acid to conserve water.

Conversion of Uric Acid to Allantoin, Allantoate, and Urea

Nucleotide Degradation

Catabolism of Pyrimidines

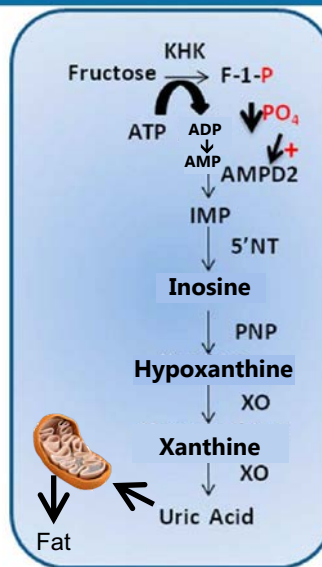
- Leads to NH_4^+ and urea
- T is degraded to succinyl-CoA.
- U & C are degraded to acetyl-CoA

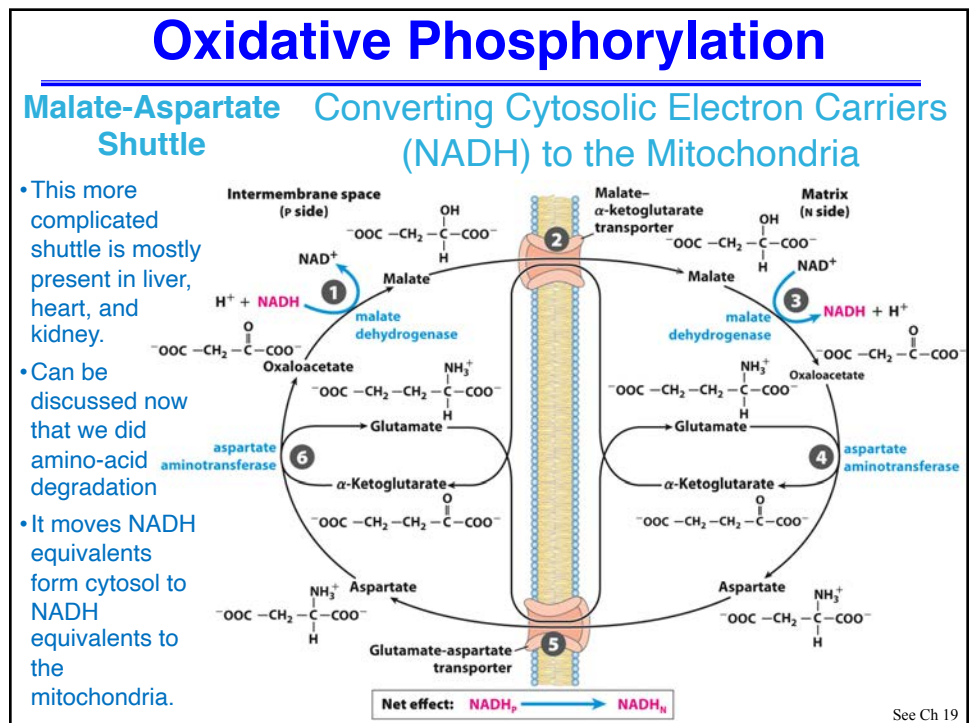
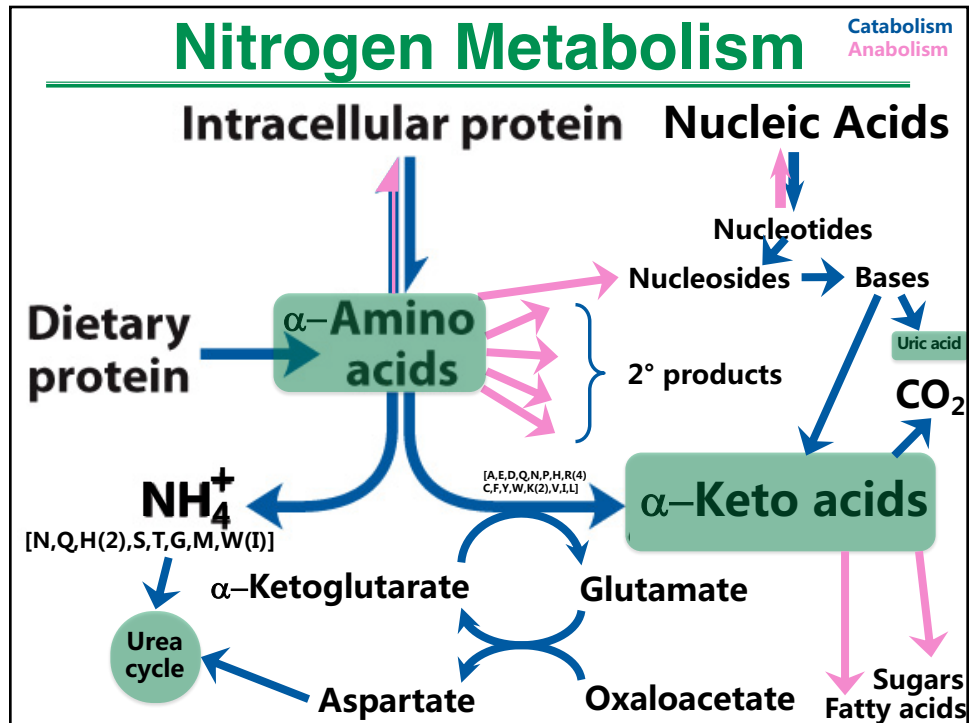


Nucleotide Degradation

Hyperuricemia: how sugar “becomes” fat

- The interesting connection between sugar metabolism and nitrogen metabolism
- What is even more interesting is that this metabolism is connected to fat metabolism as well
- The production of Uric Acid converts liver mitochondria to fatty acid synthesis by unknown mechanisms.





**End of Material for
Exam 3**