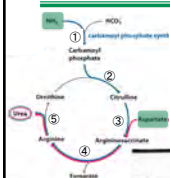


Amino Acid Catabolism: Urea Cycle

Enzymes of the Urea Cycle



See [Sapling](#) animated Figure on CPS mechanism ①

See [Sapling](#) animated Figure on ASS mechanism ③

	Enzyme	Compartment	Activity	M _r	pH opt	K _m , mM	Equilibrium Constant	Tissue Distribution
	N-acetylglutamate synthetase, EC 2.3.1.1	Mitochondrial matrix	0.30–1.49	200,000	8.5	Glu, 3.0 Ac CoA 0.7 Arg, 0.01	Irreversible	Liver, intestine, kidney (trace), spleen
①	Carbamoyl phosphate Synthetase, EC 6.3.4.16	Mitochondrial matrix	279 [†]	310,000 dimer	6.8–7.6	NH ₄ , 0.8 HCO ₃ , 6.7 Mg ATP, 1.1 NAG, 0.1	Irreversible	Liver, intestine, kidney (trace)
②	Ornithine transcarbamylase, EC 2.1.3.3	Mitochondrial matrix	6600	108,000 trimer	7.7	CP, 0.16 Orn, 0.40	$\frac{(\text{Cit})(\text{p})}{(\text{Orn})(\text{CP})} = 10^5$	Liver, intestine, kidney (trace)
③	Argininosuccinic acid synthetase, EC 6.3.4.5	Cytosol	90	185,000 tetramer	8.7	Asp, .03 Cit, .03	$\frac{(\text{ASA})(\text{AMP})(\text{Mg PP})_2(2\text{H})}{(\text{Cit})(\text{Asp})(\text{Mg ATP})} = 0.89^{\dagger}$	Liver, kidney, fibroblasts, brain (trace)
④	Argininosuccinase, EC 4.3.2.1	Cytosol	220	173,200 tetramer	7.5	Asp, 0.017 Cit, 0.016 ATP, 0.041	$\frac{(\text{Arg})(\text{lumarate})}{(\text{ASA})} = 11.4 \times 10^{-3}$	Liver, kidney, brain, fibroblasts
⑤	Arginase, EC 3.5.3.1	Cytosol	86,600	107,000 tetramer	9.5	Arg, 10.5	Irreversible	Liver, erythrocytes, kidney, lens, brain (trace)

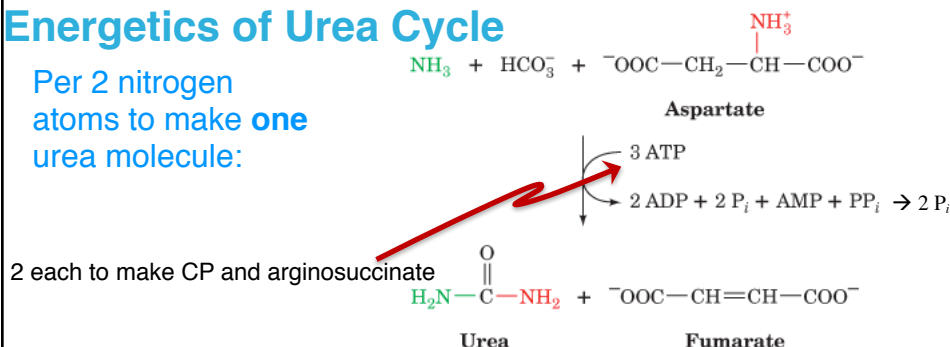
[†]Enzyme activity is expressed as micromoles per hour per gram wet weight. Apart from the equilibrium constants, the values described are those of human liver.
[†]The monomers may have substantial catalytic activity.¹¹

AT pH = 7.0.
Source: Table assembled from Rainer,⁴ Snodgrass,⁸ Meijer and Hensgens,¹⁰ Jackson, et al.,¹² Beaudet et al.,¹³ Lusty,¹³ and Bachman et al.¹⁴

Amino Acid Catabolism: Urea Cycle

Energetics of Urea Cycle

Per 2 nitrogen atoms to make one urea molecule:



Net is 4 ATP equivalents

[Its not actual ATPs used, its how many it takes to re-cycle the products (2 ADP + 1 AMP)]

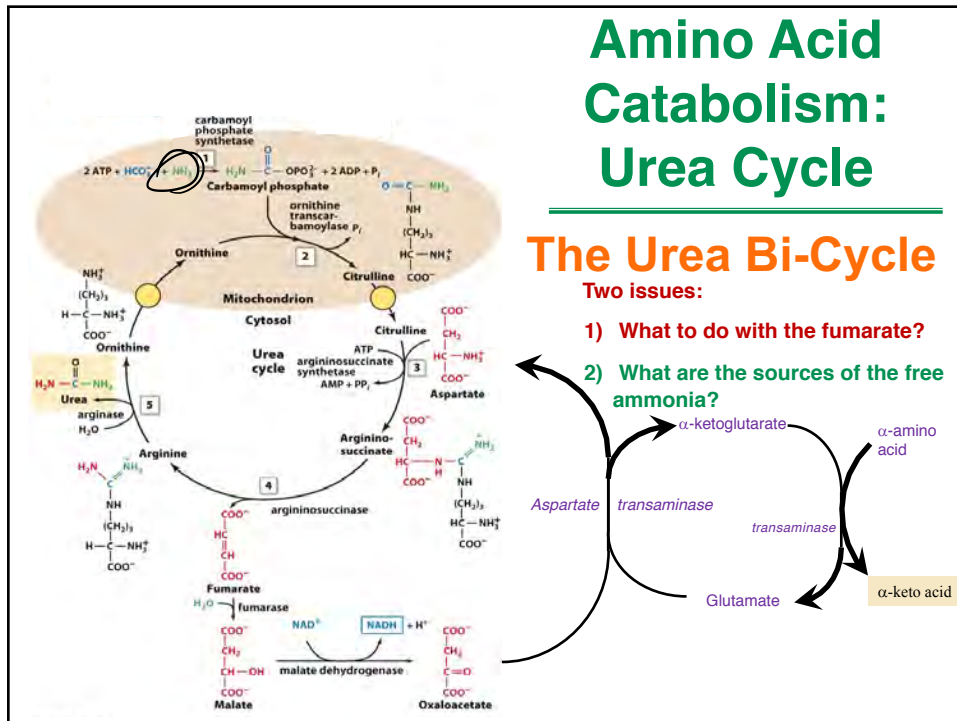
- But what about this fumarate?
- What is its fate?
- How do you regenerate Asp to keep the cycle going?

Amino Acid Catabolism: Urea Cycle

The Urea Bi-Cycle

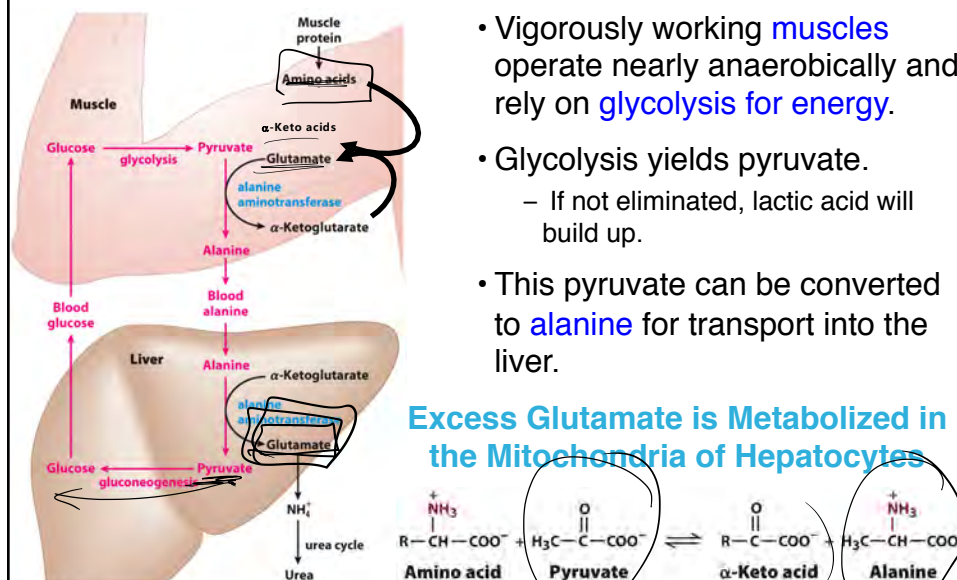
Two issues:

- 1) What to do with the fumarate?
- 2) What are the sources of the free ammonia?



Amino Acid Catabolism: Urea Cycle

The Glucose-Alanine Cycle

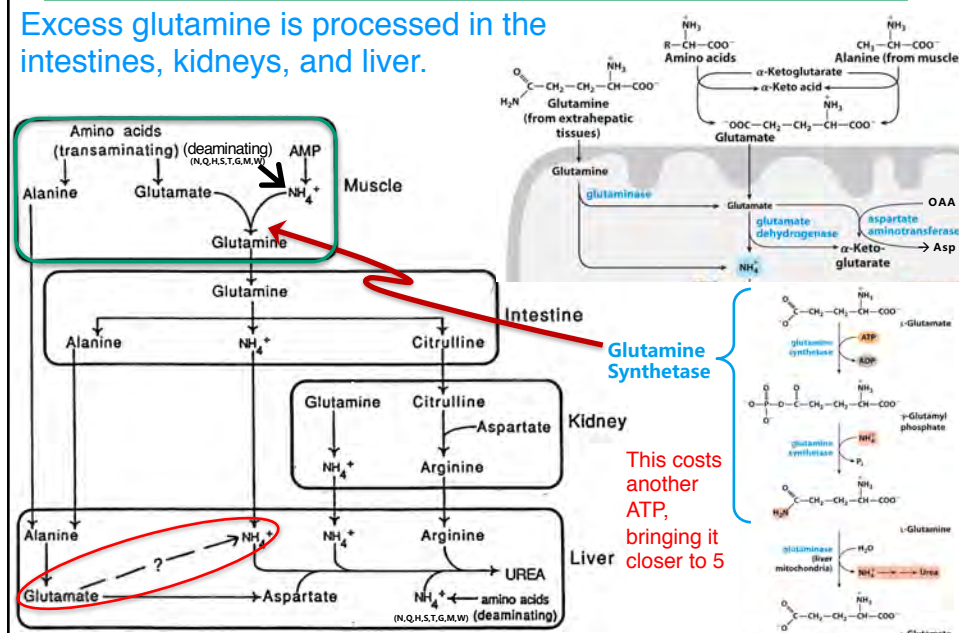


Excess Glutamate is Metabolized in the Mitochondria of Hepatocytes

Excess glutamine is processed in the intestines, kidneys, and liver.

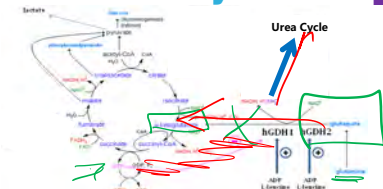


Excess glutamine is processed in the intestines, kidneys, and liver.



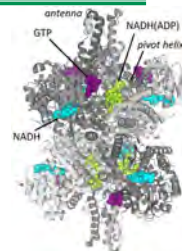
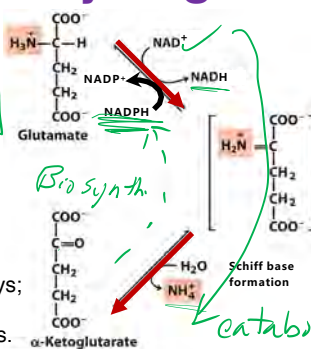
Amino Acid Catabolism: Urea Cycle

Regulation of the Urea Cycle



- The pool of **Glu** is a critical junction between biosynthesis and catabolism
- The enzyme represents a key link between **catabolic** and **anabolic** pathways; when biosynthesis is needed, its off and Glu provides for amino-acid biosynthesis. When energy is needed, the ammonia is pulled off to provide for the urea cycle and oxidation of carbon skeletons.
- Can use either NAD^+ in the catabolic direction (**oxidative deamination**), or NADPH in the biosynthetic direction.

Glutamate Dehydrogenase



Allosteric inhibitors:
GTP, ATP, Palmitoyl-CoA

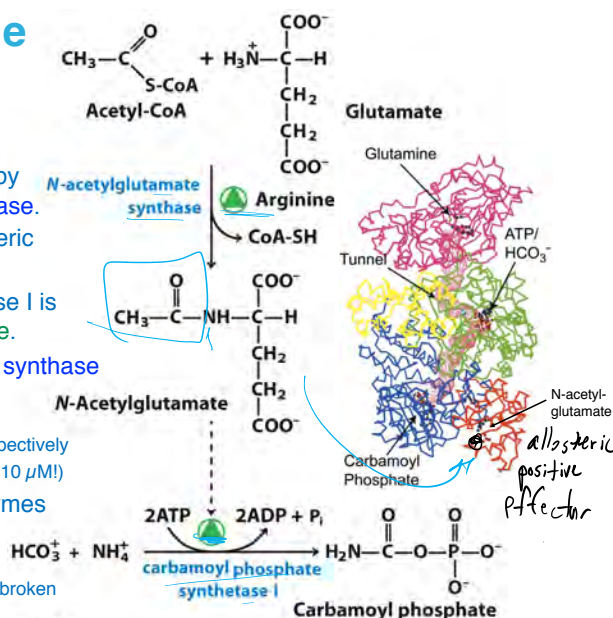
Allosteric Activators:
ADP, Leucine, Isoleucine, Valine

- GDH has a high K_m value ($>1 \text{ mM}$) for ammonia, so it mostly functions to provide the initial material for the urea cycle; ammonia, using NAD^+ , which is high when energy charge is low.
- Ammonia is processed into **urea** for excretion. Carbon skeletons are oxidized for ATP.

Amino Acid Catabolism: Urea Cycle

Regulation of the Urea Cycle

- Urea production is controlled by **carbamoyl phosphate synthetase**.
- N-acetylglutamate** is an allosteric activator.
- Carbamoyl phosphate synthetase I is activated by **N-acetylglutamate**.
- Formed by **N-acetylglutamate synthase**
 - when glutamate and acetyl-CoA concentrations are high
 - K_m values are 1 and 0.7 mM, respectively
 - activated by arginine (K_i value of $10 \mu\text{M}$)
- Expression of urea cycle enzymes increases when needed.
 - high-protein diet
 - starvation, when protein is being broken down for energy



Amino Acid Catabolism: Urea Cycle

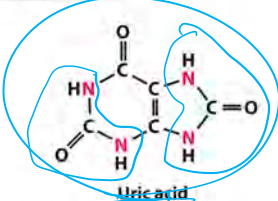
Excretory Forms of Nitrogen: Fates of Nitrogen in Organisms

NH_4^+
 Ammonia (as ammonium ion)

$\text{H}_2\text{N}-\text{C}(=\text{O})-\text{NH}_2$
 Urea

Ammonotelic animals: most aquatic vertebrates, such as bony fishes and the larvae of amphibia

Ureotelic animals: many terrestrial vertebrates; also sharks


 Uric acid

Uricotelic animals: birds, reptiles

Ure-otelic

Uric-otelic

- Plants conserve almost all the nitrogen.
- Many aquatic vertebrates release ammonia to their environment.
 - passive diffusion from epithelial cells
 - active transport via gills
- Many terrestrial vertebrates and sharks excrete nitrogen in the form of urea.
 - Urea is far less toxic than ammonia.
 - Urea has very high solubility.
 - Requires lots of water
- Some animals such as birds and reptiles excrete nitrogen as uric acid.
 - Uric acid is rather insoluble.
 - Excretion as paste allows the animals to conserve water.
- Humans and great apes excrete both urea (from amino acids) and uric acid (from purines).

Amino Acid Degradation

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; T,M)

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

2. Pyruvate Family
 - a. PLP reactions

3. α-ketobutyric Family
 - a. 1-C metabolism

D. Dedicated (F,Y; K,W; V,I,L)

1. Aromatic Family
 - a. oxidases/oxygenases
2. α-ketoadipic Family
3. Branched-chain Family