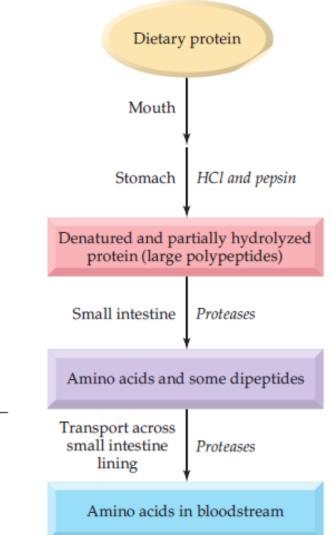


meilodsjeM Protein & Amino Acid

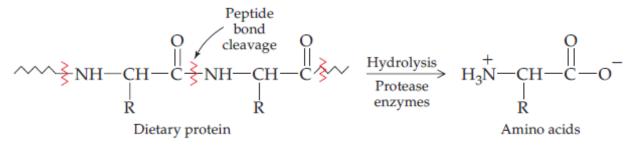
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Digestion of Proteins

- What is protein digestion?
- Where digestion begins?
- Stomach; HCI (2-3) & pepsinogen (polypeptides)
 - Acid and autocatalytic activation

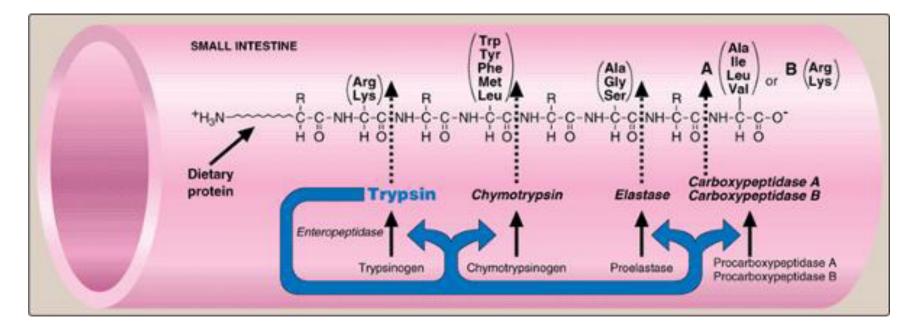


Hydrolysis of peptide bonds



Digestion of Protein

- Pancreatic (specific). Secretion mediated by cholecystokinin & secretin
- Enteropeptidase (mucosal) activates trypsinogen (the common activator of all the pancreatic zymogens). Result in amino acids and small peptides
- Intestinal lining enzymes, aminopeptidases



Absorption & transport of amino acids & dipeptides

Free amino acids (Na⁺-cotransport)

Di- & tripeptides (H+-cotransport) followed by hydrolysis

Only free amino acids are found in the portal vein

From cells to the bloodstream through active transport

Active transport is grouped (at least 7 transport systems); excess in one can cause deficiency in another!

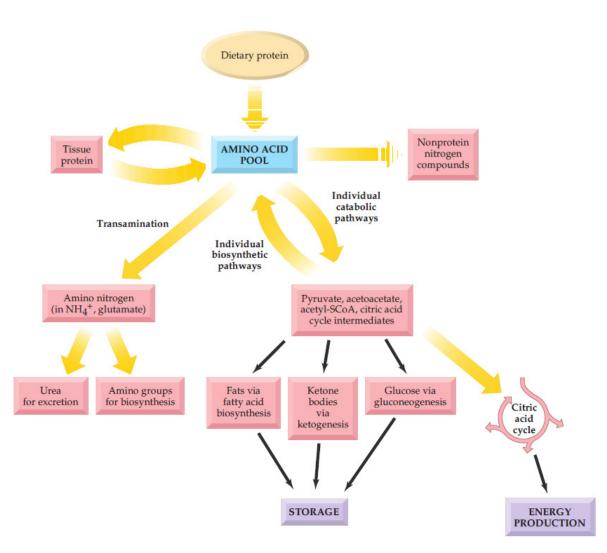
The small intestine & the proximal tubule of the kidney have common transport systems for amino acid uptake

Amino Acid Metabolism overview

- Amino acids are unlike fats and carbohydrates!
- Three sources; diet, synthesis, degradation
- Three destinations; synthesis, precursors, conversion
- Catabolism is of two phases; the amino group & the carbon skeleton
- Each amino acid is degraded via its own unique pathway but the general scheme is similar
- The metabolism involves two important concepts: amino acid pool and protein turn-over

Amino Acid Pool

- ■Where is it? How much is it? (≈100g)
- Supply: degradation, diet, synthesis
- Demand: synthesis, precursors, conversion
- In healthy, well-fed individuals, steady state, nitrogen balance

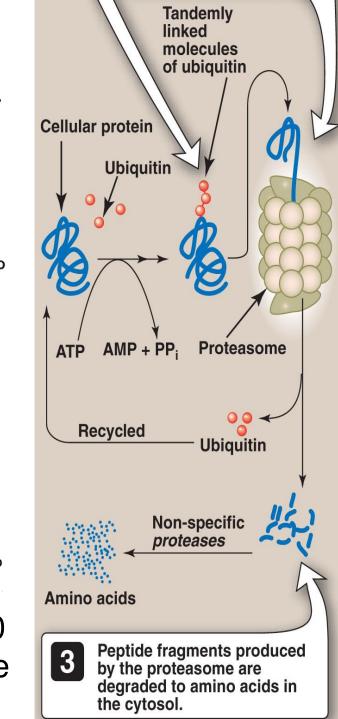


Protein Turn-over

- Most proteins in the body are in a constant remodeling.
- What regulates protein concentration in cells? synthesis or degradation?
- Turn over is about 300-400 g of protein per day, varies widely for individual proteins (minutes, hours, days, weeks, years)
- Two protein degradation pathways; ATP-dependent vs. independent, proteasomes vs. lysosomes, intracellular vs. extracellular

ATP dependent

- Ubiquitin-proteasome proteolytic pathway
- Ubiquitin is a small globular protein, nonenzymic
 ⁺H₃N-^c_aCH-^c_acH-^c_a
- Covalent attachment, polyubiquitination, proteasome complex, ubiquitins are recycled
- Is protein degradation random?
- Signals can be: oxidation of aa., ubiquitination, N-terminal residue; Ser (20 hrs) vs. Asp (3 min), PEST sequences are short lived



CH2

CH2

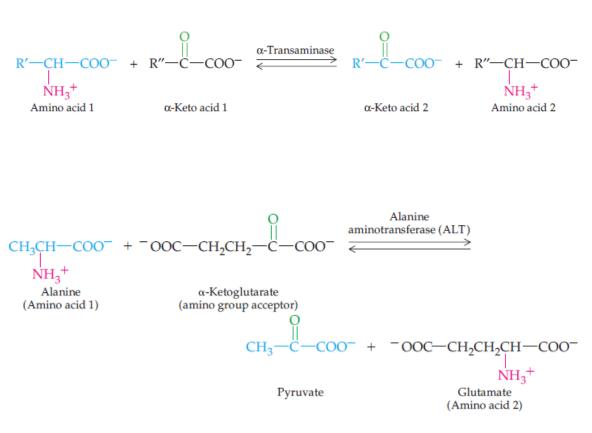
CH,

General scheme for amino acid catabolism

- ▶ Removal of amino group → Use of nitrogen (synthesis of Ncompounds → Passage of nitrogen (urea cycle) → carbon atoms convert to compounds (citric acid cycle or glucose formation)
- No storage of nitrogen-containing compounds & ammonia is toxic to cells
- N-containing compounds (NO, hormones, neurotransmitters, NAD⁺, heme, purine & pyrimidine bases)
- Carbon portion that converted to TCA cycle compounds is available for several pathways; energy (10-20%), TAGs (lipogenesis), glycogen (gluconeogenesis & glycogen synthesis), or ketone bodies

Catabolism of The Amino Group

- Removal of amino group
- Transamination
- Several aminotransferases (transaminases)
- Most are specific for αketogluterate, easily reversible depending on concentrations (regulation)



The products are an α-keto acid (derived from the original amino acid) & glutamate

Catabolism of **The Amino Group**

- Glutamate deamination
 - Oxidative deamination (α -ketogluterate)
 - Transamination (nonessential amino acids)

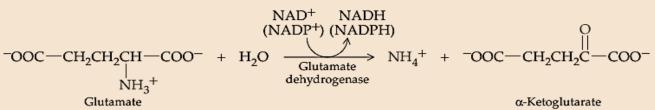
Alanine

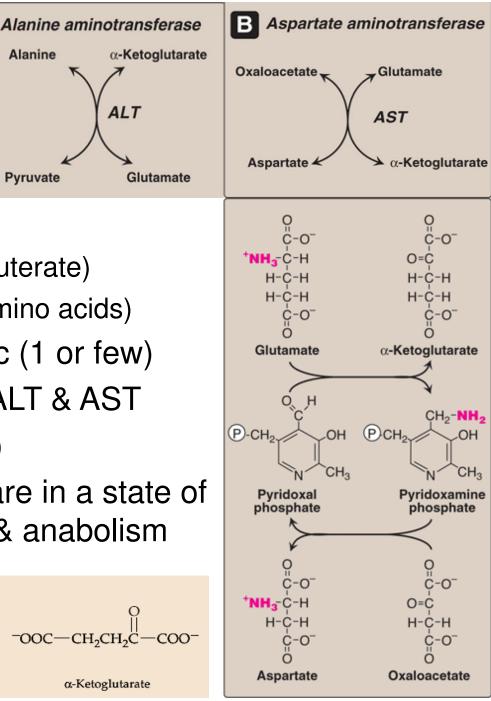
Pvruvate

ALT

Glutamate

- Aminotransferases are specific (1 or few)
- The two most important are; ALT & AST
- All require PLP coenzyme (B_6)
- Aminotransferases reactions are in a state of equilibrium, aid in catabolism & anabolism

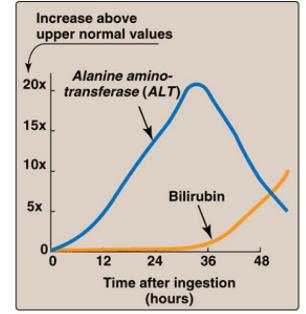


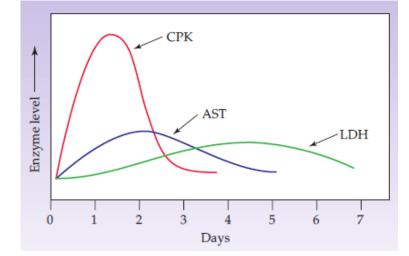


Diagnostic value of plasma aminotransferases

AST & ALT

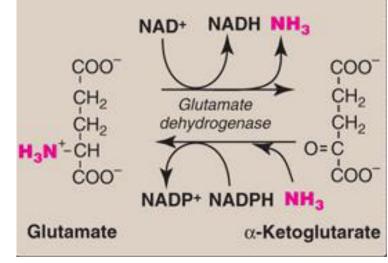
- Conditions that cause extensive cell necrosis
 - Viral hepatitis, toxic injury, & prolonged circulatory collapse
- ALT is more specific for liver disease
- AST is more sensitive for liver disease
- Elevated serum bilirubin
- Nonhepatic disease: Aminotransferases may be elevated but they are clinically different





Oxidative deamination of amino acids: Glutamate DH

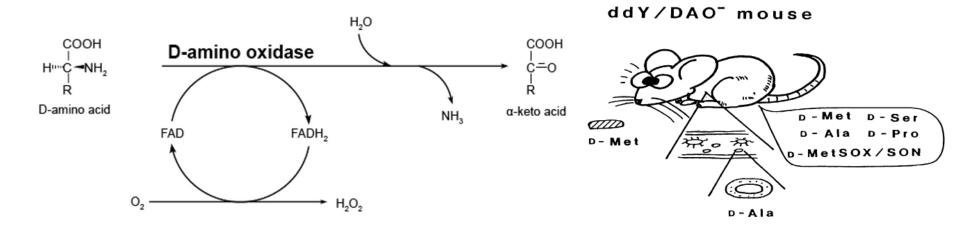
- The result is free amonia
- Glutamate is the only amino acid that undergoes rapid oxidative deamination
 - Amino groups of most aa. can be released as ammonia
- Occurs primarily in liver & kidney
- Provides α-ketogluterate (energy) & ammonia (urea)
- Use either NAD⁺ or NADP⁺ as a coenzyme



- ADP is an allosteric activator
- The direction depends on: conc. of glutamate, α-ketoglutarate, & ammonia, and the ratio of oxidized to reduced coenzymes

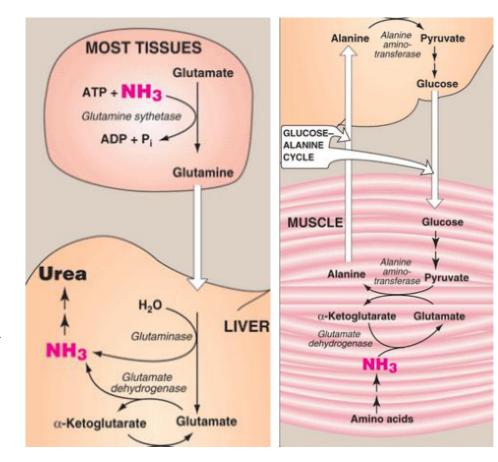
Oxidative deamination of amino acids: D-amino acid Oxidase

- Plants & microorganisms
- FAD-dependent peroxisomal enzyme
- Catalyzes oxidative deamination
- Results in α-keto acids that can either be reaminated to Lisomers, or catabolized for energy



Transport of ammonia to the liver

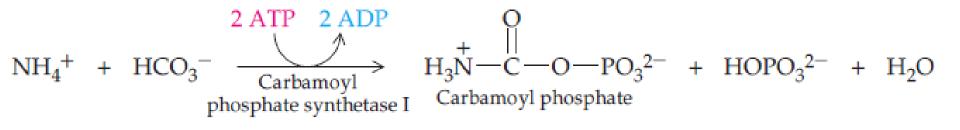
- Two mechanisms;
- I. Glutamine synthetase
 - ✓In most tissues → blood → liver (glutaminase)
- 2. Glucose-alanine cycle,
 ✓ Primarily by muscle
 - ✓ Pyruvate transamination (alanine) → blood → liver →
 - oTransamination (puruvate)
 - Gluconeogenesis
 (glucose), back to muscle



The Urea Cycle

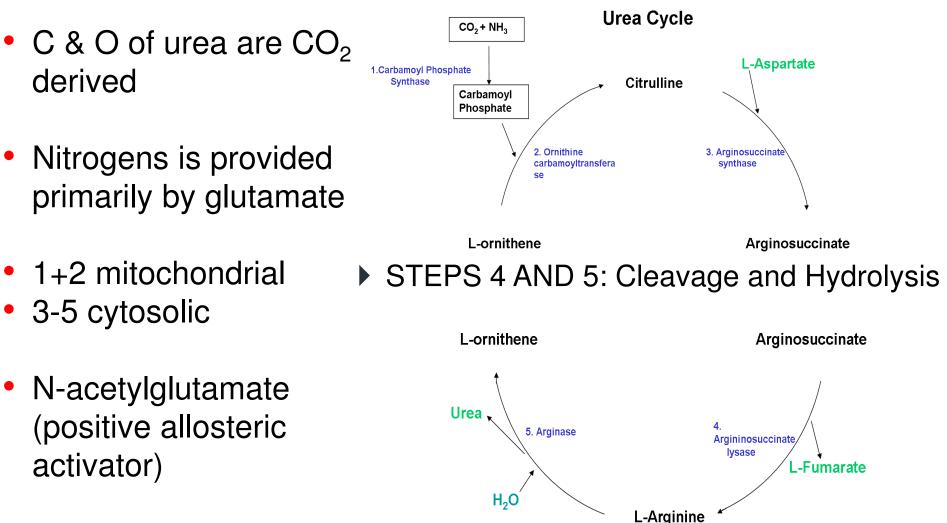
- What is it? Why do we need it? What fish do? Is it applicable in mammals? What mammals do?
- Where does it occur? Where does it go? Is energy needed?
- Accounts for 90% of the nitrogencontaining components of urine

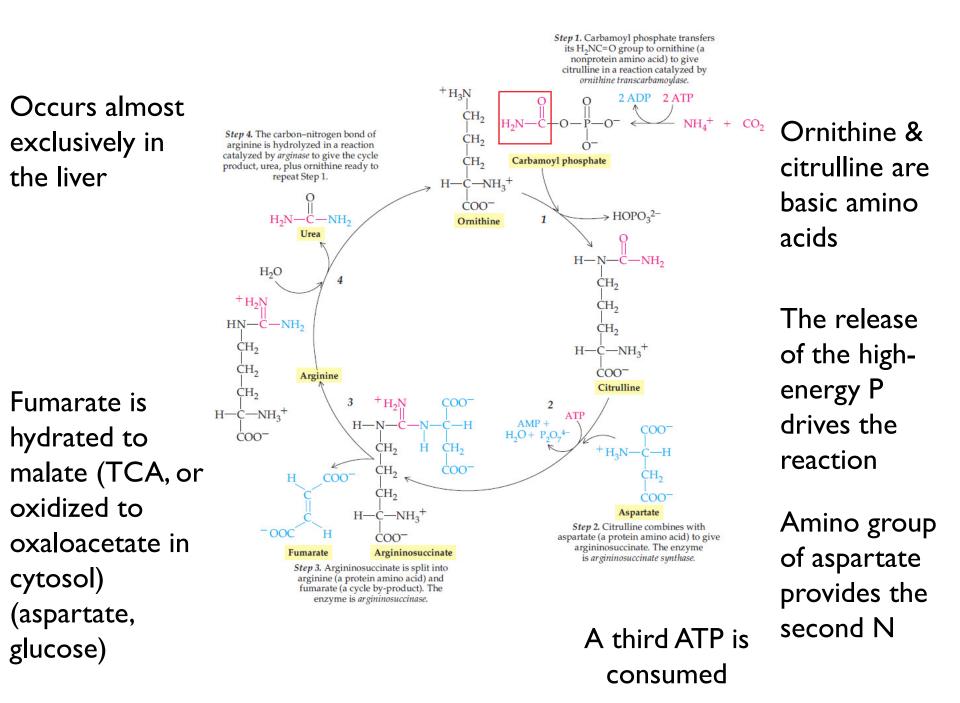




UREA CYCLE

STEPS 2 AND 3: Building Up a Reactive Intermediate





Net Result of the Urea Cycle

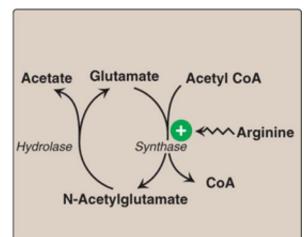
 $Asp + NH_3 + CO_2 + 3ATP \rightarrow Urea + Fumarate + 2ADP + AMP + 3H_2O$

- Breaking of four high-energy phosphate bonds (large –ΔG, irreversible)
- Production of fumarate
- Glutamate is the immediate precursor of both ammonia (through oxidative deamination by glutamate dehydrogenase) & aspartate nitrogen (through transamination of oxaloacetate by AST)
- Formation of urea from the C of CO₂, NH₄⁺, and aspartate, followed by biological elimination through urine
- Small portion to the intestine, cleaved to CO₂ and NH₃ by bacterial urease, feces and blood, kidney failure, hyperammonemia, neomycin

Regulation of the Urea Cycle

 $Asp + NH_3 + CO_2 + 3ATP \rightarrow Urea + Fumarate + 2ADP + AMP + 3H_2O$

- Regulation occurs at the level of the ratelimiting step (CPSI)
- N-Acetylglutamate is synthesized by Nacetylglutamate synthase, arginine is an activator (fed state)



$$NH_4^+ + HCO_3^- \xrightarrow[phosphate]{2 ATP 2 ADP} \xrightarrow[phosphate]{0} + H_3^{O} + H_3$$

Metabolism of Ammonia

- Ammonia is produced by all tissues, slightly elevated concentrations (hyperammonemia) are toxic to CNS
- Sources of ammonia: amino acids (food and transamination with oxidative deamination), kidneys (renal glutaminase and glutamate dehydrogenase), Intestinal glutaminase, bacterial urease, catecholamines, nitrogenous bases
- Transport of ammonia: very low levels in the blood, rapid action of liver, alanine or glutamine (primarily in muscle, liver and brain, glutamine synthetase), deaminated by glutaminase
- Disposal of ammonia: urea in liver, to kidneys, urine

Diseases

- Hyperammonemia:
 - Normal serum ammonia (5–50 µmol/L)
 - Liver compromised (genetic or acquired): can reach 1000 µmol/L)!
 - Medical emergency (CNS toxicity)
- Acquired: either acute (viral hepatitis, ischemia, or hepatotoxins), cirrhosis (alcoholism), or chronic hepatitis
- Hereditary: five enzymes, (1:30,000 live births), autosomal recessive. Ornithine transcarbamoylase deficiency (X-linked, most common, males)
- Immediate treatment (hemodialysis) vs. long-term treatment (low-protein diet and frequent small meals)