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Fatty Acids

Monday, January 14, 2008

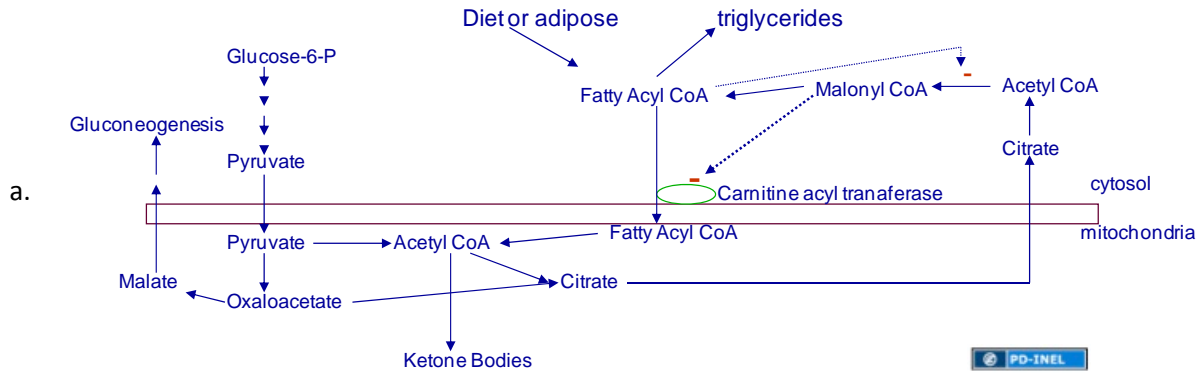
9:30 AM

1. Why does glucagon stimulation of liver cause an increase in acetyl-CoA levels in mitochondria?
 - a. Drop in blood glucose leads to glucagon levels increasing, insulin levels decreasing
 - b. Gluconeogenesis is stimulated
 - c. FA are released from adipose
 - d. Increase in FA in liver
 - e. FA oxidation increases
 - f. Increase in gluconeogenesis leads to decrease in overall TCA cycle
 - g. Increase in Acetyl CoA concentration because production exceeds TCA cycle capacity
2. What are the two ketone bodies produced by liver? How is acetoacetyl-CoA formed in the liver? What controls the conversion of acetoacetyl-CoA to HMG-CoA?
 - a. Acetoacetate + NADH \rightarrow beta-hydroxybutyrate (+NAD⁺)
 - b. Mechanism
 - i. Tholase: 2 Acetyl-CoA \rightarrow acetoacetyl-CoA
 - ii. HMG CoA synthase: Acetoacetyl-CoA + H₂O + Acetyl-CoA \rightarrow HMG-CoA
 - 1) Inhibited by succinyl-CoA
 - 2) Mitochondrial
 - iii. HMG-CoA Lyase: HMG-CoA \rightarrow Acetoacetate + Acetyl-CoA
 - c. HMG CoA Synthase is inhibited by succinyl-CoA
3. Why does an increase in the flux of the TCA cycle result in a decrease in ketone body synthesis?
 - a. Increased TCA cycle \rightarrow increased succinyl-CoA \rightarrow inhibition of HMG-CoA Synthase
 - b. Signal to increase gluconeogenesis \rightarrow shift away from TCA \rightarrow decrease in succinyl-CoA
4. How does an increase in an acetyl-CoA levels lead to an increase in formation of HMG-CoA?
 - a. Acetyl-CoA reverses inhibition of HMG-CoA Synthase by succinyl-CoA
 - b. Ketogenesis increased in fasting, prolonged exercise, high fat diet, fetal suckling
 - c. Acetyl-CoA increases transcription of HMG-CoA synthase gene; insulin can reverse increase
5. How are acetoacetate and B-hydroxybutyrate metabolized in muscle and brain?
 - a. Beta-hydroxybutyrate dehydrogenase: beta-hydroxybutyrate + NAD⁺ \rightarrow acetoacetate + NADH
 - b. 3-ketoacyl-CoA transferase: Acetoacetate + Succinyl-CoA \rightarrow Acetoacetyl-CoA + Succinate
 - c. Thiolase: Acetoacetyl-CoA + CoA \rightarrow 2 Acetyl-CoA
6. What enzyme is present in muscle but absent from liver?
 - a. 3-ketoacyl-CoA transferase
 - b. Absence of enzyme prevents futile cycle
7. Why do ketone bodies increase in an uncontrolled diabetic? What is the relationship between liver and adipose tissue?
 - a. Lack of insulin \rightarrow glucagon secretion \rightarrow
 - i. High blood glucose
 - ii. Increase in glucose synthesis
 - iii. Increase in FA release from adipose
 - b. Liver
 - i. Fatty acids \rightarrow acetyl-CoA \rightarrow ketone bodies
 - ii. Acetyl-CoA + OAA \rightarrow citrate is decreased
 - iii. Leads to even more glucose production
 - c. Muscle
 - i. Metabolism of ketone bodies decreased
 - ii. Lack of insulin decreases GLUT4 transport \rightarrow no source of OAA to combine w/ Acetyl-CoA for use in TCA cycle
 - d. Starvation
 - i. Increase in gluconeogenesis, release of FA from adipose, ketone body production
 - ii. Depletion of TCA intermediates in muscle

- iii. Reduces capacity of muscle to use acetyl CoA
 - iv. Ketone bodies increase in blood
 - v. Ketoacidosis
 - 1) Not as severe as in diabetics b/c there is still some insulin to FA release from adipose is moderated
 - 2) Glucose can still enter muscle to some extent allowing greater operation of TCA cycle and better use of Acetyl CoA
8. What tissues are most active in the synthesis of fatty acids?
- a. Liver and adipose tissue
 - b. All tissues synthesize FA
9. How is acetyl-CoA transported out of mitochondria for fatty acid synthesis?
- a. Citrate synthase: Acetyl-CoA + OAA \rightarrow citrate
 - b. Carried through citrate transporter on inner mito membrane
 - c. Citrate lyase: citrate + CoA + ATP \rightarrow Acetyl-CoA + OAA + ADP
 - d. Malate dehydrogenase: OAA + NADH + H⁺ \rightarrow Malate + NAD⁺
 - e. Malic Enzyme: Malate + NADP⁺ \rightarrow Pyruvate + NADPH + H⁺ + CO₂
 - f. Pyruvate is exported back into mitochondria via pyruvate transporter
 - g. Pyruvate + CO₂ + ATP \rightarrow OAA + ADP + Pi
10. How is NADPH generated for use in fatty acid biosynthesis?
- a. Generated in conversion of malate to pyruvate
 - b. Used in reduction step of FA synthesis
11. What is the sequence of reaction carried out by the fatty acid synthase complex? How is malonyl-CoA used? When does acetyl-CoA enter the sequence? What is required for the reduction processes? Is CO₂ incorporated into fatty acids?
- a. Acetyl CoA Carboxylase: Acetyl-CoA + ATP + HCO₃⁻ \rightarrow Malonyl-CoA + ADP + Pi + H⁺
 - i. Biotin cofactor carries CO₂ from HCO₃⁻ and attaches it to acetyl CoA
 - ii. Regulated (see question 12)
 - b. FA Synthase complex carries out steps of synthesis ketoacyl-ACP synthase
 - i. Acetyl-CoA binds to cysteine -SH on ketoacyl-ACP synthase
 - ii. Malonyl-CoA binds to phosphopantetheine on ACP
 - iii. Condensation reaction between malonyl and acetyl, releases CO₂
 - iv. Reduction using NADPH
 - v. Dehydration
 - vi. Reduction using NADPH
 - vii. Saturated acyl lengthened by two carbons
 - c. Requires pantothenic acid, NADPH, biotin
 - d. CO₂ is not incorporated into FA
 - e. Once at palmitic acid, chain is released by thioesterase
12. How is acetyl-CoA carboxylase regulated? What are the effects of citrate and palmitoyl-CoA? What is the role of phosphorylation and how is it regulated?
- a. Begins as protomer (several subunits)
 - b. Citrate causes polymerization of subunits into long chain, malonyl-CoA and palmitoyl-CoA inhibit polymerization
 - c. Phosphorylation inactivates polymer
 - i. Palmitoyl CoA \rightarrow AMP Kinase \rightarrow inactive carboxylase
 - 1) AMPK activated via phosphorylation
 - 2) AMP and Acyl-CoA activate AMPK Kinase
 - 3) Dephosphorylated/inactivated by PPP
 - ii. Glucagon \rightarrow PKA \rightarrow inactive Acetyl CoA Carboxylase
 - iii. Dephosphorylated by PPP
 - d. Transcriptional regulation
 - i. High carb diet \rightarrow increased transcription of gene \rightarrow increased FA synthesis
 - ii. High fat diet \rightarrow reduced transcription of gene, b/c fat in diet is same as what we

synthesize

13. How are fatty acid oxidation and fatty acid synthesis coordinately regulated?



- b. Glucagon --> increase gluconeogenesis --> decrease in citrate --> decreased malonyl CoA --> increase in carnitine acyl transferase --> increase in FA ox --> increase in ketone bodies
 Glucagon --> increase in cAMP --> increase in PKA --> inactivated ACC --> decrease in malonyl CoA
- c. Increase in FA --> decrease in malonyl CoA --> increase in FA transport --> increase in FA Ox
- d. Insulin --> increase in ACC --> increase in malonyl CoA --> inhibition of FA transport --> decrease FA Ox --> increase FA synthesis
- e. Increase in AMP --> increase in AMPK --> inactive ACC --> decrease in malonyl-CoA and FA synthesis --> increase in FA transport and FA Ox