

Cardiac metabolism

- Heart is **highly oxidative** (elevated rate of ATP hydrolysis by Oxidative phosphorylation), there is a fine equilibrium between the work the heart has to do and the energy to be synthesized.
- Metabolism of heart is designed to generate a large amount of ATP by oxidative phosphorylation

▼ Three key components of cardiac metabolism

1. Capture and utilization of substrates to use them in the TCA cycle (citric acid/kreb's)
2. Oxidative phosphorylation in mitochondria
3. Phosphocreatine-creatine kinase system (transferring phosphate from ATP to creatine to make PC (phosphocreatine which is a source of energy under high demand))

- Metabolism of heart can use O₂ up to 80-90% of ETC capacity, but at resting state, heart operates at 15-20% of maximum oxidative capacity

▼ Energy sources of heart under basal aerobic conditions

1. **60% from Fatty acids**
 - Synthesising capacity of fatty acids in heart is low
 - heart depends on influx of fatty acids from blood vessels
 - So, rate of FA consumption is determined by concentration of non-esterified fatty acids in plasma
2. **35% from carbohydrates**
3. **5% from amino acids and ketone bodies**

▼ Uses of energy (ATP) in heart

- 60-70% used for **muscle contraction**
- 30-40% for **Ca²⁺ ATPase** in sarcomplasmic reticulum and **other pumps**

▼ General steps of metabolic pathways in heart

- 10-40%: Oxidative decarboxylation of **pyruvate in glycolysis** → Acetyl CoA (discussed below)
- 60-90%: **B-oxidation of fatty acids** → Acetyl CoA (discussed in lecture 2)
- Acetyl CoA → Citric acid cycle → Forms NADH and FADH₂ → Deliver electrons to electron transport chain → ATP

▼ Carbohydrate metabolism

- Depends on glycogen stores and exogenous glucose
 - **Glycogen stores are small** (30mmol/g)
 - **Exogenous glucose is transported by GLUT4** (found in sarcolemma and translocated to membrane in response to Insulin, Work demand, and Ischemia) **GLUT1** plays an accessory role
 - Factors regulating transport: Transmembrane gradient of glucose + Content of GLUT4

▼ **Glycolytic pathway** (Glucose-6-phosphate + NAD⁺ → Pyruvate and NADH)

- Generates **2 ATP per glucose**
- In Anaerobic conditions: Pyruvate → Lactic acid
- In Aerobic conditions: Pyruvate + NADH → CO₂ + NAD⁺ (in mitochondrial matrix)

- Only 2% of ATP is produced in glycolytic pathway, but it's important in Anerobic or ischemic status
- In heart failure and hypertrophy, shift of metabolism towards carbohydrates not FAs

▼ Regulatory steps

▼ Glyceraldehyde-3-phosphate dehydrogenase (GAPDH)

- Major regulatory step
- Produces NADH
- Inhibited by excess NADH (-ve feedback), Activated by NAD+
- Accumulation of NADH and lactate in ischemia → Stop oxidative metabolism and lactate production

+ ☹ ▼ PFK-1: KEY regulatory enzyme

- Catalyzes second irreversible step
- Activated by low forms of energy (ADP, AMP, Pi (inorganic phosphate)) + Fructose 2,6-bisphosphate (formed by PFK-2)
- Inhibited by high forms of energy (ATP and low pH)

▼ Pyruvate fates in mitochondria

1. Oxidatively **decarboxylated** by Pyruvate dehydrogenase (PDH) → **Acetyl CoA**

- Control of PDH (mitochondrial multicomplex) activity is essential in control of glucose metabolism
- Controlled by work, substrate and hormones

2. **Carboxylated** by Pyruvate carboxylase → **oxaloacetate**

3. (in cytosol) **Reduced** by Lactate dehydrogenase → **Lactate**

- Lactate is then released into blood by transporter (this has a critical role in maintaining cellular pH)
- During starvation, lactate can be recycled into pyruvate (produces 2.5 ATP since it forms NADH) → This pyruvate is then burned aerobically in Citric acid cycle (produces 12.5 ATP per cycle)

▼ Fates of Glycolytic intermediates (pathways that don't make ATP)

▼ PPP pathway

- **Glucose-6-phosphate** from hexokinase reaction enters → **Produces NADPH** (in oxidative phase) + **5-carbon sugars** (in non-oxidative phase)
 - **NADPH** important for antioxidant effect (as it maintains level of reduced glutathione)
 - **Ribose-5-phosphate** (non-oxidative phase product) → substrate for nucleic acid synthesis
 - **Xylulose-5-phosphate** (non-oxidative product) → Transcriptional signaling molecule

▼ Polyol pathway

- G6P → Sorbitol, by aldose reductase
- Role of this pathway is **unknown**, but diabetic patients have increased flux to this pathway + Associated with abnormal glucose metabolism and cardiac dysfunction
- Increased aldose reductase: implicated in myocardial response to ischemia-reperfusion injury

▼ Hexosamine biosynthetic pathway

- Fructose-6-phosphate → Uridine diphosphate-N-Acetylglucosamine, by glutamine fructose-6-phosphate amidotransferase
- Product of this reaction (uridine...) Participates in O-linked glycosylation reactions of proteins
 - **Observed in diabetes**, could be responsible for alteration of insulin sensitivity and fatty acid oxidation