

# Aerobic Respiration

BCH 100 – Introductory Biochemistry · Dr. Radi

## By the end of this unit, you can...

- Explain the pyruvate dehydrogenase complex – the link reaction to acetyl-CoA – its cofactors and its allosteric + covalent regulation
- Walk all eight reactions of the TCA cycle with the running NADH/FADH<sub>2</sub>/GTP ledger, and explain its regulated, amphibolic, and anaplerotic roles
- Describe the pentose phosphate pathway – its oxidative and non-oxidative phases and its NADPH and ribose-5-phosphate products
- Trace electron flow through the four ETC complexes, driven by reduction potential, pumping protons to build the gradient
- Explain chemiosmosis and ATP synthase's rotary catalysis, the NADH shuttles, and the total ATP yield from one glucose
- Describe the regulation of respiration, uncouplers (thermogenin), and ETC poisons (cyanide, rotenone)

# Today's route



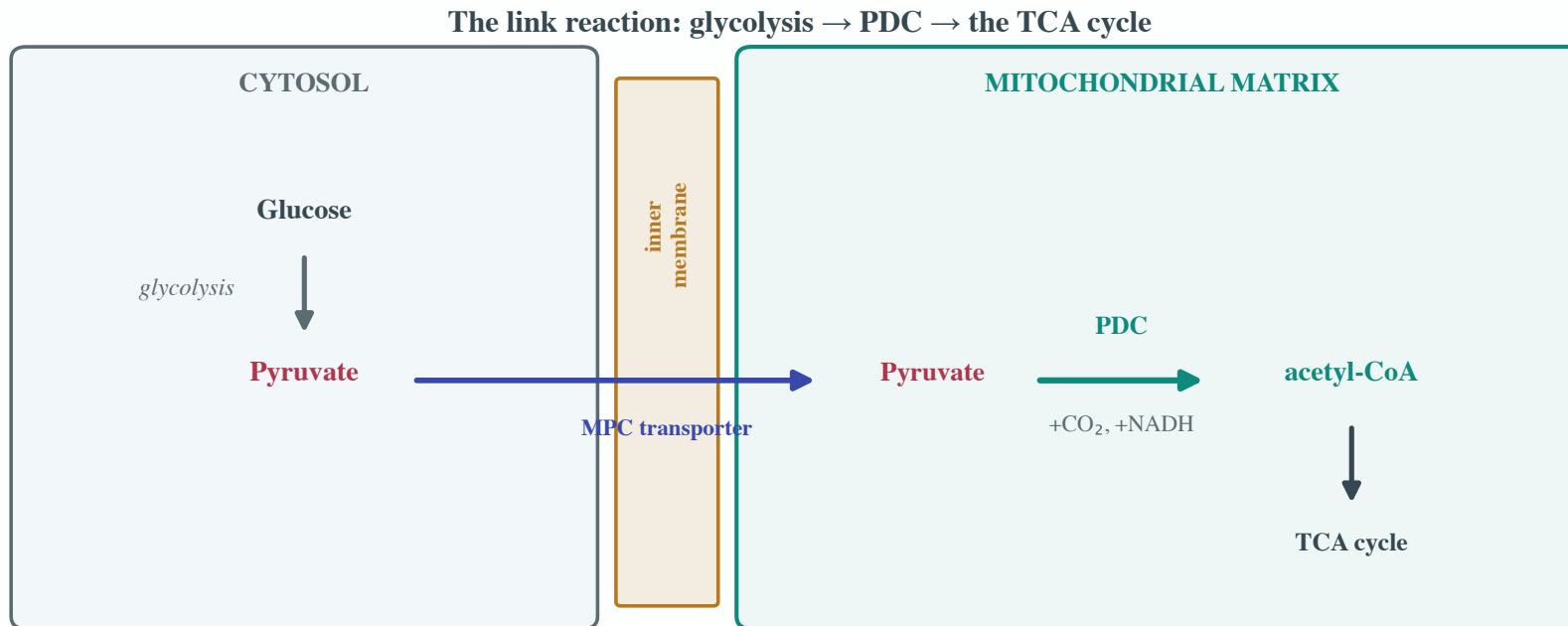
1. The Pyruvate Dehydrogenase Complex
2. The TCA Cycle
3. The Pentose Phosphate Pathway
4. The Electron Transport Chain
5. Oxidative Phosphorylation
6. Regulation, Poisons & Uncouplers

# 1 • The Pyruvate Dehydrogenase Complex

"The bridge from glycolysis to the TCA cycle – meet the giant complex that turns pyruvate into acetyl-CoA."

# The bridge to aerobic metabolism

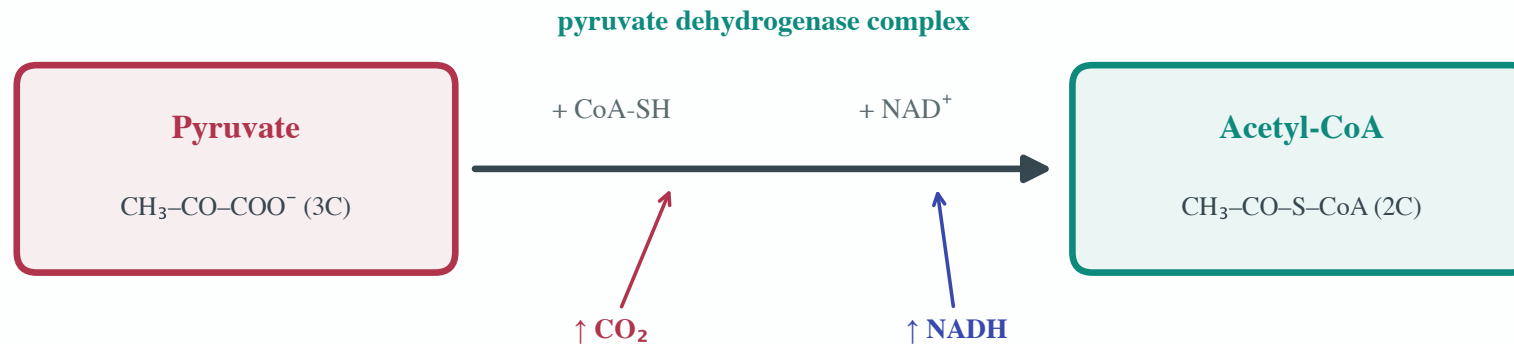
Glycolysis ends with **pyruvate** in the **cytosol** – but the real energy harvest happens in the **mitochondrion**. So pyruvate is shuttled across the inner membrane (by the **MPC transporter**) into the **matrix**, where both the **PDC** and the **TCA cycle** live. There the **pyruvate dehydrogenase complex** turns pyruvate into **acetyl-CoA** – the fuel the cycle actually burns.



# The link reaction

**Pyruvate + CoA + NAD<sup>+</sup> → acetyl-CoA + CO<sub>2</sub> + NADH.** In one move the PDC does three things: **chops off a carbon** (as CO<sub>2</sub>), **oxidizes** what's left (banking an **NADH**), and **attaches CoA** to make the reactive **acetyl-CoA**. It's **irreversible** – the true point of no return for glucose oxidation.

Oxidative decarboxylation — lose a carbon, bank an NADH, activate the rest as acetyl-CoA



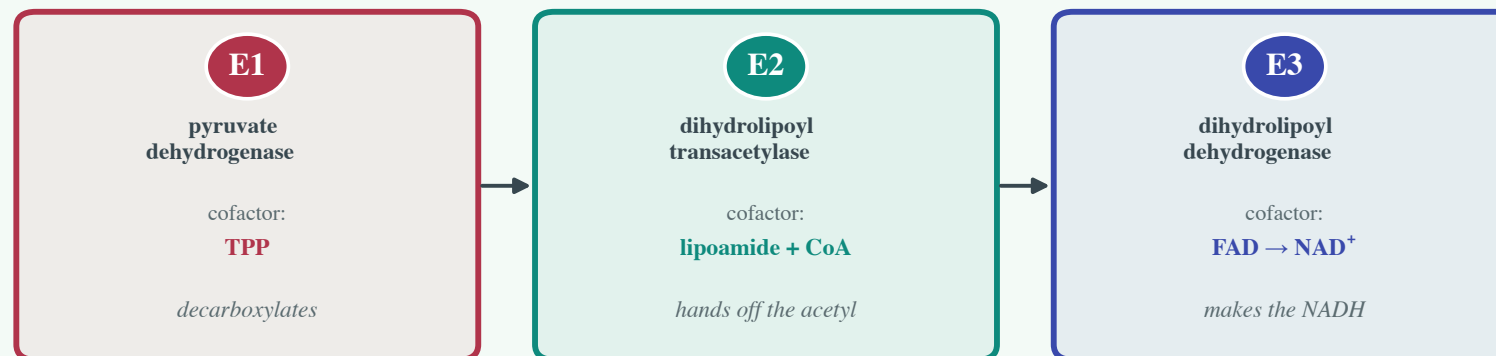
*Irreversible — the true committed step of glucose oxidation.*

# One complex, three enzymes, five cofactors

The PDC is a **giant** – three enzymes working as one, with **five** cofactors handing the substrate down an assembly line.

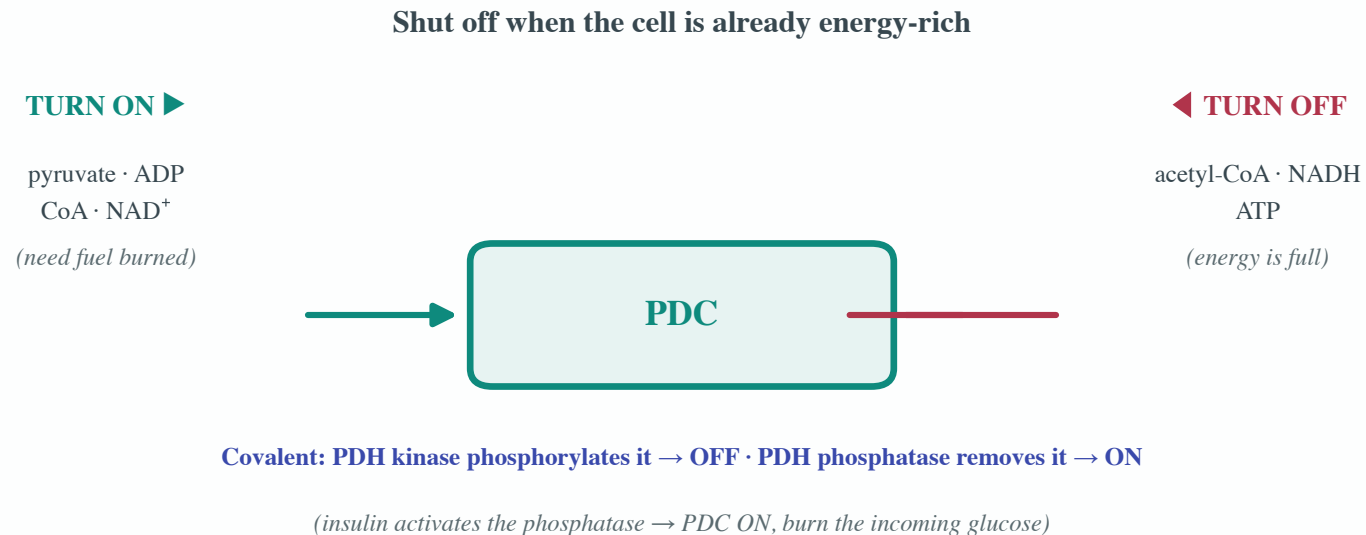
- **E1** (with **TPP**) decarboxylates pyruvate
- **E2** (with **lipoamide + CoA**) hands off the acetyl group
- **E3** (with **FAD → NAD<sup>+</sup>**) resets the machine and makes the **NADH**
- Remember the cofactors: "**Tender Loving Care For Nancy**" → **TPP · Lipoamide · CoA · FAD · NAD<sup>+</sup>**

One giant complex, three enzymes, five cofactors



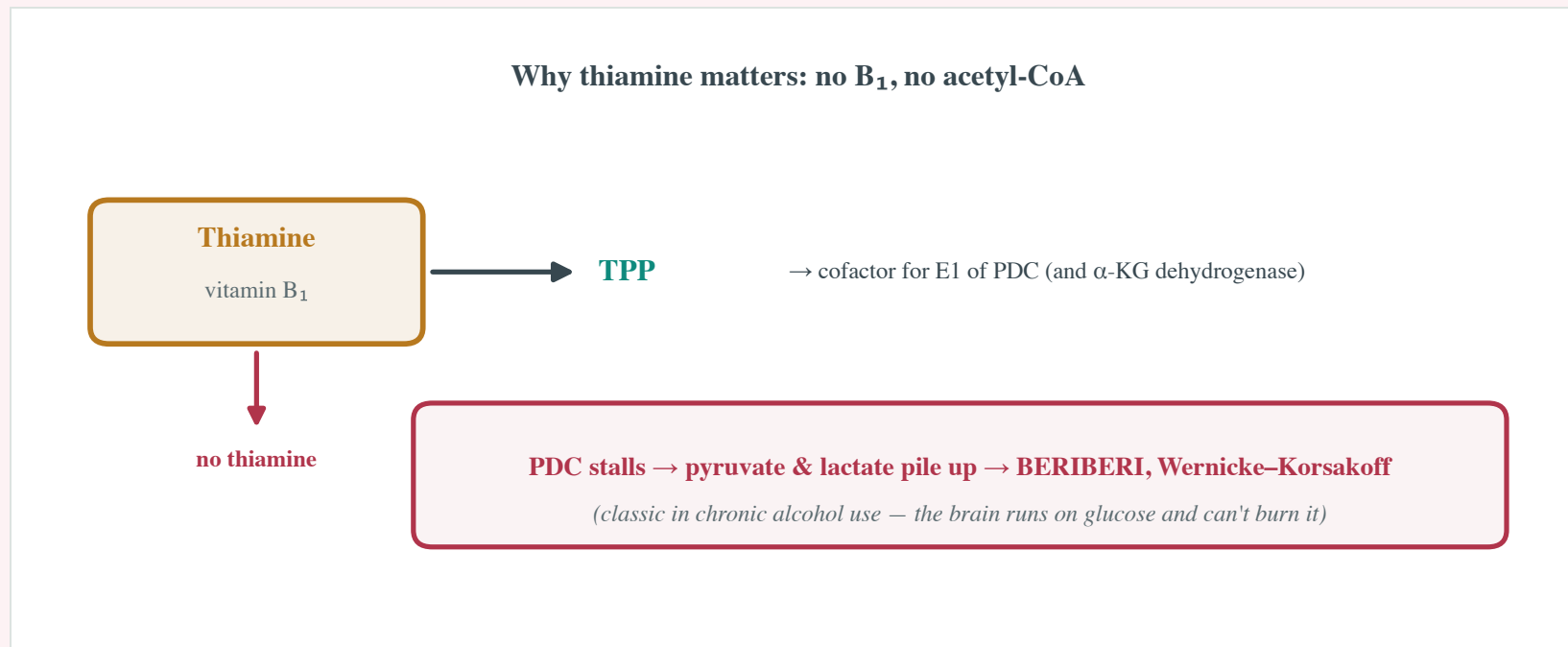
# Turned off when you're full

The PDC is a **committed, irreversible** step, so it's **tightly regulated** – and the theme is simple: **don't burn fuel you don't need**. Its own products (**acetyl-CoA, NADH**) and **ATP** shut it **off**; **pyruvate** and **ADP** turn it **on**. A **kinase/phosphatase** pair adds the covalent layer – **phosphorylation switches it OFF**, and **insulin** (fed) flips the phosphatase back **on** to burn the incoming glucose.



# No B<sub>1</sub>, no acetyl-CoA

**TPP** – the E1 cofactor – is made from **thiamine (vitamin B<sub>1</sub>)**. Without it, the PDC (and  $\alpha$ -ketoglutarate dehydrogenase) **stall**, so pyruvate and lactate **pile up**. The result is **beriberi** and **Wernicke-Korsakoff** syndrome – classically in **chronic alcohol use**, where the glucose-hungry brain suddenly can't burn its fuel.

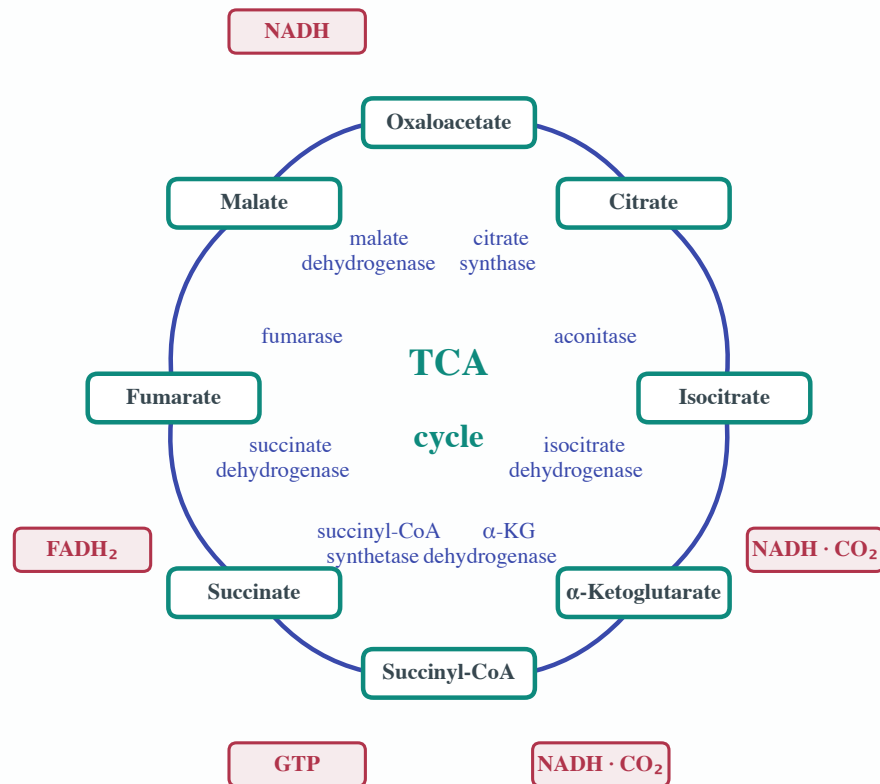


## 2 • The TCA Cycle

"NOT the Krebs's cycle – the TCA cycle! Walk all eight reactions, bank 3 NADH + 1 FADH<sub>2</sub> + 1 GTP per turn, and meet the hub of all metabolism."

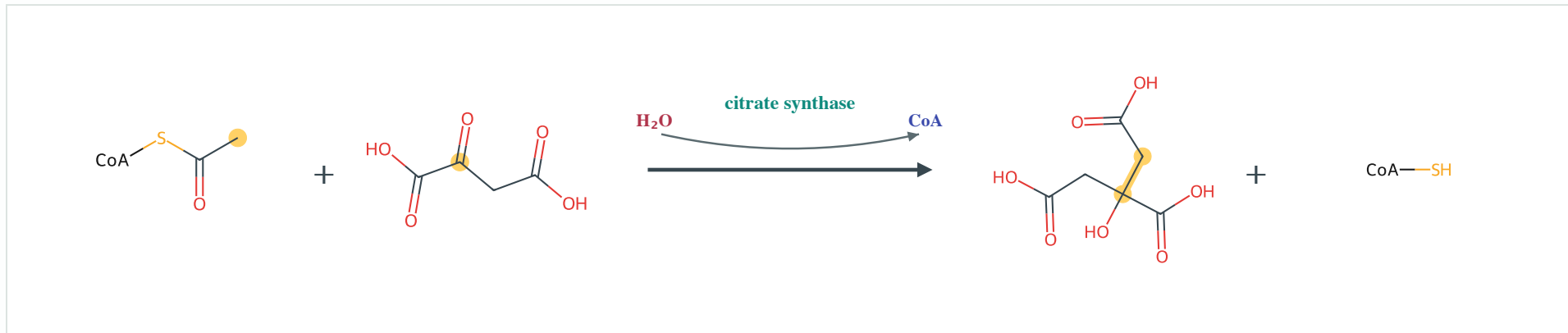
# NOT the Kreb's cycle – the TCA cycle!

Acetyl-CoA hands its **two carbons** to the cycle, which oxidizes them all the way to **two CO<sub>2</sub>** – and captures the energy as **reducing power**. Eight enzymes, one full turn, and oxaloacetate comes back out ready to grab the next acetyl-CoA. Watch what gets tapped off each step.



## Step 1: Citrate synthase

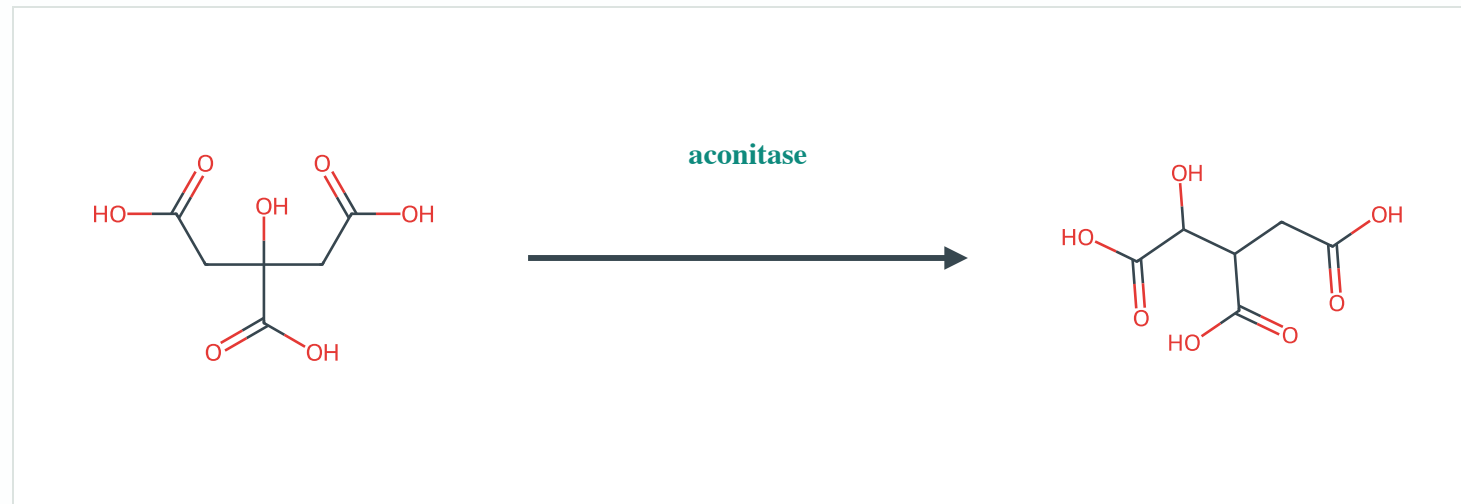
**Acetyl-CoA + oxaloacetate → citrate.** The 2-carbon acetyl group is welded onto the 4-carbon **oxaloacetate** to make the 6-carbon **citrate**, and the CoA is released. This first step is **irreversible** and a key **control point**.



no carriers yet • running: —

## Step 2: Aconitase

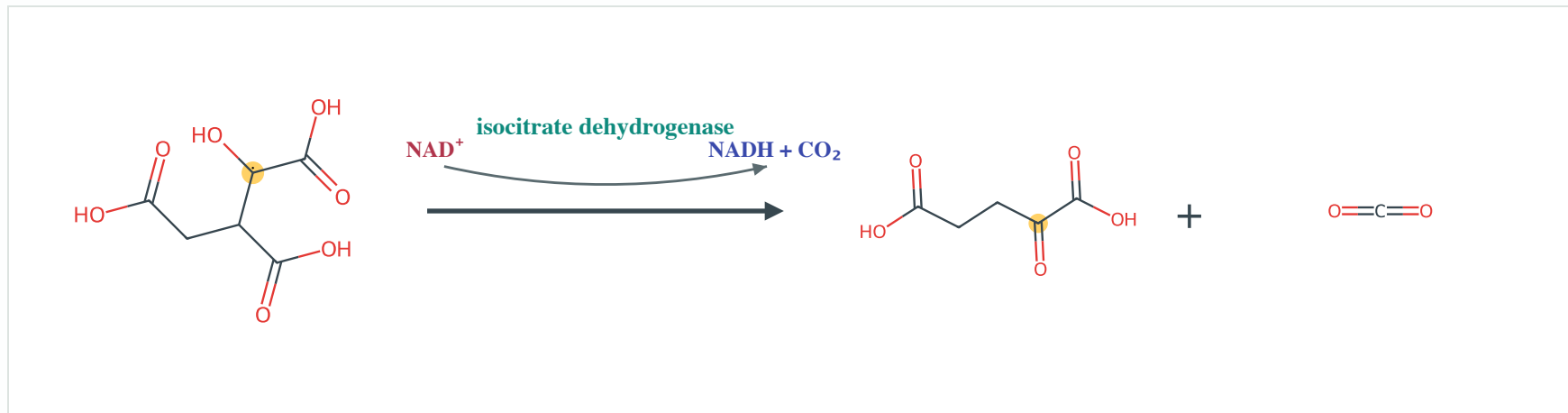
**Citrate → isocitrate.** Just a rearrangement – the **hydroxyl moves** to a neighboring carbon (via a **cis**-aconitate intermediate). Why? Citrate's -OH is on a carbon that can't be oxidized; isocitrate's can. Setting up the next step. **Reversible.**



no carriers yet • running: –

## Step 3: Isocitrate dehydrogenase

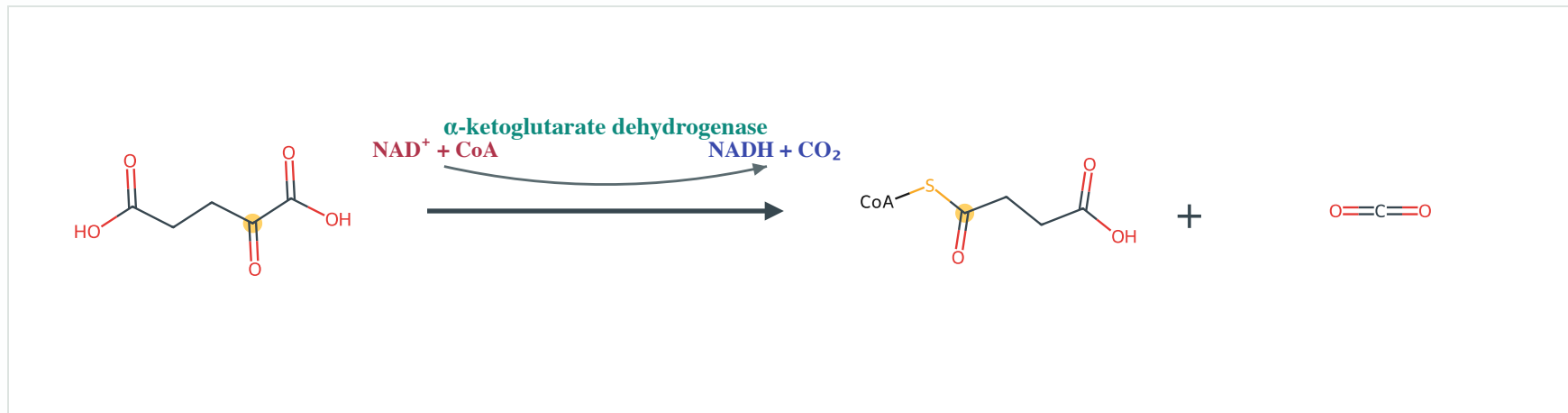
**Isocitrate** → **α-ketoglutarate** + **CO<sub>2</sub>**. The **first oxidation** and the **first CO<sub>2</sub> lost**: isocitrate is oxidized (making our **first NADH**) and immediately decarboxylated. This is the cycle's **rate-limiting, most-regulated** step.



**NADH +1 · CO<sub>2</sub> ↑ · running: 1 NADH**

## Step 4: $\alpha$ -Ketoglutarate dehydrogenase

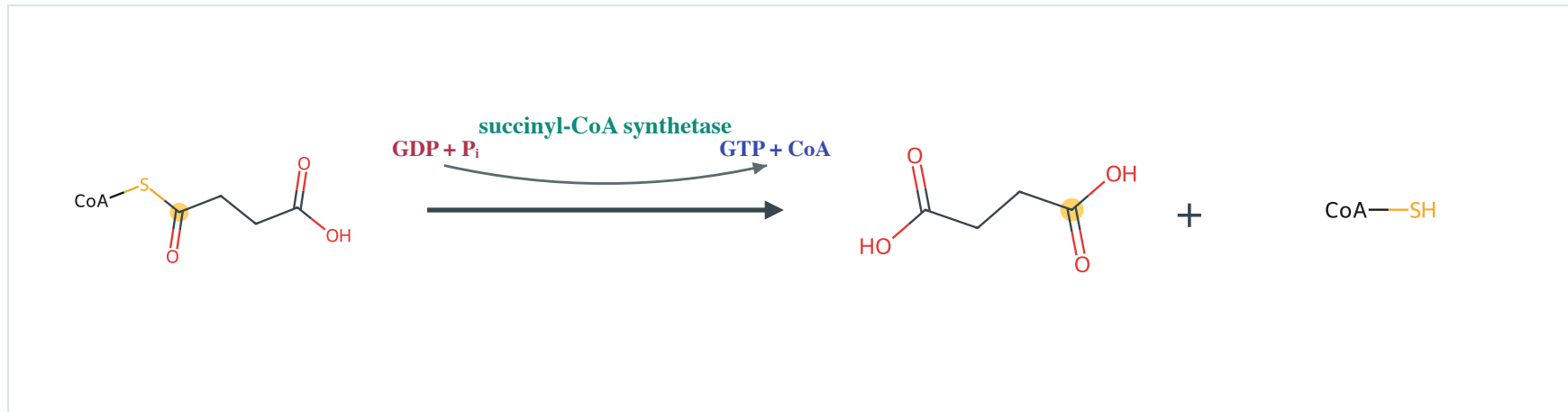
$\alpha$ -Ketoglutarate  $\rightarrow$  succinyl-CoA + CO<sub>2</sub>. The **second oxidative decarboxylation** – a giant complex just like the PDC (same 5 cofactors!). It banks a **second NADH**, releases the **second CO<sub>2</sub>**, and attaches CoA to make the high-energy thioester **succinyl-CoA**.



**NADH +1 · CO<sub>2</sub> ↑ · running: 2 NADH**

## Step 5: Succinyl-CoA synthetase

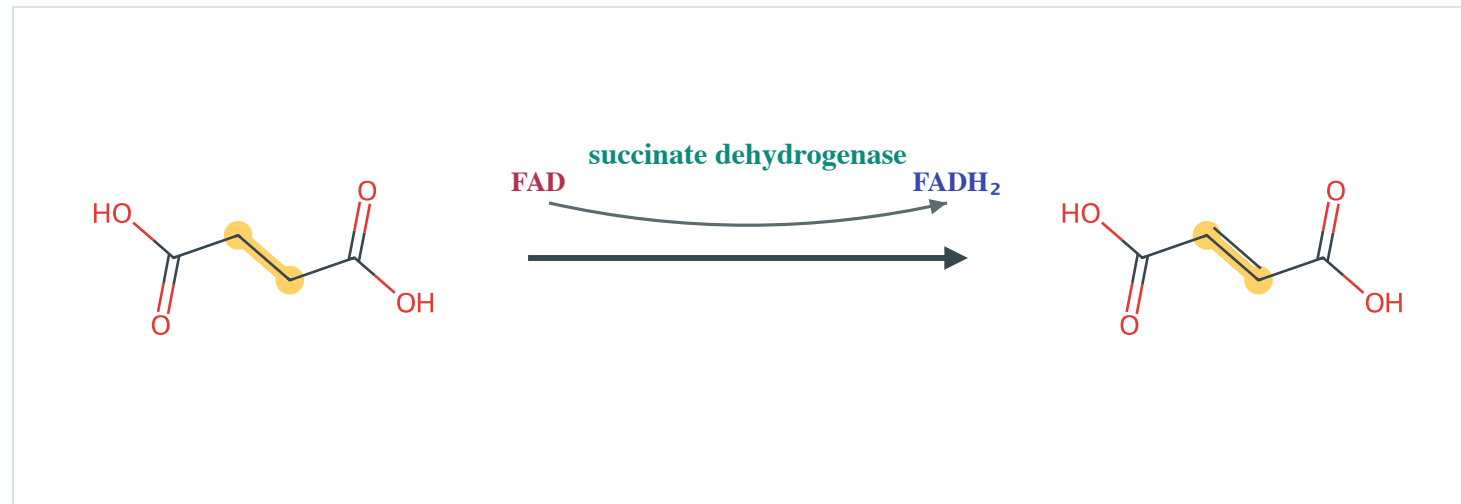
**Succinyl-CoA → succinate + GTP.** The thioester bond is **high-energy** – snapping it powers a **substrate-level phosphorylation**, making one **GTP** (= 1 ATP). This is the cycle's **only** direct high-energy phosphate. **Reversible.**



**GTP +1** • running: 2 NADH, 1 GTP

## Step 6: Succinate dehydrogenase

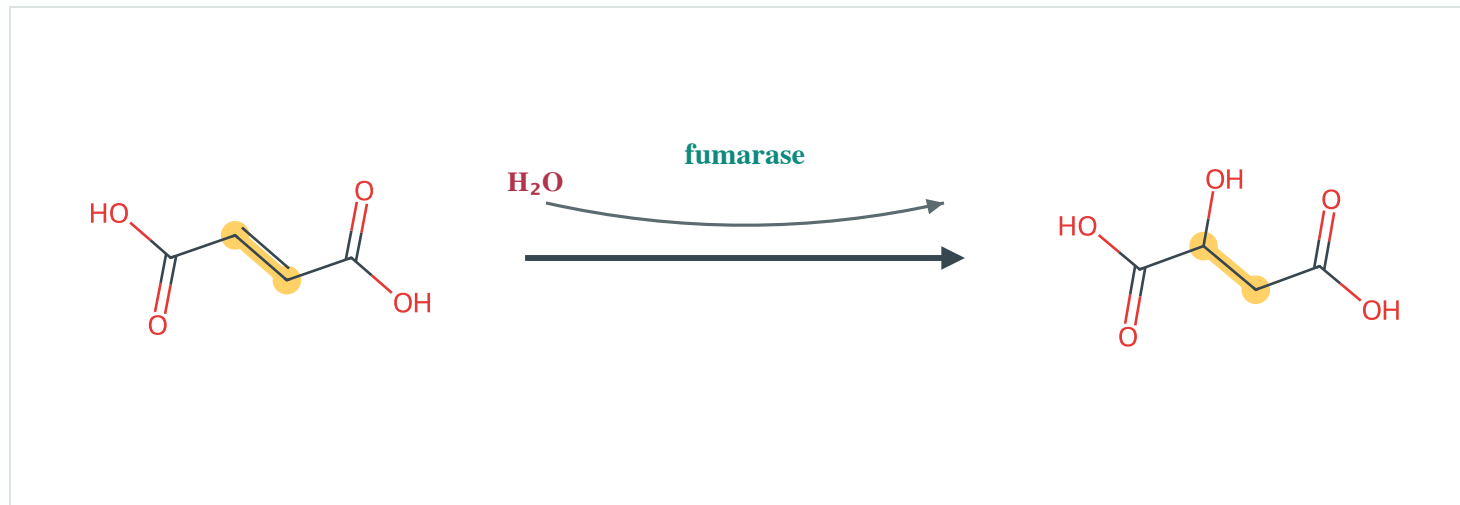
**Succinate → fumarate.** An oxidation that forms a **double bond** – and its electrons are a bit lower-energy, so they go to **FAD → FADH<sub>2</sub>** instead of NAD<sup>+</sup>. Fun fact: this enzyme is **embedded in the inner membrane** – it **is** Complex II of the electron transport chain!



**FADH<sub>2</sub> +1** • running: 2 NADH, 1 FADH<sub>2</sub>, 1 GTP

## Step 7: Fumarase

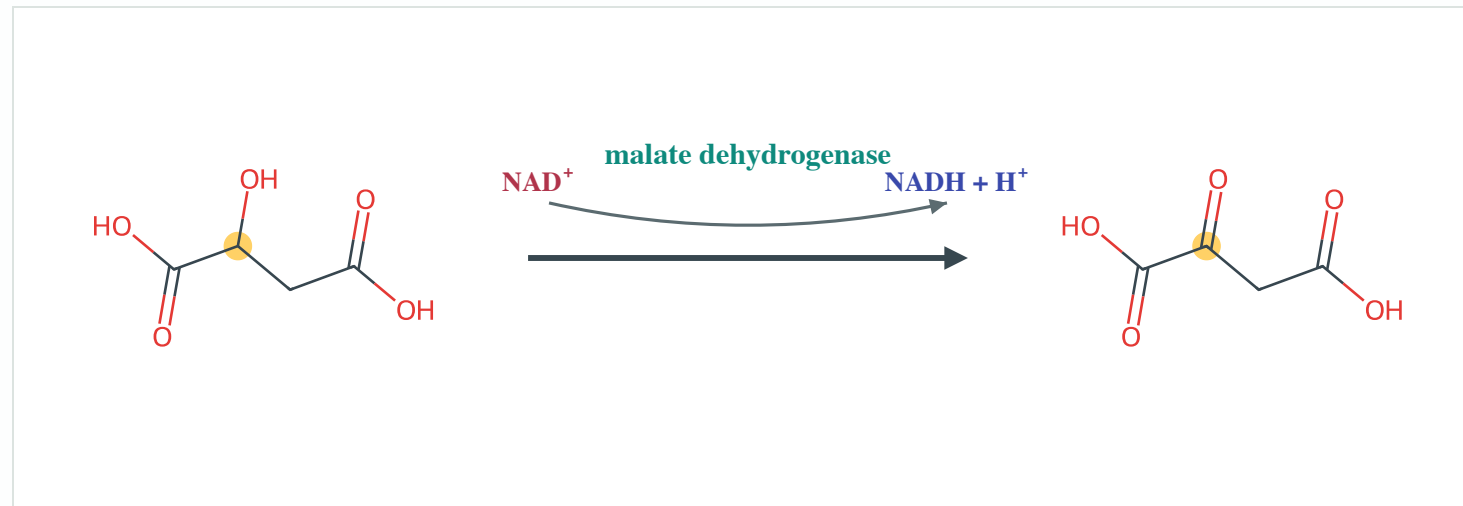
**Fumarate + H<sub>2</sub>O → malate.** A simple **hydration** – water adds across the double bond, putting a hydroxyl back on. No energy captured; we're setting up the final oxidation. **Reversible.**



**no carriers • running: 2 NADH, 1 FADH<sub>2</sub>, 1 GTP**

## Step 8: Malate dehydrogenase

**Malate → oxaloacetate.** The last oxidation banks a **third NADH** and regenerates **oxaloacetate** – ready to accept the next acetyl-CoA. The cycle is closed! Per turn we've made **3 NADH + 1 FADH<sub>2</sub> + 1 GTP** and released **2 CO<sub>2</sub>**.

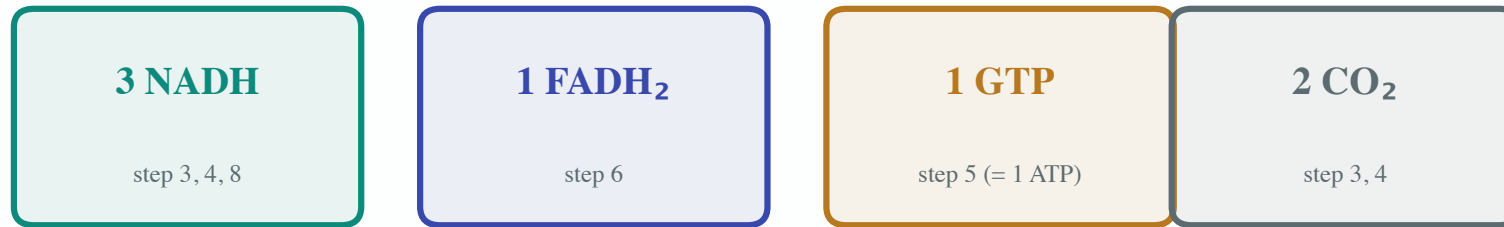


**NADH +1** • running: **3 NADH, 1 FADH<sub>2</sub>, 1 GTP** → per turn

# Tallying one turn

So one acetyl-CoA yields **3 NADH + 1 FADH<sub>2</sub> + 1 GTP**. The GTP is a **direct** ATP – but the real prize is the **reducing power**: those NADH and FADH<sub>2</sub> are worth **~9 more ATP** once they reach the electron transport chain. That's where aerobic metabolism gets its huge payoff.

**Per turn (one acetyl-CoA): 3 NADH + 1 FADH<sub>2</sub> + 1 GTP**

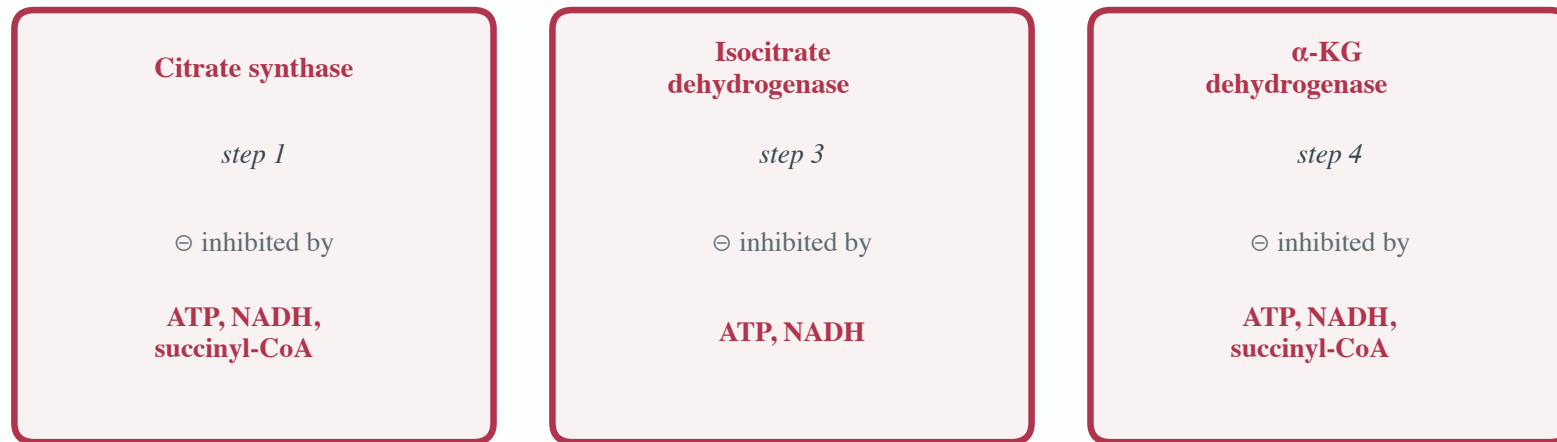


The GTP is direct ATP — but the real prize is the NADH & FADH<sub>2</sub>,  
cashied in for ~9 more ATP when they hit the electron transport chain.

# Regulation: read the energy charge

Three enzymes are the control points – **citrate synthase**, **isocitrate dehydrogenase**, and  **$\alpha$ -ketoglutarate dehydrogenase**. All are **slowed** when the cell is energy-rich (high **ATP**, **NADH**), and sped up by **ADP** and  **$\text{Ca}^{2+}$**  (the working-muscle signal). Don't oxidize fuel you don't need.

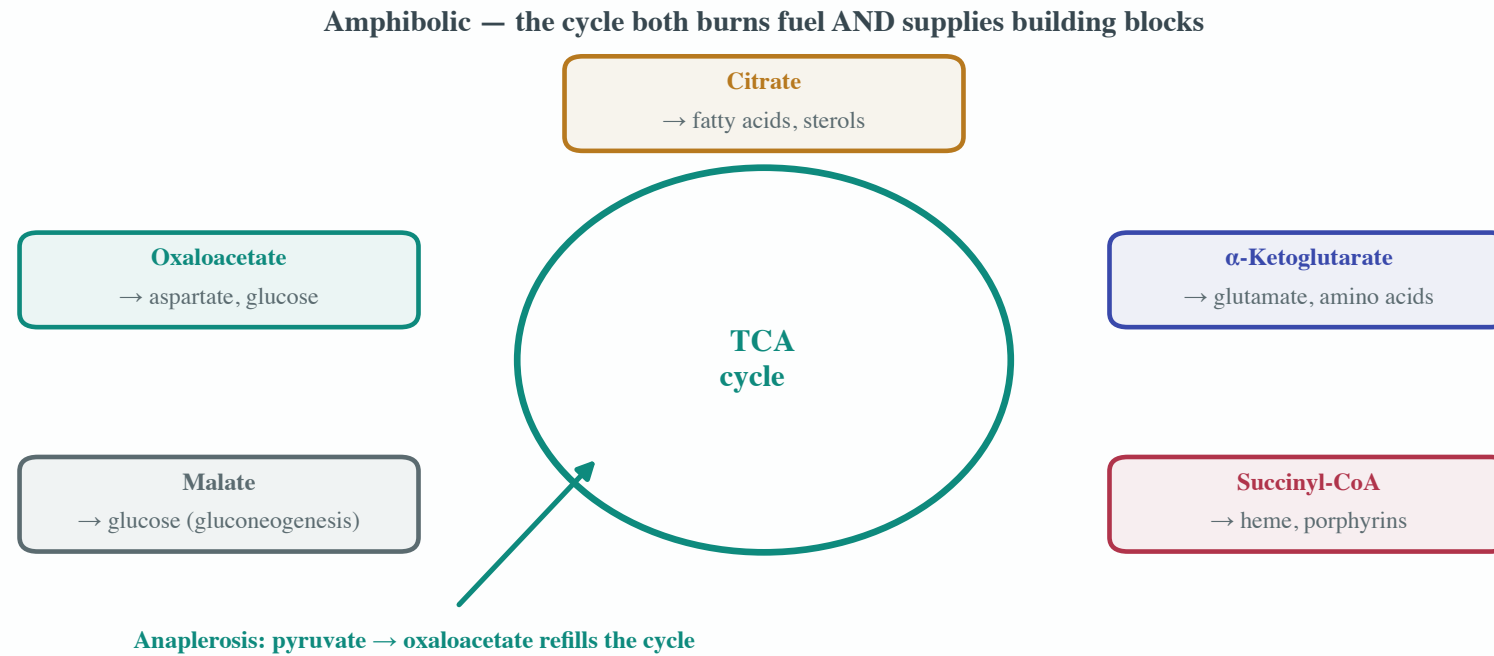
Slowed when the cell is energy-rich (high ATP / NADH)



*ADP and  $\text{Ca}^{2+}$  (a working-muscle signal) speed it back up.*

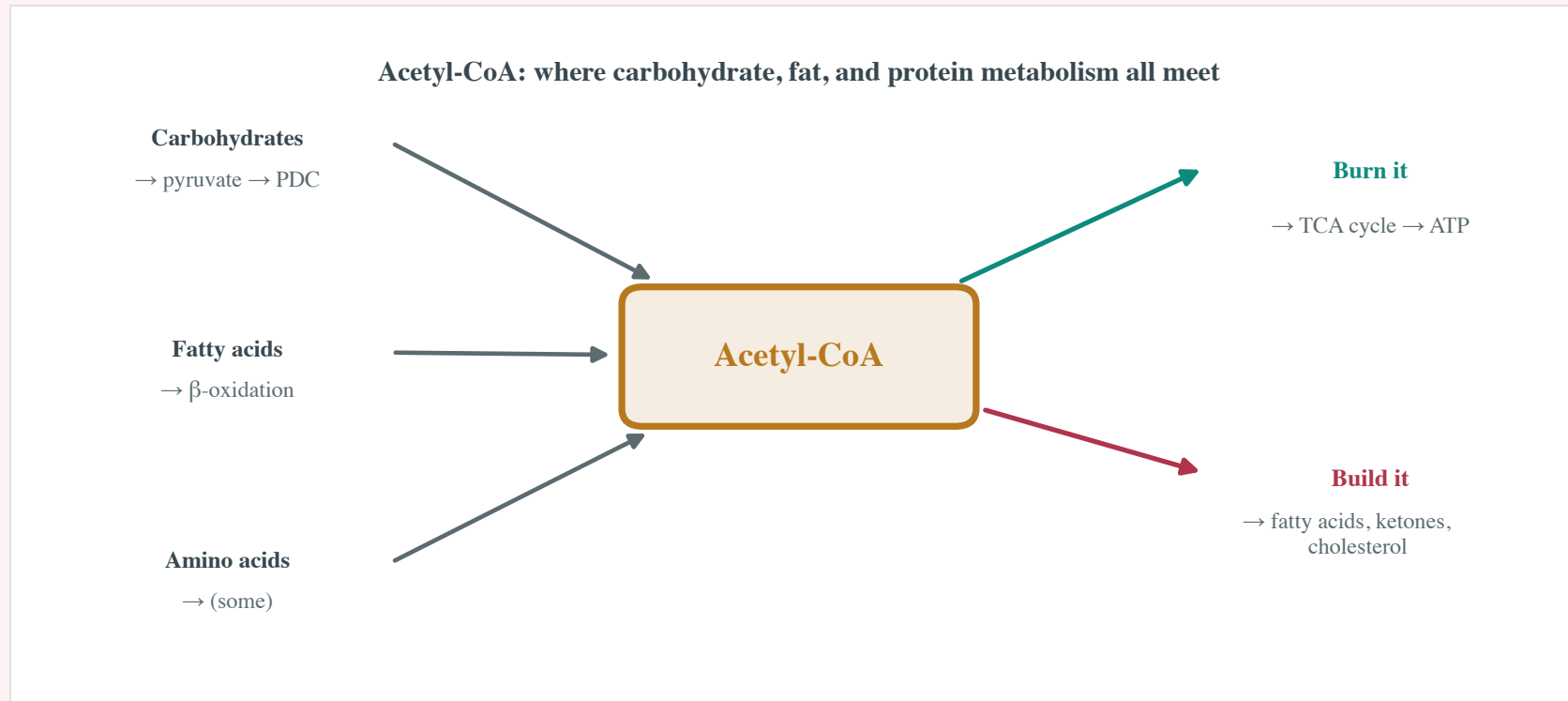
# Amphibolic – it builds as it burns

The TCA cycle isn't **only** for energy. Its intermediates get **pulled off for biosynthesis** – citrate → **fatty acids**,  $\alpha$ -ketoglutarate → **amino acids**, succinyl-CoA → **heme**. When they're drained, **anaplerotic** reactions top the cycle back up (pyruvate → oxaloacetate). It runs **both directions of metabolism at once**.



# Acetyl-CoA: the crossroads of everything

**Carbs, fats, and proteins all funnel into acetyl-CoA** – then it either **burns** in the TCA cycle or gets **built** into fatty acids, ketones, and cholesterol. One catch: because acetyl-CoA's two carbons **leave as CO<sub>2</sub>** each turn, you can't make **net** glucose from it – which is exactly **why you can't turn fat into sugar**.



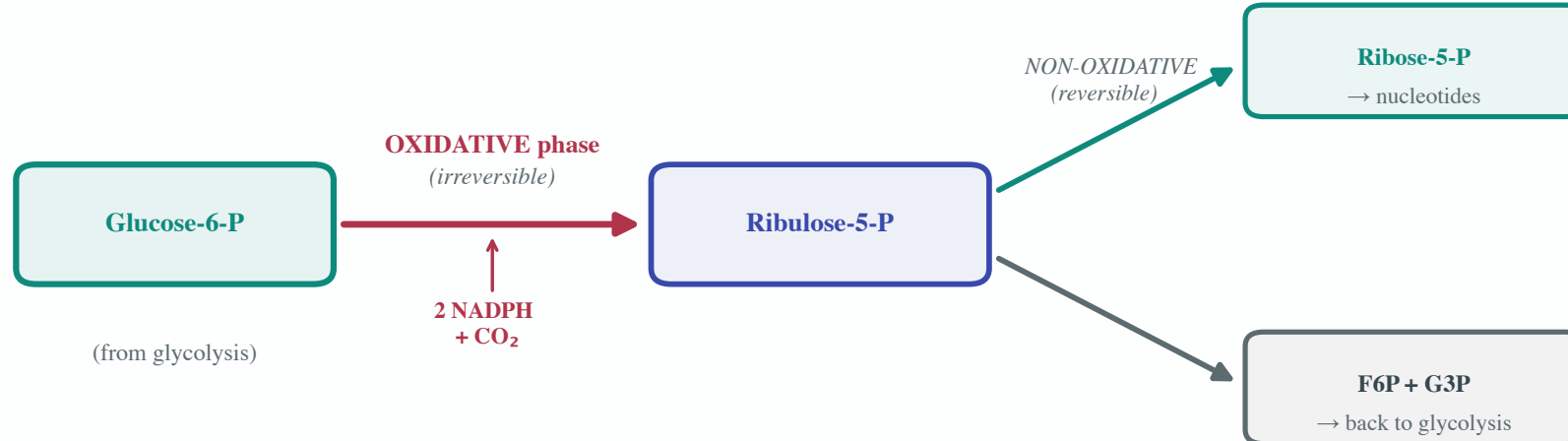
## 3 · The Pentose Phosphate Pathway

"A branch off glycolysis that makes something ATP can't – NADPH for building and defense, plus ribose for your DNA."

# Not everything is about ATP

Glucose-6-phosphate can take a **detour** off glycolysis: the **pentose phosphate pathway**. It runs in **two phases** – an **oxidative** phase (irreversible) that makes **NADPH**, and a **non-oxidative** phase (reversible) that shuffles sugars around. The 5-carbon **ribulose-5-P** in the middle can become **ribose-5-P** or fold **back into glycolysis**.

The pentose phosphate pathway: NADPH + ribose, off the glycolysis trunk



# Two products worth having

**NADPH** looks like NADH but does a totally different job: it's **reducing power for building** – fatty acids, cholesterol – and for **antioxidant defense** (it keeps glutathione charged). **Ribose-5-phosphate** is the sugar backbone of your **nucleic acids** (**RNA** directly; **DNA** once it's reduced) and of cofactors like ATP and NAD<sup>+</sup>. Energy? That's glycolysis's job – this pathway supplies **materials**.

Not ATP — NADPH is different: it's for building and for defense

## **NADPH**

reducing power for **BUILDING**

- fatty-acid & cholesterol synthesis
- glutathione → antioxidant defense

## **Ribose-5-phosphate**

the sugar for nucleic acids

- DNA & RNA
- ATP, NAD<sup>+</sup>, CoA backbones

# It flexes to what the cell needs

The beauty of the PPP is that it's **flexible**. Need only **ribose** (a dividing cell)? Run just the non-oxidative phase. Need **NADPH** (a fat-making liver cell)? Run oxidative, then recycle the carbons back into glycolysis. Need **both**, or **NADPH plus ATP**? There's a mode for that too — **four paths**, one pathway.

Four modes — the cell dials NADPH vs ribose vs ATP to its need

## Need RIBOSE only

*(dividing cells)*

non-oxidative only — make R5P  
from glycolysis intermediates

## Need BOTH

*(balanced)*

oxidative phase alone gives  
NADPH + ribose-5-P

## Need NADPH only

*(fat synthesis)*

oxidative → recycle carbons  
back into glycolysis

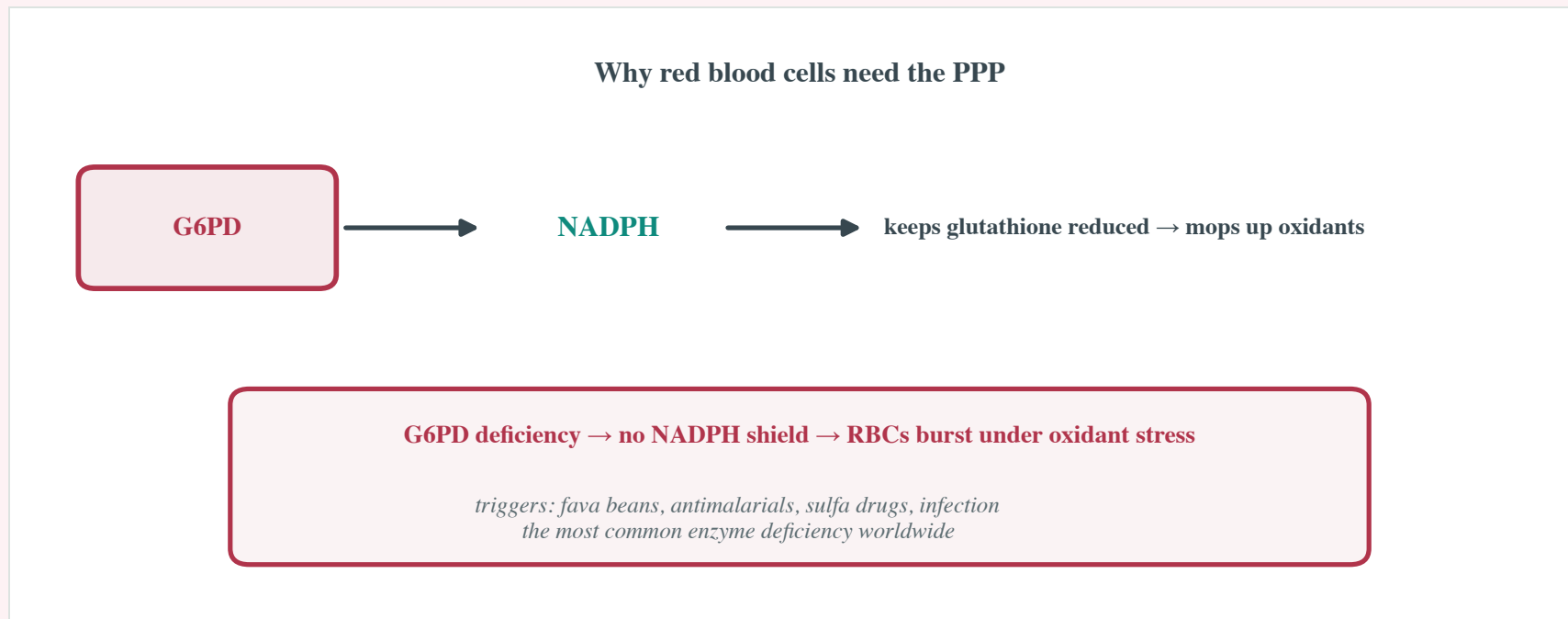
## Need NADPH + ATP

*(energy too)*

oxidative NADPH, then carbons  
→ glycolysis → pyruvate

# Why your red blood cells depend on it

**G6PD** (glucose-6-phosphate dehydrogenase) starts the oxidative phase – so it's the **NADPH supply**. Red blood cells use that NADPH to keep **glutathione** charged and neutralize oxidants. In **G6PD deficiency** – the world's **most common enzyme defect** – that shield fails, and RBCs **burst** when hit by oxidative stress: **fava beans**, antimalarials, sulfa drugs, or infection.

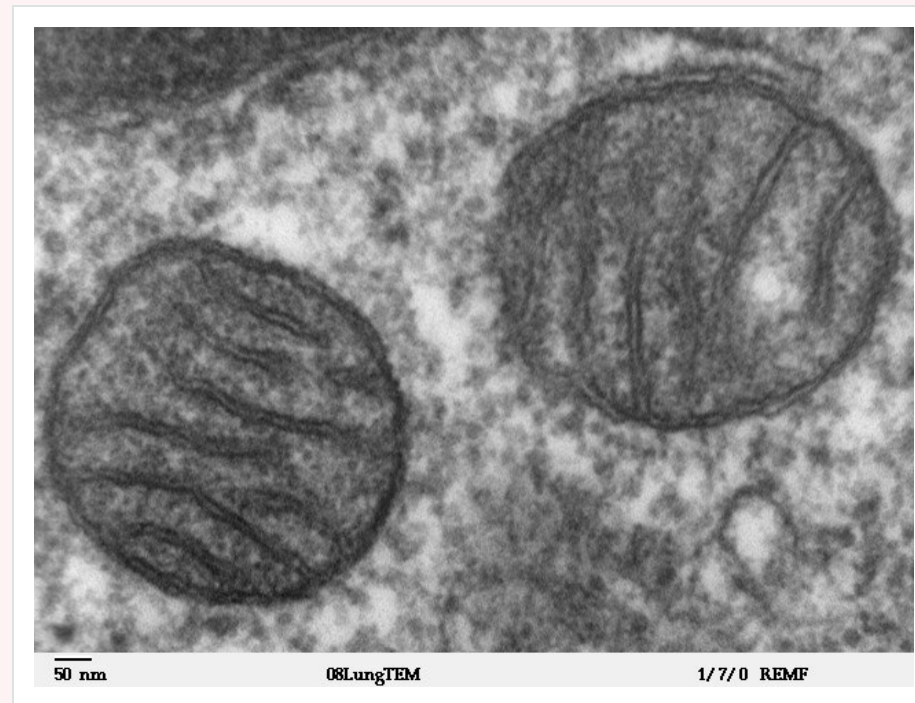


## 4 • The Electron Transport Chain

"Cash in all that NADH and  $\text{FADH}_2$  – watch electrons tumble down four complexes to oxygen, pumping protons the whole way."

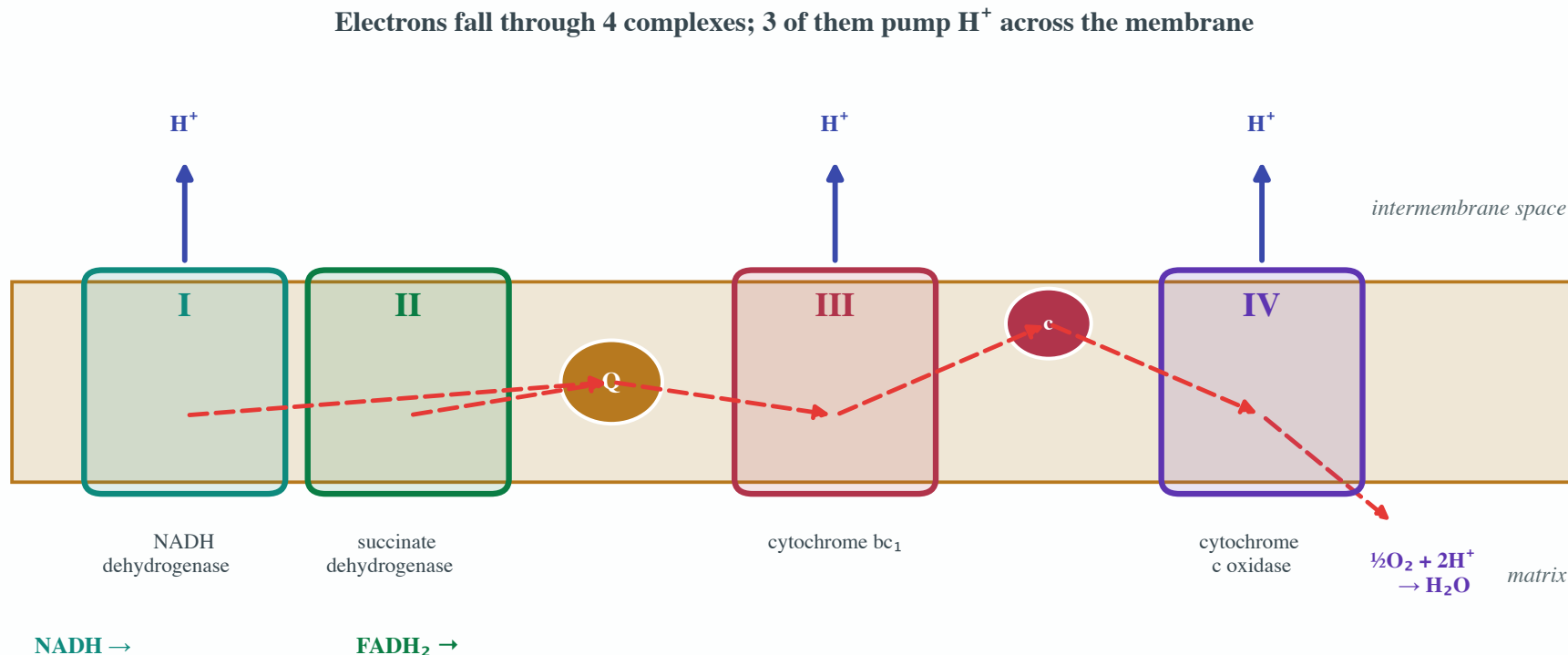
## Where the real ATP is made

Glycolysis and the TCA cycle made only a little ATP directly – their real haul was **NADH and FADH<sub>2</sub>**. Now we cash them in. It all happens on the **inner mitochondrial membrane**, folded into **cristae** to pack in as much surface area (and as many chains) as possible. You can see those folds in this real electron micrograph.



# Four complexes, one chain

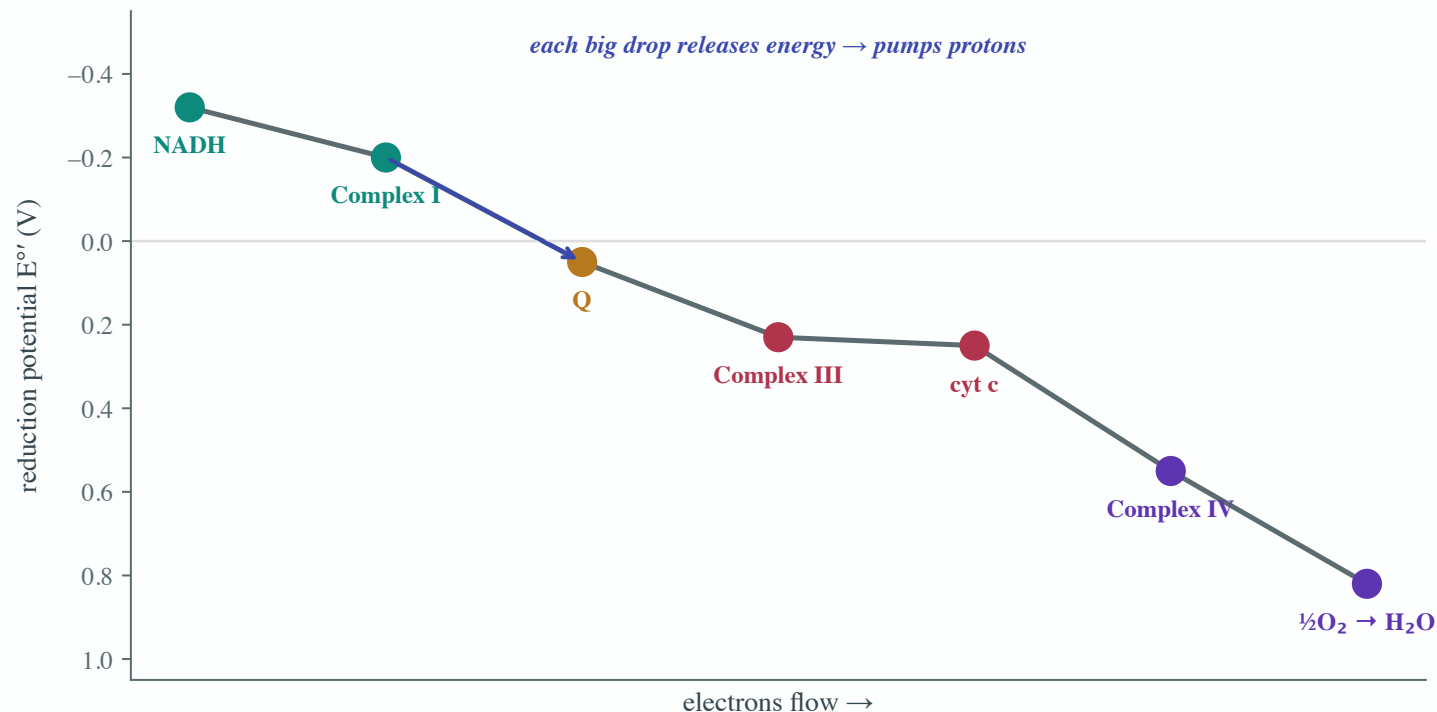
The chain is **four protein complexes** in the membrane. Electrons are handed from one to the next – NADH drops them at **Complex I**, FADH<sub>2</sub> at **Complex II** – and mobile carriers **Q** (ubiquinone) and **cytochrome c** ferry them along. Three of the complexes (**I, III, IV**) use the released energy to **pump H<sup>+</sup>** out into the intermembrane space.



*the red dashes are electrons · blue arrows are protons pumped up*

# Downhill all the way to oxygen

Why do electrons flow **this** direction? Each carrier has a **reduction potential** – a "grabbiness" for electrons. Electrons fall from **low** potential (**NADH**) to **high** (**O<sub>2</sub>**, the greediest of all), and every big drop **releases energy** to pump protons. At the end, **Complex IV** hands them to **O<sub>2</sub>**, making **water** – which is why you must **breathe**.



# NADH vs FADH<sub>2</sub>: entry matters

Here's why NADH is worth more than FADH<sub>2</sub>. **NADH** enters at **Complex I** and drives **all three** proton pumps. **FADH<sub>2</sub>** enters later, at **Complex II**, so it **skips Complex I** – only two pumps. Fewer protons pumped means less ATP: about **2.5** for NADH versus **1.5** for FADH<sub>2</sub>.

Where you enter decides how much ATP you're worth

## NADH

enters at Complex I · passes 3 pumping complexes (I, III, IV)

≈ 2.5 ATP

## FADH<sub>2</sub>

enters at Complex II · skips Complex I — only 2 pumps (III, IV)

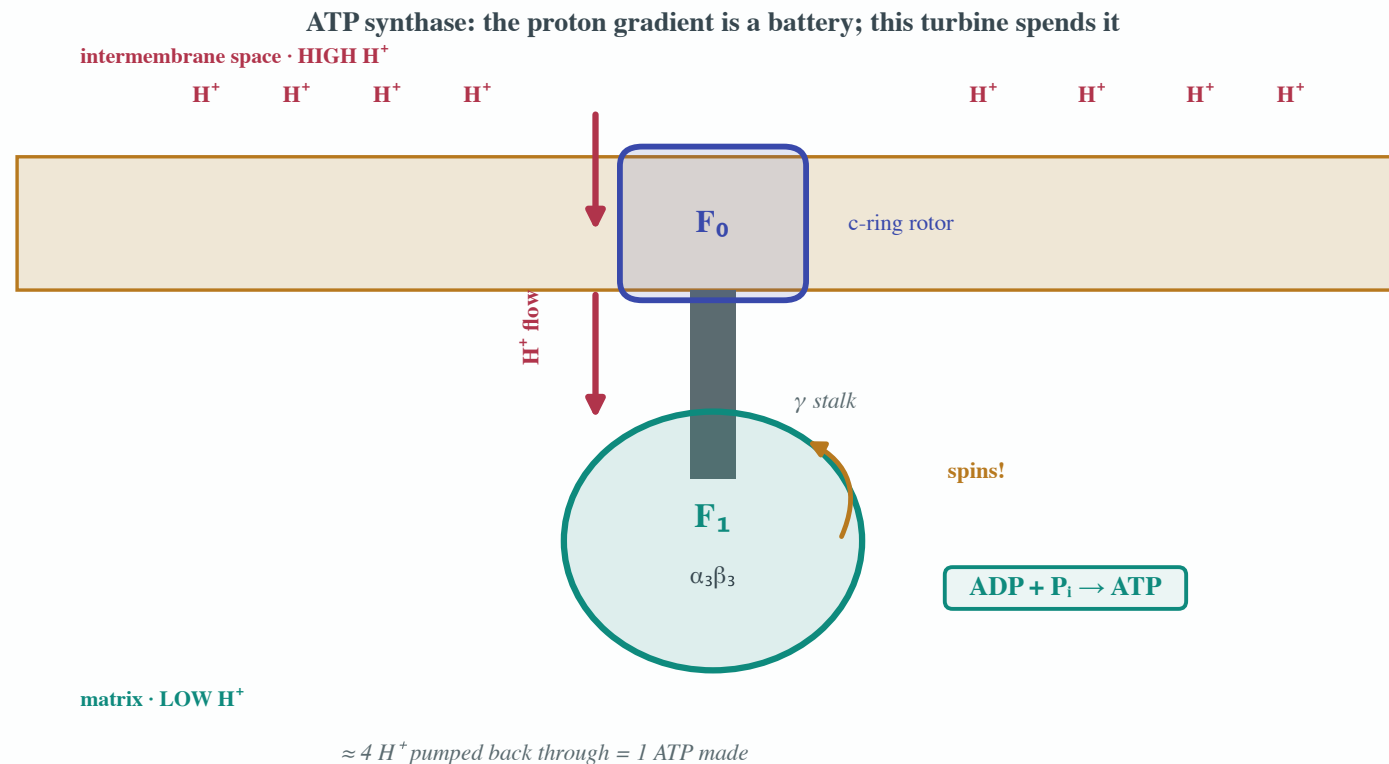
≈ 1.5 ATP

## 5 · Oxidative Phosphorylation

"The payoff – the proton gradient spins a molecular turbine into ATP. Add it all up: one glucose, ~30 ATP."

# The gradient is a battery

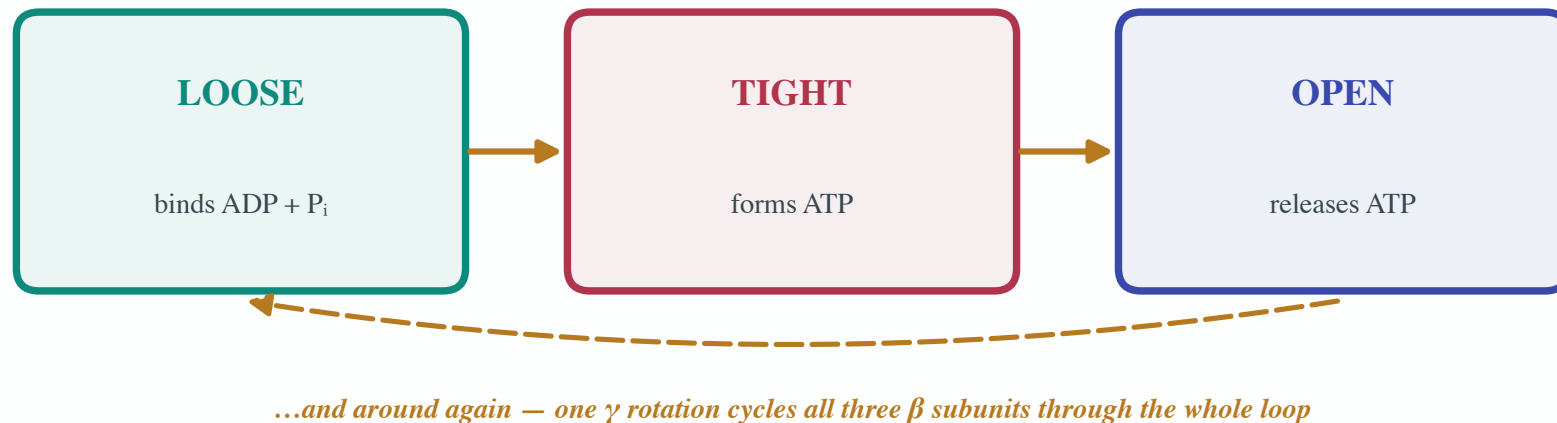
All that proton-pumping charged up a **gradient** across the inner membrane – more  $H^+$  outside than in. That's **stored energy** (the proton-motive force), and it wants to flow back. The only door back in is through **ATP synthase**, and as protons rush through, they **spin it like a turbine** – that's **chemiosmosis**, and it's where most of your ATP is made.



# How a spinning motor makes ATP

ATP synthase has two parts: **F<sub>o</sub>** (a **rotor** in the membrane that the protons turn) and **F<sub>1</sub>** (the  **$\alpha_3\beta_3$**  head in the matrix that does the chemistry), joined by a  **$\gamma$  stalk**. As  $\gamma$  spins, it shoves each  **$\beta$  subunit** through three shapes – **Loose** (grab ADP + P<sub>i</sub>), **Tight** (squeeze them into ATP), **Open** (let ATP go). Mechanical motion → chemical bond.

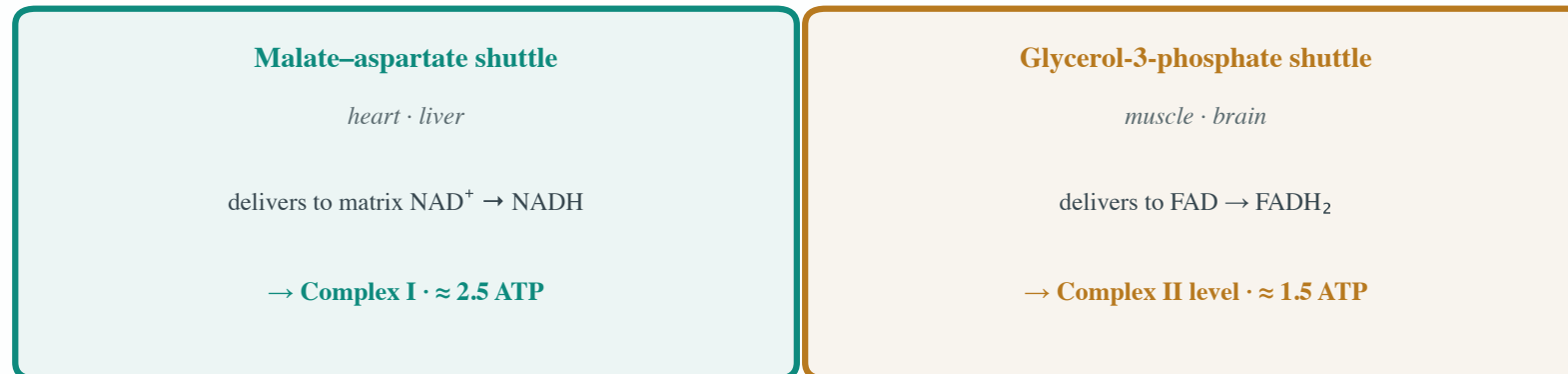
The binding-change mechanism — mechanical rotation, chemical ATP



# Getting cytosolic NADH into the game

One problem: the **NADH made in glycolysis** is stuck in the **cytosol** – the inner membrane won't let it cross. So its **electrons** are ferried in by a **shuttle**. The **malate-aspartate** shuttle (heart, liver) hands them to **NADH** (worth ~2.5 ATP); the **glycerol-3-phosphate** shuttle (muscle, brain) hands them to **FADH<sub>2</sub>** (only ~1.5 ATP).

Cytosolic NADH gets in by shuttle — and the shuttle sets its ATP value



*That's why "ATP per glucose" is a range, not a fixed number.*

# Adding it all up

Here's the whole harvest from **one glucose**. A little **direct** ATP (2 from glycolysis, 2 GTP from the TCA cycle) – but the bulk comes from cashing in **10 NADH + 2 FADH<sub>2</sub>** at the electron transport chain. All told: **about 30–32 ATP**, depending on which shuttle carried the cytosolic NADH. **That's** why we breathe.

| stage                      | Adding it all up: one glucose, fully oxidized |                                     |
|----------------------------|-----------------------------------------------|-------------------------------------|
|                            | direct                                        | carriers → ETC                      |
| Glycolysis                 | <b>2 ATP</b>                                  | <b>2 NADH</b>                       |
| Pyruvate → acetyl-CoA (×2) | —                                             | <b>2 NADH</b>                       |
| TCA cycle (×2 turns)       | <b>2 GTP</b>                                  | <b>6 NADH + 2 FADH<sub>2</sub></b>  |
| <b>totals</b>              | <b>4 ATP</b>                                  | <b>10 NADH + 2 FADH<sub>2</sub></b> |

**GRAND TOTAL ≈ 30–32 ATP per glucose**

4 direct + (10 NADH × 2.5) + (2 FADH<sub>2</sub> × 1.5) = 32  
slightly less if the glycerol-P shuttle is used

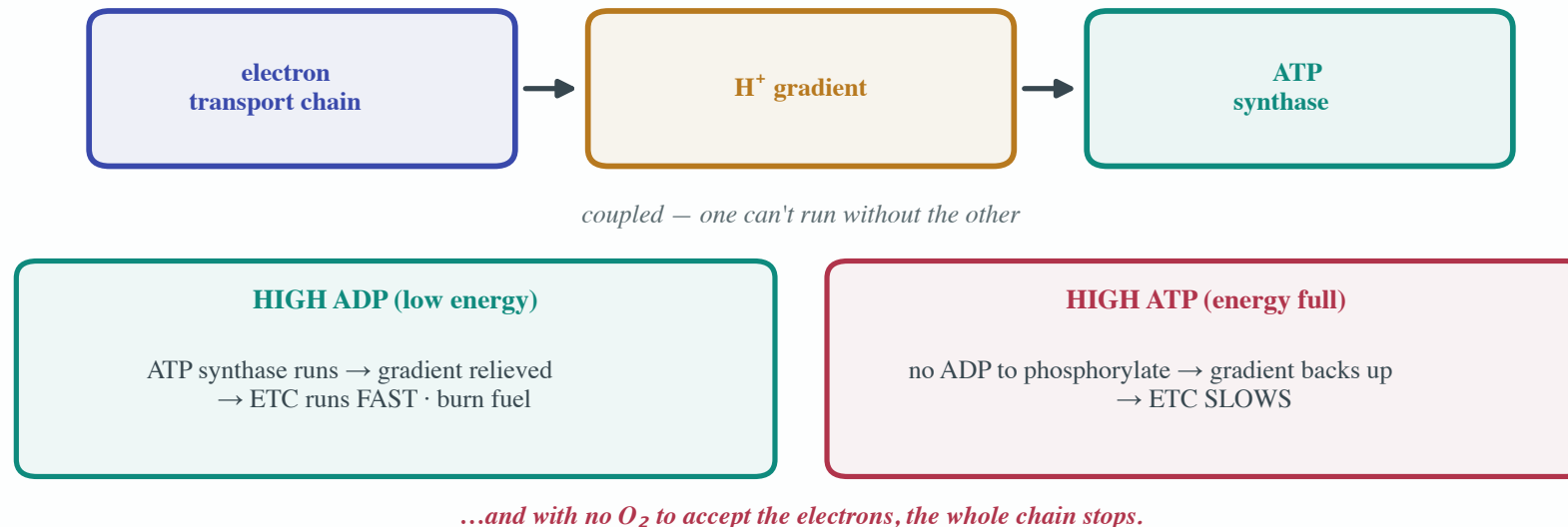
## 6 · Regulation, Poisons & Uncouplers

"What sets the pace of respiration – and what happens when you poison it or short-circuit it for heat."

# What sets the pace: ADP

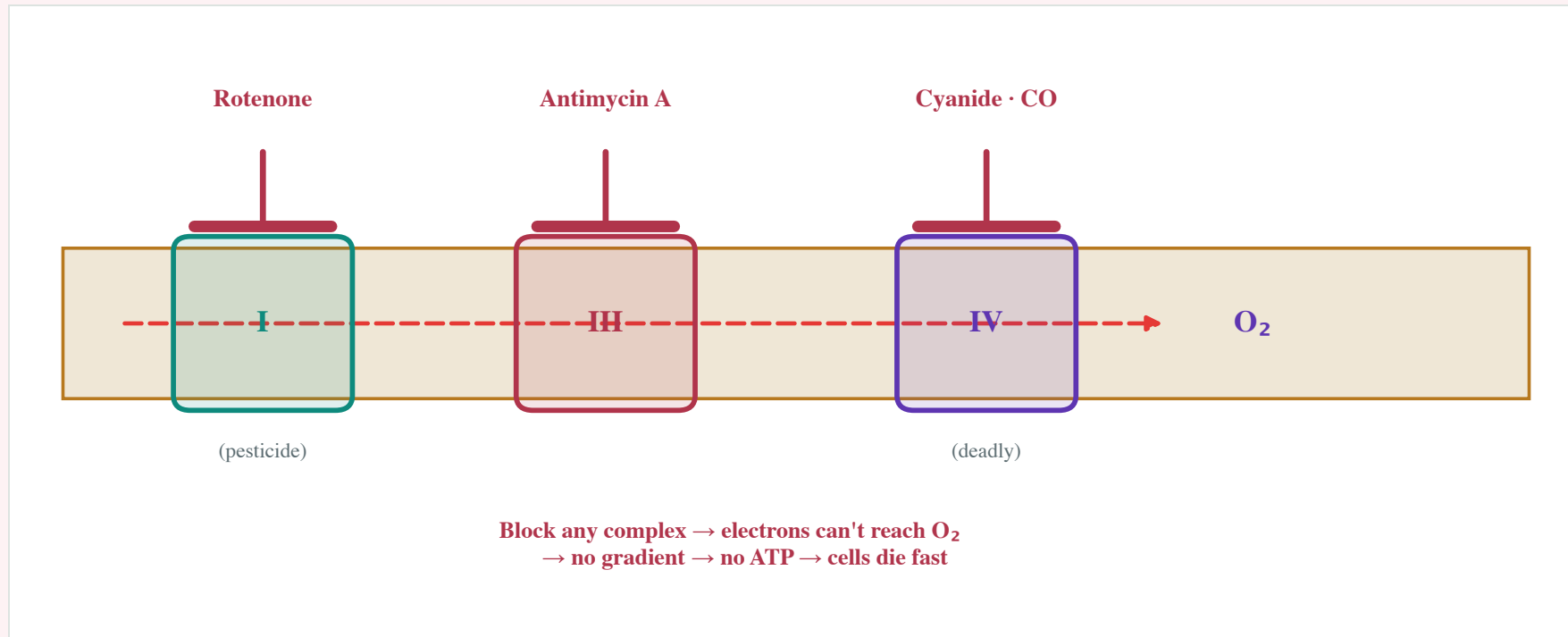
The electron transport chain and ATP synthase are **coupled** – chained together by the proton gradient, so **one can't run without the other**. That makes **ADP the throttle** (called **respiratory control**): when **ADP is high** (you're low on energy), ATP synthase runs, the gradient is relieved, and the chain burns **fast**. When **ATP is high**, everything **slows**. And with **no O<sub>2</sub>** to catch the electrons at the end, the whole chain simply **stops**.

Respiratory control — the whole system runs only when ATP is needed



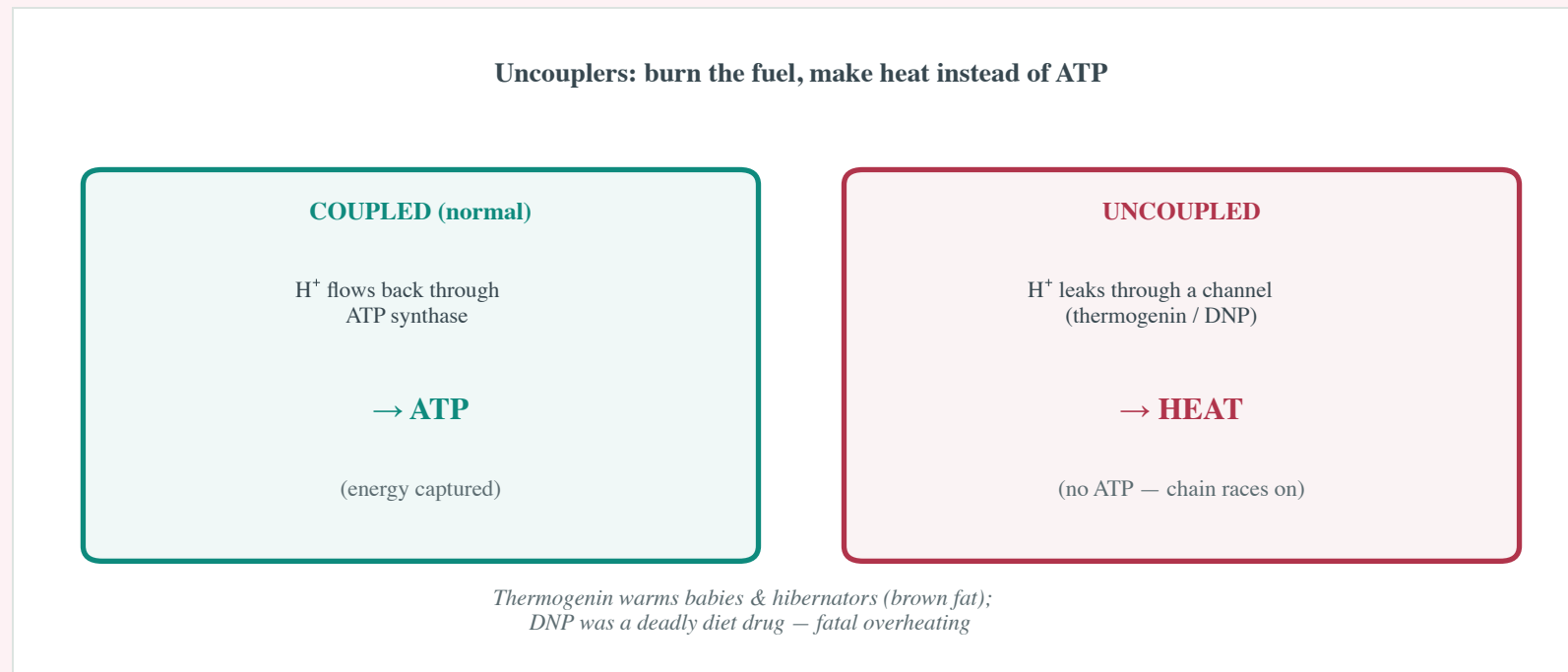
# Poisons that stop the chain

Because every complex passes electrons to the next, **blocking any one** jams the whole line. **Rotenone** (a pesticide) blocks **Complex I**; **antimycin A** blocks **Complex III**; **cyanide** and **carbon monoxide** block **Complex IV**. Electrons can't reach **O<sub>2</sub>**, the gradient collapses, ATP synthesis halts – and cells die **fast**. This is exactly how cyanide kills.



# Uncouplers: heat instead of ATP

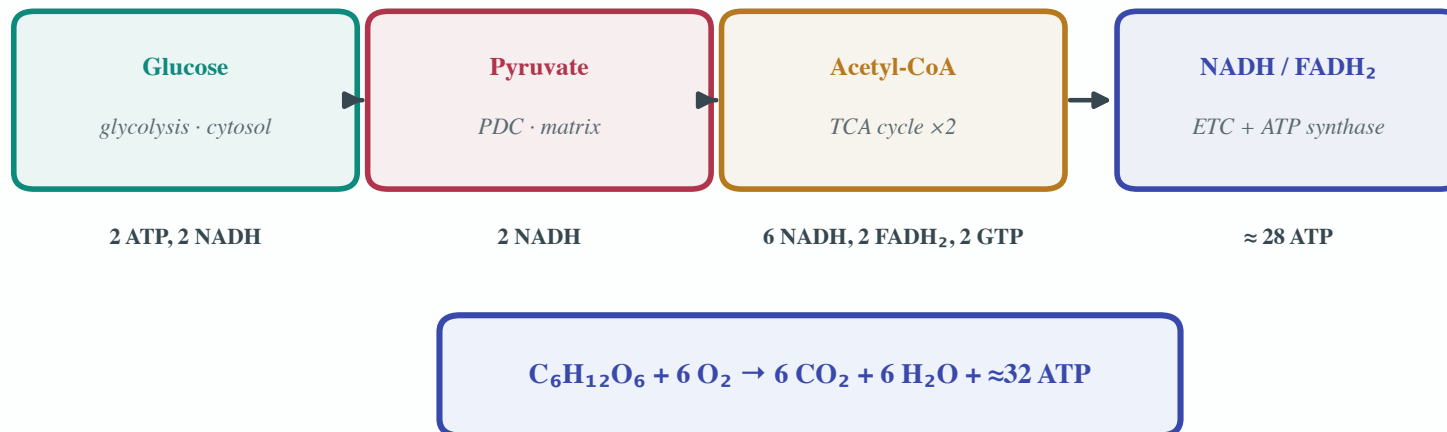
There's another way to wreck it: an **uncoupler** lets **H<sup>+</sup> leak back across** the membrane **without** passing through ATP synthase – so the energy comes out as **heat**, not ATP. Your **brown fat** does this **on purpose** with **thermogenin** to keep babies warm; the diet drug **DNP** did it too – and cooked people to death.



# The whole story, one glucose

Step back and look at it all: **glucose** is split in the **cytosol**, its pyruvate shipped into the **mitochondrion**, oxidized by the **TCA cycle**, and the harvested **NADH and FADH<sub>2</sub>** cashed in at the **electron transport chain** to spin out ATP. One glucose, fully burned to **CO<sub>2</sub> and water**, yields about **32 ATP**. **That's** why you breathe.

The whole story: one glucose → carbon dioxide, water, and ~32 ATP



# Can you...?

- ☐ explain the pyruvate dehydrogenase complex – the link reaction to acetyl-CoA – its cofactors and its allosteric + covalent regulation?
- ☐ walk all eight reactions of the TCA cycle with the running NADH/FADH<sub>2</sub>/GTP ledger, and explain its regulated, amphibolic, and anaplerotic roles?
- ☐ describe the pentose phosphate pathway – its oxidative and non-oxidative phases and its NADPH and ribose-5-phosphate products?
- ☐ trace electron flow through the four ETC complexes, driven by reduction potential, pumping protons to build the gradient?
- ☐ explain chemiosmosis and ATP synthase's rotary catalysis, the NADH shuttles, and the total ATP yield from one glucose?
- ☐ describe the regulation of respiration, uncouplers (thermogenin), and ETC poisons (cyanide, rotenone)?

If any box stays empty, the practice site has a drill for it.

