

KEY

I. Short Answer

A. Lactate produced in the muscle under anaerobic conditions is transported through the blood to the liver, where it reacts with lactate dehydrogenase and proceeds through gluconeogenesis; the resulting glucose is released back into the bloodstream.

What is the name of this cycle?

Cori cycle

B. Even though gluconeogenesis in the liver should be favorable ($\Delta G^\circ < 0$), the $\Delta G'$ of the reaction is even more negative after a period of intense exercise when the levels of lactate produced in the body may be quite high. Why is this so?

$$\Delta G' = \Delta G^\circ + RT \ln (\text{products/reactants})$$

Since lactate conc. is high and this is the reactant for the pathway, the $\Delta G'$ is more negative than the ΔG° (AKA "substrate loading")

C. Why does the glucose produced from gluconeogenesis in the liver not "turn around" and get directed right back into the glycolytic pathway?

The K_m for glucose by glucokinase is high; that is, it is not a good substrate for glycolysis.

D. Under anaerobic conditions, yeast will grow to a high density only if there is a very high concentration of sugar, whereas under aerobic conditions, they can achieve the same density with much less sugar. What is the (i) name and (ii) physiological basis of this effect?

(i) Pasteur Effect.

(ii) 38 vs 2 ATPs can be produced per glucose under aerobic vs anaerobic conditions, respectively. Thus, the yeast "need" less food.

E. During glycolysis, fructose-1,6-bisphosphate activates another enzyme in the glycolytic pathway. What is the (i) name of this enzyme and (ii) what is the name given to this type of activation?

(i) Pyruvate Kinase

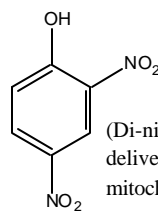
(ii) Feed-forward activation

F. Enzymes such as hexokinase and citrate synthase undergo a large conformational change to exclude water from their active site. What is the term used to describe this conformational change?

Induced-Fit

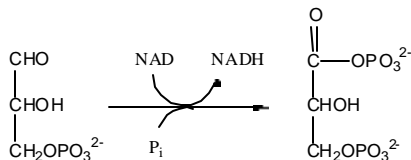
G. The structure of the following molecule belongs to a class of factors known as what?

Uncouplers.



(Di-nitrophenol - can deliver H⁺s across the mitoch. inner membrane)

II. Draw the structures of the reactants and products of the reaction catalyzed by glyceraldehyde-3-phosphate dehydrogenase.

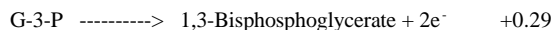


Using the following table, calculate the E'° for this reaction.

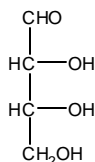
Oxidant	Reductant	n	E'° , V
Acetate + CO ₂ + 2H ⁺	Pyruvate + H ₂ O	2	-0.70
Succinate + CO ₂ + 2H ⁺	α-Ketoglutarate + H ₂ O	2	-0.67
Acetate + 3H ⁺	Acetaldehyde + H ₂ O	2	-0.60
O ₂	O ₂ ⁻	1	-0.45
Ferredoxin (oxidized)	Ferredoxin (reduced)	1	-0.43
2H ⁺	H ₂	2	-0.42
Acetoacetate + 2H ⁺	β-Hydroxybutyrate	2	-0.35
Pyruvate + CO ₂ + H ⁺	Malate	2	-0.33
NAD ⁺ + H ⁺	NADH	2	-0.32
NADP ⁺ + H ⁺	NADPH	2	-0.32
FMN (enzyme-bound) + 2H ⁺	FMNH ₂ (enzyme-bound)	2	-0.30
Lipoate (oxidized) + 2H ⁺	Lipoate (reduced)	2	-0.29
1,3-Bisphosphoglycerate + 2H ⁺	Glyceraldehyde-3-phosphate + P _i	2	-0.29
Glutathione (oxidized) + 2H ⁺	2 Glutathione (reduced)	2	-0.23
FAD + 2H ⁺	FADH ₂	2	-0.22
Acetaldehyde + 2H ⁺	Ethanol	2	-0.20
Pyruvate + 2H ⁺	Lactate	2	-0.19
Oxaloacetate + 2H ⁺	Malate	2	-0.17
α-Ketoglutarate + N ₂ + 2H ⁺	Glutamate + H ₂ O	2	-0.14

Methylene blue (oxidized) + 2H ⁺	Methylene blue (reduced)	2	0.01
Fumarate + 2H ⁺	Succinate	2	0.03
CoQ + 2H ⁺	CoQH ₂	2	0.04
Cytochrome b (+3)	Cytochrome b (+2)	1	0.07
Dehydroascorbate + 2H ⁺	Ascorbate	2	0.08
Cytochrome c ₁ (+3)	Cytochrome c ₁ (+2)	1	0.23
Cytochrome c (+3)	Cytochrome c (+2)	1	0.25
Cytochrome a (+3)	Cytochrome a (+2)	1	0.29
½O ₂ + H ₂ O	H ₂ O ₂	2	0.30
Ferricyanide	Ferrocyanide	2	0.36
Nitrate + 2H ⁺	Nitrite + H ₂ O	1	0.42
Cytochrome a ₃ (+3)	Cytochrome a ₃ (+2)	1	0.55
Fe (+3)	Fe (+2)	1	0.77
½O ₂ + 2H ⁺	H ₂ O	2	0.82

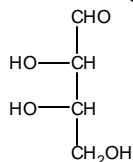
Note: E₀ is the standard reduction potential at pH 7 and 25°C, n is the number of electrons transferred, and each potential is for the partial reaction written as follows: Oxidant + ne⁻ → reductant.



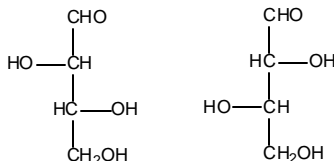
III. Draw the (A) enantiomeric and (B) diastereomeric forms of the following sugar:



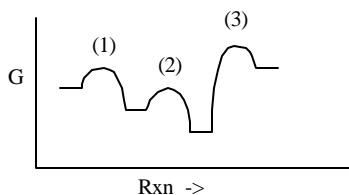
A. "mirror image"



B.



IV. The following depicts an energy diagram of a reaction catalyzed by an enzyme that we have discussed:



A. Which step is rate limiting? 3

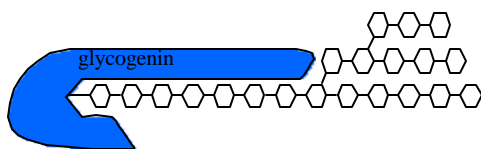
B. Which enzyme that we discussed does this diagram most closely resemble?

F₁F₀ ATP Synthase

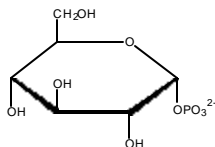
C. Ignoring the 3rd step, if reactions 1 and 2 describe a short biosynthetic pathway (A + B making product C), how might you drive the process in reverse? That is, what "tricks" are used in living systems to make this happen? Give 2 examples.

Hormonal control Substrate Loading Utilization of different compartments Utilization of by-pass rxns
Channelling Changes in enzyme conc. Cleave good leaving group...

V. The following is a representation of a short glycogen polymer:



Draw the structure of the product when glycogen phosphorylase acts on this polymer:



How many of these products can be formed before the action of de-branching enzyme?

Needs to stop 4 from branch. So, 1. (From bottom chain)

Assuming that de-branching enzyme is not active, what is the theoretical maximum number of ATPs that can be generated under aerobic conditions from the action of phosphorylase on this polymer?

1 glucose-1-p <----> 1 glucose-6-p
 | Glycolysis 3 ATPs each
 | NADH---> 2x3 = 6 ATPs each
 | TCA ---> 12 x 2 = 24 ATPs each
 | Pyr.d.h. 2NADH x 3 = 6 ATPs each

VI. What are the names of the following co-factors?

