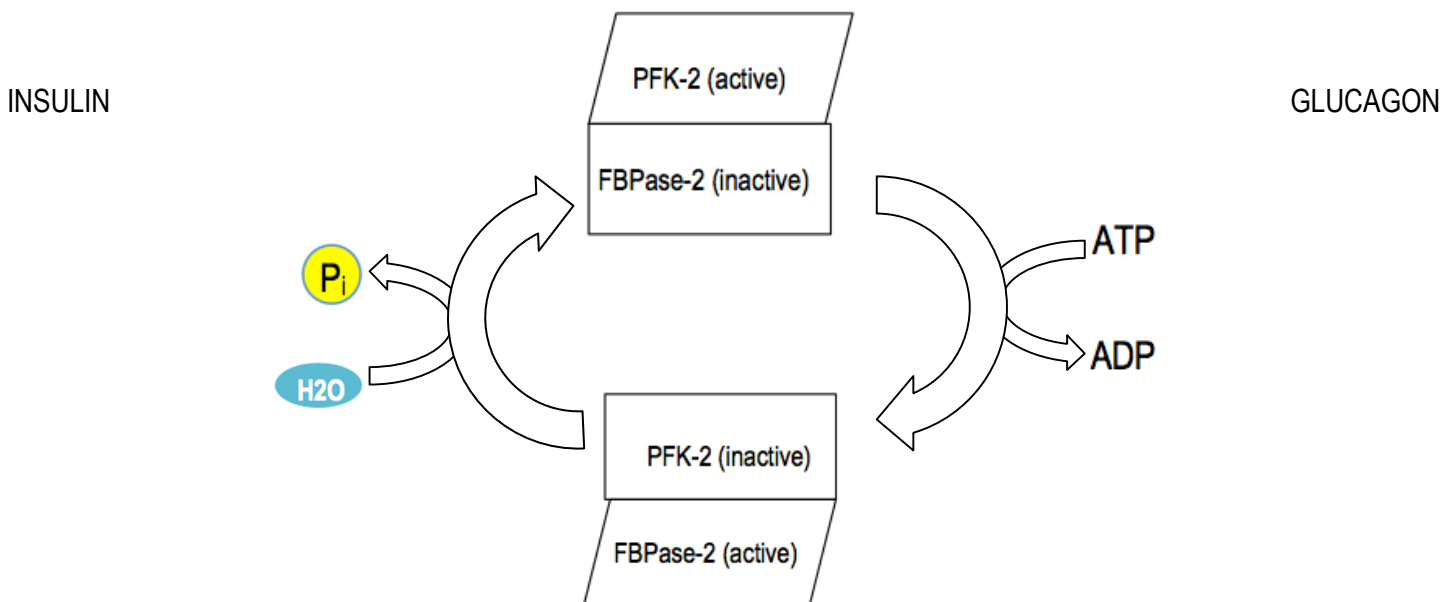


CONCEPT: METABOLIC REGULATION

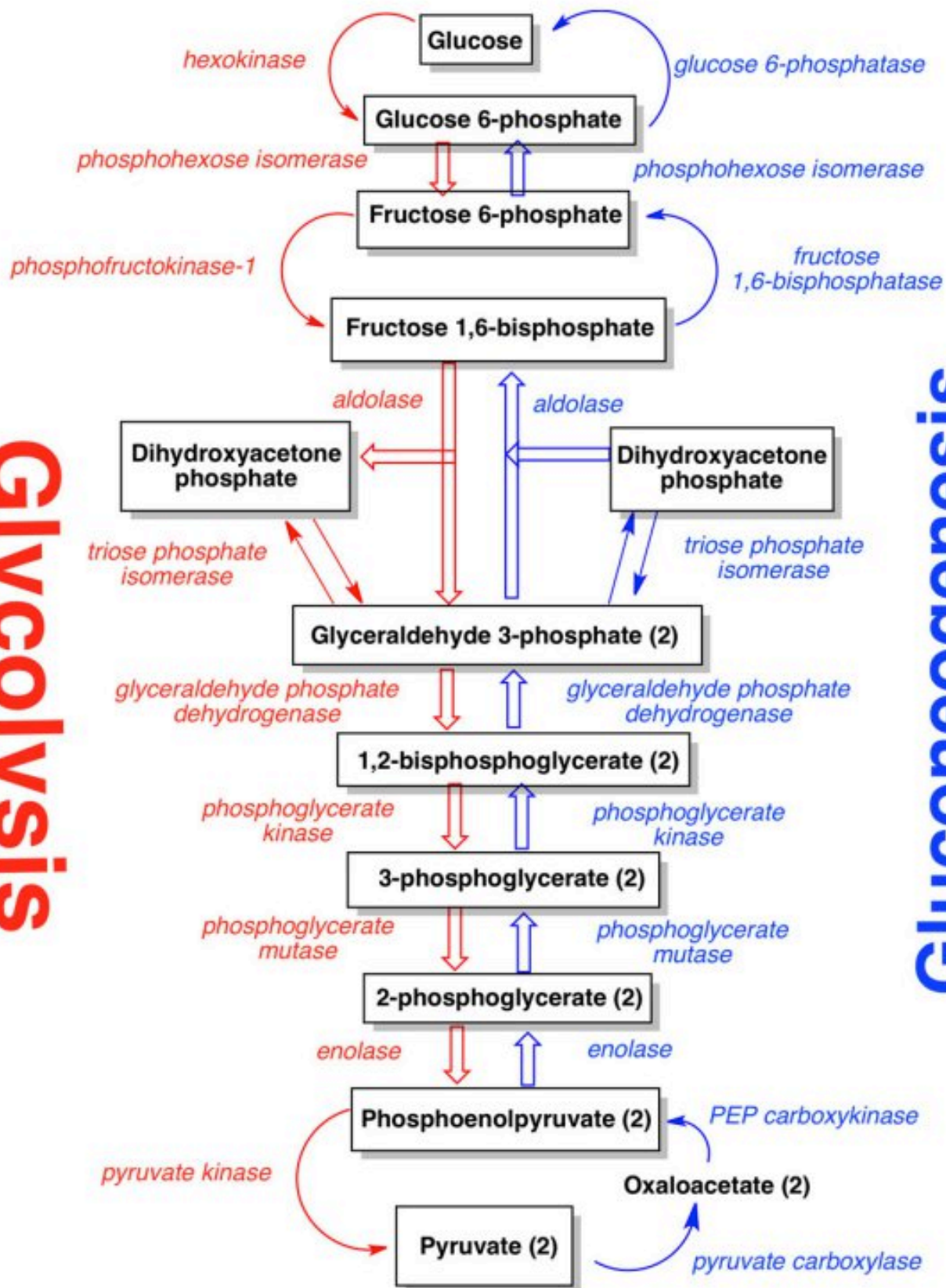
- Phosphofructokinase-1 (PFK-1) – catalyzes the commitment step of glycolysis
 - Has allosteric regulatory sites that bind ATP; high [ATP] lowers enzyme activity
 - Citrate is transported out into cytoplasm for biosynthesis, and high concentrations will inhibit PFK-1
 - ADP and AMP stimulate the enzyme's activity
 - Fructose 2,6-bisphosphate is an allosteric activator, and the most important regulator of PFK-1
 - PFK-1 is considerably less active without fructose 2,6-bisphosphate
 - Fructose 2,6-bisphosphate is effective at extremely small concentrations (~1 μ M)
 - Fructose 2,6-bisphosphate inhibits fructose 1,6-bisphosphatase, PFK-1's gluconeogenic counterpart
 - AMP also inhibits fructose 1,6-bisphosphatase, and therefore inhibits gluconeogenesis
- Phosphofructokinase-2 (PFK-2) – generates fructose 2,6-bisphosphate to regulate the action of PFK-1
 - Insulin causes phosphate group to be removed from enzyme, activates PFK2
 - Glucagon leads to phosphorylation of enzyme, activating fructose bisphosphatase 2
 - Fructose 2,6-bisphosphate is broken down by fructose bisphosphatase 2; part of same protein as PFK-2
 - Phosphoprotein phosphatase dephosphorylates PFK-2; stimulated by xylulose 5-phosphate



- Pyruvate kinase – last enzyme of glycolysis; only in the liver does glucagon cause PKA to phosphorylate and inactivate it
 - Stimulated by fructose 1,6-bisphosphate, an upstream glycolytic substrate
 - Inhibited by ATP, acetyl-CoA, and long-chain fatty acids (also alanine, which is derived from pyruvate)
- Pyruvate carboxylase is activated by acetyl-CoA

CONCEPT: METABOLIC REGULATION

Glycolysis



Gluconeogenesis