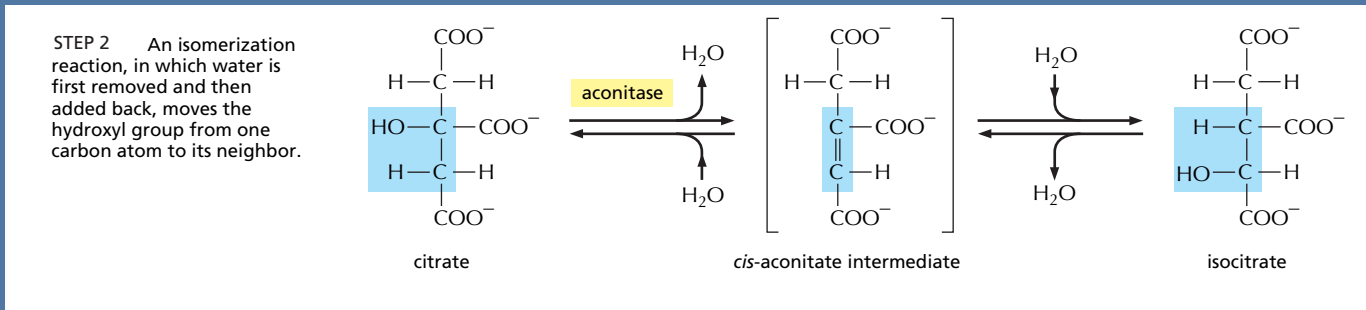
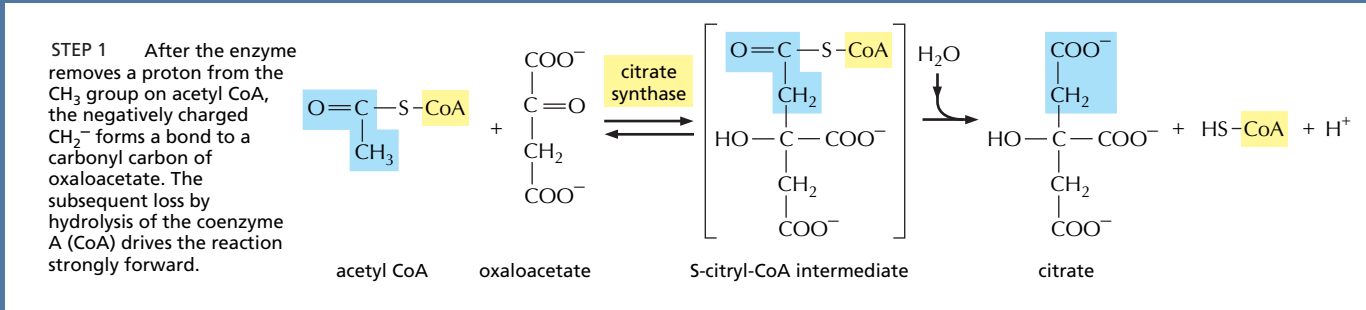
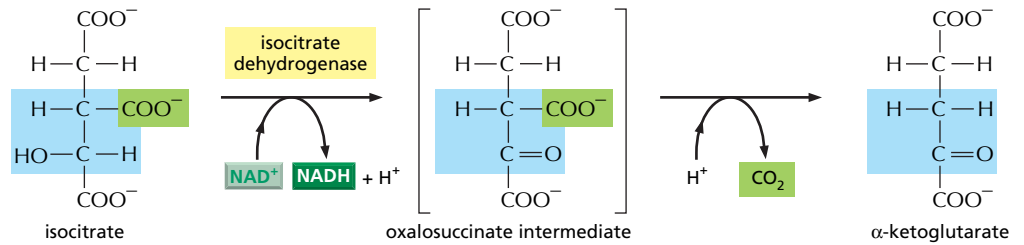


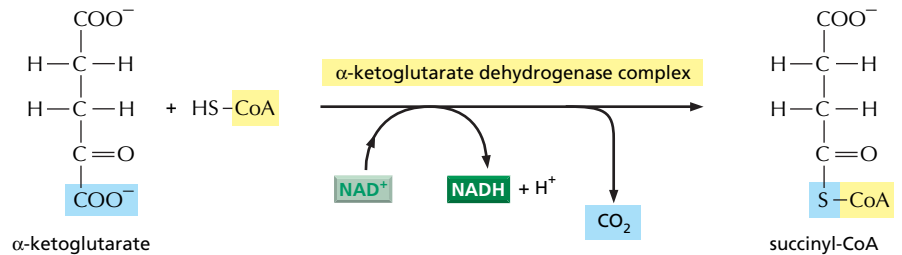
Details of the eight steps are shown below. For each step, the part of the molecule that undergoes a change is shaded in blue, and the name of the enzyme that catalyzes the reaction is in a yellow box.



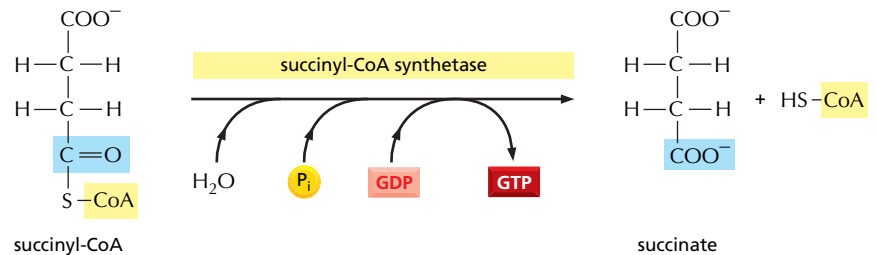
**STEP 3** In the first of four oxidation steps in the cycle, the carbon carrying the hydroxyl group is converted to a carbonyl group. The immediate product is unstable, losing  $\text{CO}_2$  while still bound to the enzyme.



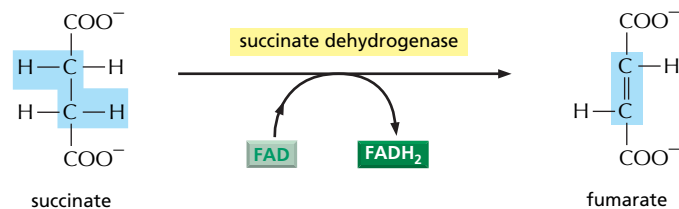
**STEP 4** The  $\alpha$ -ketoglutarate dehydrogenase complex closely resembles the large enzyme complex that converts pyruvate to acetyl CoA (pyruvate dehydrogenase). It likewise catalyzes an oxidation that produces NADH,  $\text{CO}_2$ , and a high-energy thioester bond to coenzyme A (CoA).



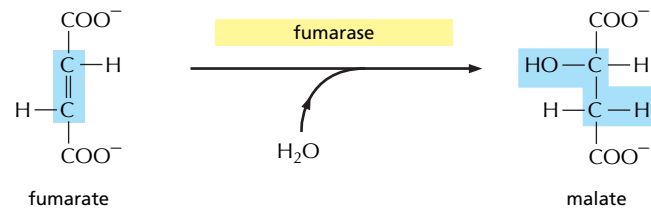
**STEP 5** A phosphate molecule from solution displaces the CoA, forming a high-energy phosphate linkage to succinate. This phosphate is then passed to GDP to form GTP. (In bacteria and plants, ATP is formed instead.)



**STEP 6** In the third oxidation step in the cycle, FAD removes two hydrogen atoms from succinate.



**STEP 7** The addition of water to fumarate places a hydroxyl group next to a carbonyl carbon.



**STEP 8** In the last of four oxidation steps in the cycle, the carbon carrying the hydroxyl group is converted to a carbonyl group, regenerating the oxaloacetate needed for step 1.

