

Phosphate mediates electron transfer in pyruvate oxidase from *Lactobacillus plantarum*

Kai Tittmann, Ralph Golbik, Gerhard Hübner

Department of Biochemistry, Martin-Luther-University Halle- Wittenberg, Germany

Sandro Ghisla

Department of Biology, University of Konstanz, Germany

Introduction

Pyruvate oxidase from *Lactobacillus plantarum* (LpPOX, EC 1.2.3.3) is homotetrameric flavoenzyme with a subunit molecular mass of 65.5 kDa composed 603 residues. Each subunit contains one tightly and noncovalently bound FA thiamin diphosphate (ThDP), the biologically active form of vitamin B₁, and Mg²⁺ anchoring the diphosphate moiety of ThDP. In the presence of phosphate and oxygen LpPOX catalyses the oxidative decarboxylation of pyruvate yielding carbon dioxide, acetylphosphate and hydrogen peroxide.

The catalytic cycle of LpPOX comprises i) the deprotonation at the C2 of ThDP, the covalent binding of pyruvate to ThDP, iii) the decarboxylation of the then formed 2-lactyl-ThDP (LThDP) intermediate to yield the carbanion/enamine of hydroxyethyl-ThDP (HEThDP), iv) an intramolecular two step electron transfer from HEThDP to FAD and finally v) the reoxidation of FADH₂ by oxygen and vi) phosphorolysis/hydrolysis of the 2-acetyl-ThDP (AcThDP) intermediate (1).

A kinetic analysis of single steps of catalysis using FAD absorbance revealed phosphorolysis of the AcThDP intermediate as well as the reoxidation of the reduced FAD to be partially rate-limiting. In the presence of phosphate no transient radical FAD species can be observed in the course of the reductive half-reaction (2).

Here, we present stopped flow and titrimetric results of the influence of the substrate phosphate on the reductive and oxidative half-reaction as well as the stabilisation of radical FAD species in the enzyme.

Results and discussion

Reductive half-reaction in the presence and absence of phosphate

The time course of the anaerobic reduction of enzyme-bound FAD in LpPOX by substrate pyruvate is clearly dependent on the presence of phosphate (Figure 1). In the presence, the progress curve shows a pronounced lag phase followed by a monophasic rapid decrease in absorbance at 457 nm. With phosphate absent, however, the

course of the reductive half reaction is composed of a *lag* phase and *two* distinguished consecutive phases.

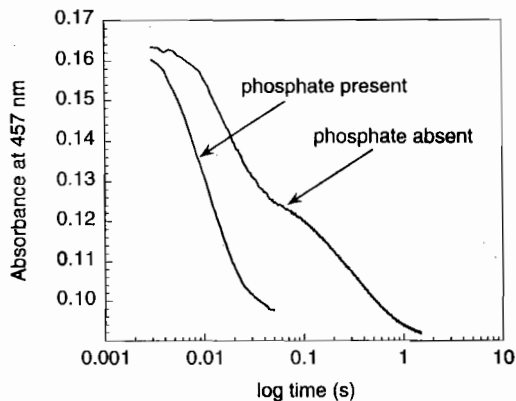


Figure 1: Time course of the *LpPOX* reaction with pyruvate at saturating concentration under anaerobic conditions in the presence and absence of phosphate at 25 °C.

A kinetic analysis of single steps in catalysis (2) revealed the reversible covalent binding of pyruvate to ThDP (k_{on} , k_{off}) as well as the decarboxylation of LThDP (k_{dec}) to be nearly independent of phosphate, whereas the intramolecular redox reaction of HThDP and FAD is remarkably slower and composed of two phases (k_{red1} , k_{red2}) with no phosphate (Table 1).

Table 1: Rate Constants of Elementary Steps of *LpPOX* Catalysis in Dependence on the Presence of Phosphate.

	k_{on} ($\text{M}^{-1} \text{s}^{-1}$)	k_{off} (s^{-1})	k_{dec} (s^{-1})	k_{red1} (s^{-1})	k_{red2} (s^{-1})
<i>LpPOX</i> in 0.1 M PIPES, pH 6.0 at 25 °C	$(2.1 \pm 0.2) \cdot 10^5$	14 ± 6	82 ± 5	138 ± 12	3.5 ± 0.1
<i>LpPOX</i> in 0.2 M KPP,	$(6.5 \pm 0.2) \cdot 10^4$	20 ± 3	112 ± 20		k_{red}

In order to prove, whether the two step reduction of FAD by HETHDP is associated with the transient occurrence of radical FAD species, the time-resolved absorbance spectra of enzyme-bound FAD in the course of the reductive half-reaction were recorded. As depicted in figure 2A, the initial rapid phase of the reaction is accompanied by an increase at $\lambda > 500$ nm and a blue-shift of λ_{max} from 405 to 378 nm. The spectra may be interpreted as being composed of the oxidized FAD and the half-reduced FAD (semiquinone). In the second phase of the reaction (figure 2B), the reduced FAD is formed.

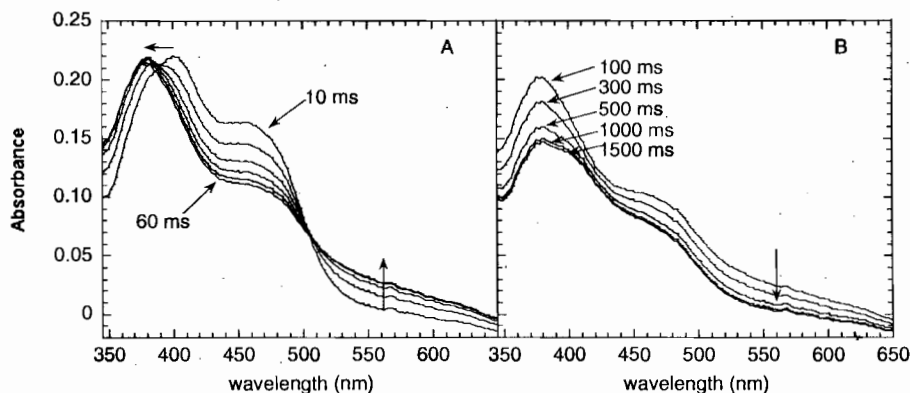


Figure 2: Time-resolved absorbance spectra of enzyme-bound FAD in *LpPOX* in the course of the reaction with pyruvate under anaerobic conditions in 0.1 M PIPES, pH 6.0 at 25 °C. A: 0-60 ms. B: 100-1500 ms.

Oxidative half-reaction in the absence and presence of phosphate

The reoxidation of the reduced FAD in *LpPOX* by oxygen (data not shown) is nearly independent of the presence of phosphate ($k_{\text{obs}} = 70 \text{ s}^{-1}$ with phosphate, $k_{\text{obs}} = 50 \text{ s}^{-1}$ without phosphate) thus supporting our suggested specific role of phosphate in the intramolecular electron transfer.

Anaerobic titration of enzyme-bound FAD with dithionite

A titration of the FAD in *LpPOX* with sodium dithionite showed (Figure 3), that the FAD semiquinone is formed both in the presence and the absence of phosphate with a comparable stabilisation of about 55 %.

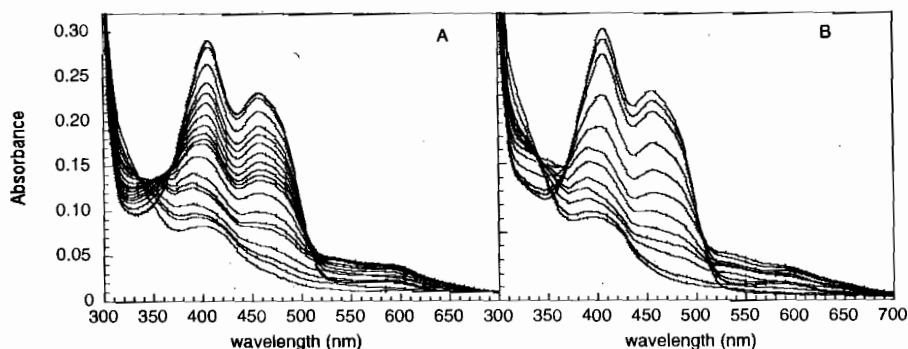


Figure 3: Anaerobic reduction of enzyme-bound FAD in *LpPOX* by sodium dithionite in either 0.2 M KPP, pH 6.0 (A) or 0.1 M PIPES, pH 6.0 (B).

Conclusions

The electron transfer from the HETHDP intermediate to FAD in *LpPOX* is dependent on the presence of phosphate. In its presence, reduction of enzyme-bound FAD proceeds via a two step single electron transfer with $k_{\text{obs}} = 422 \text{ s}^{-1}$, but no radical FAD species can be detected by using time-resolved spectroscopy. In the absence of phosphate, however, the reduction of enzyme-bound FAD is clearly biphasic and slowed. The time-resolved spectra provide evidence for a stepwise electron transfer from HETHDP to FAD with transient formation of radical FAD and an oxyethyl-ThDP species. Since the thermodynamic stabilisation of radical flavin species by *LpPOX* is independent of the presence of phosphate, the latter appears to mediate directly or facilitate electron transfer from HETHDP to FAD.

References

1. Tittmann K., Proske D., Spinka M. et al. (1998) Activation of thiamin diphosphate and FAD in the phosphate dependent pyruvate oxidase from *Lactobacillus plantarum*. *J. Biol. Chem.* **273**, 12929-12934
2. Tittmann K., Golbik R., Ghisla S., Hübner G. (2000) Mechanism of elementary catalytic steps of pyruvate oxidase from *Lactobacillus plantarum*. *Biochemistry* **39**, 10747-10754