

High-resolution structure and reaction cycle of Fatty Acid Photodecarboxylase: anatomy of a crime scene

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Fatty Acid Photodecarboxylase (FAP) is a recently discovered photoenzyme that catalyzes the conversion of fatty acids into alkane and CO₂ under light, with potential importance in green chemistry applications [1]. Its mechanism was still not fully understood and partly relied on a low-resolution crystal structure obtained from crystals with a twinning default [1]. Here, we present high-resolution crystal structures of FAP obtained in the dark and after light illumination at cryogenic temperatures (Figure 1). Combined with structural, computational, and spectroscopic techniques we are now able to provide a detailed reaction cycle of FAP. The reaction mechanism starts with an electron transfer from the fatty acid to a photoexcited oxidized flavin cofactor. Decarboxylation yields an alkyl radical, which is then reduced by back electron transfer and protonation rather than hydrogen atom transfer. Along with flavin reoxidation by the alkyl radical intermediate, a major fraction of the cleaved CO₂ unexpectedly transforms in 100 ns, most likely into bicarbonate. This is orders of magnitude faster than in solution, which indicates a catalytic step. FT-IR, structural and functional studies on variants centered on two conserved active site residues (R451 and C432) showed that R451 is essential for substrate stabilization and proton transfer. Altogether this study provides a detailed characterization of this unique enzyme and reveals a striking and unanticipated mechanistic complexity [2].

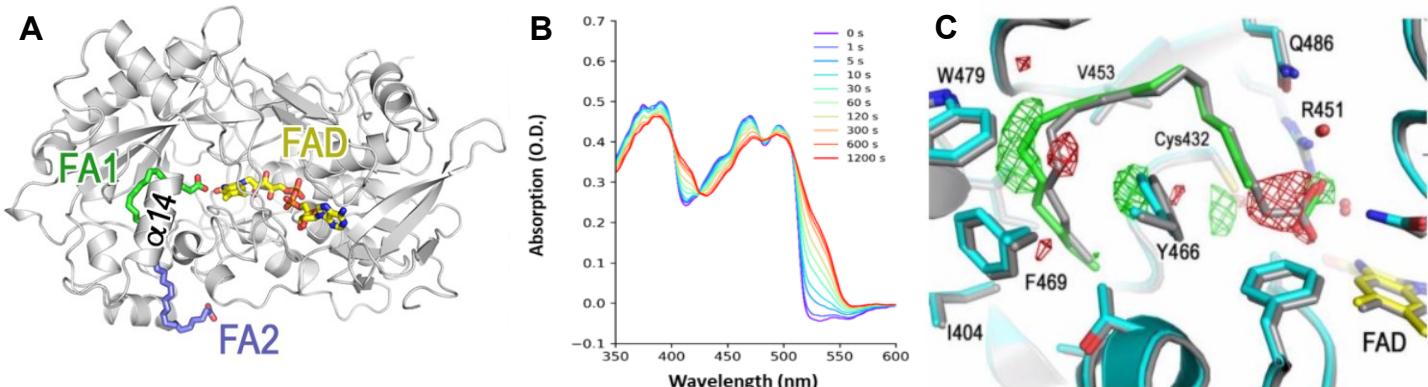


Figure 1. Overall structure of FAP and trapping of a key reaction intermediate at 100 K. (A) Overall structure of FAP, with two substrates stabilized (FA1 and FA2). (B) Building of a key reaction intermediate (FAD in a red-shifted state) along blue-light illumination at 100 K. (C) Experimental difference density map ($F_{\text{light}} - F_{\text{dark}}$) contoured at $\pm 4\sigma$ around the active site substrate, superimposed on the structures of the dark state (gray) and the red-shifted form (cyan, with FAD in yellow and alkane/CO₂ in green). CO₂ formation is clearly visible, together with the retreat of the alkane product and a rotation of the side chain of Y466.

[1] Sorigué D, Légeret B, Cuiné S, Blangy S, Moulin S, Billon E, Richaud P, Brugiére S, Couté Y, Nurizzo D, Müller P, Brettel K, Pignol D, Arnoux P, Li-Beisson Y, Peltier G, Beisson F. (2017) *Science*. **357**, 903.

[2] Sorigué, D., K. Hadjidemetriou, S. Blangy, G. Gotthard, A. Bonvalet, N. Coquelle, P. Samire, A. Aleksandrov, L. Antonucci, A. Benachir, S. Boutet, M. Byrdin, M. Cammarata, S. Carbojo, S. Cuiné, R. B. Doak, L. Foucar, A. Gorel, M. Grünbein, E. Hartmann, R. Hienerwadel, M. Hilpert, M. Kloos, T. J. Lane, B. Légeret, P. Legrand, Y. Li-Beisson, S. L. Y. Moulin, D. Nurizzo, G. Peltier, G. Schirò, R. L. Shoeman, M. Sliwa, X. Solinas, B. Zhuang, T. R. M. Barends, J.-P. Colletier, M. Joffre, A. Royant, C. Berthomieu, M. Weik, T. Domratcheva, K. Brettel, M. H. Vos, I. Schlichting, P. Arnoux, P. Müller, F. Beisson (2021) *Science* **372**, 148.

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