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An Engineered Self-sufficient Biocatalyst Enables Scalable Production of Linear Alpha Olefins from Carboxylic Acids

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ABSTRACT: Fusing the decarboxylase OleT_{JE} and the reductase domain of P450BM₃ creates a self-sufficient protein, OleT-BM₃R, which is able to efficiently catalyze oxidative decarboxylation of carboxylic acids into linear α -olefins (LAOs) under mild aqueous conditions using O₂ as the oxidant and NADPH as the electron donor. The compatible electron transfer system installed in the fusion protein not only eliminates the need for auxiliary redox partners, but also results in boosted decarboxylation reactivity and broad substrate scope. Coupled with the phosphite dehydrogenase based NADPH regeneration system, this enzymatic reaction proceeds with improved product titers of up to 2.51 g L⁻¹ and volumetric productivities of up to 209.2 mg L⁻¹ h⁻¹ at low catalyst loadings (~0.02 mol%). With its stability and scalability, this self-sufficient biocatalyst offers a nature-friendly approach to deliver LAOs.

KEYWORDS: Green chemistry, Biocatalysis, Protein engineering, Oxidative decarboxylation, α -Olefins

Linear α -olefins (LAOs) are next-generation fuels and key feedstock chemicals for the production of surfactants, lubricants, detergents, and polymers.¹⁻⁵ Currently, these olefins are predominantly produced from petroleum via ethylene oligomerization that only yields even-numbered terminal olefins.⁶⁻⁹ Considering dwindling fossil fuel reserves, pursuing a renewable route to access LAOs is of urgency.^{10,11} Pleasingly, abundant and bioavailable fatty acids (FAs) may provide an alternative source for the sustainable production of LAOs through straightforward decarbonylative dehydration or oxidative decarboxylation,¹² which can lead to both even- and odd-numbered LAOs. To achieve this goal, transition-metal catalysts including palladium, rhodium, iridium, and iron have been developed.¹³⁻²⁰ While impressive, these approaches usually suffer from harsh reaction conditions ($T \geq 110$ °C) and require in situ distillation of the product to maintain acceptable α -selectivity. Furthermore, activation of the FAs with stoichiometric anhydrides is often needed, resulting in extra waste. Nowadays, intensifying environmental concerns urge scientists to investigate chemical transformations using green solvents, such as water, or eco-friendly methods, including electrochemistry²¹⁻²⁴ and biocatalysis.²⁵⁻²⁸ Accordingly, we aimed for a green strategy to supply LAOs from FAs.

Recently, a cytochrome P450 enzyme, OleT_{JE}, was found to catalyze the oxidative decarboxylation of saturated FAs into terminal olefins using H₂O₂ as the oxidant.²⁹ H₂O₂ deactivated this biocatalyst at millimolar concentration and these H₂O₂-dependent OleT_{JE} systems had relatively low catalytic efficiency and narrow substrate scope (FAs \geq C8).³⁰⁻³⁵ When coupled with redox partners, OleT_{JE} could catalyze the same reaction with O₂ as the oxidant and NAD(P)H as the electron donor.³⁶⁻³⁹ Addition of putidaredoxin reductase and putidaredoxin (CamAB) to the reaction could increase the reactivity and turnover number of OleT_{JE}.³⁷⁻³⁹ However, preparation of the CamAB-containing lysate and calibration of the lysate versus OleT_{JE} complicated the reaction system and would limit its application. Previous effort to create a self-sufficient enzyme by fusing OleT_{JE} to P450RhF reductase domain achieved H₂O₂-independent catalysis, but this engineered enzyme unfortunately displayed low activity, only being able to decarboxylate FAs C₁₂-C₁₈.³⁶ In this study, a more compatible electron transfer system is installed in an OleT_{JE}-containing fusion protein, resulting in a self-sufficient biocatalyst with increased decarboxylation reactivity. Our approach, employing O₂ as the

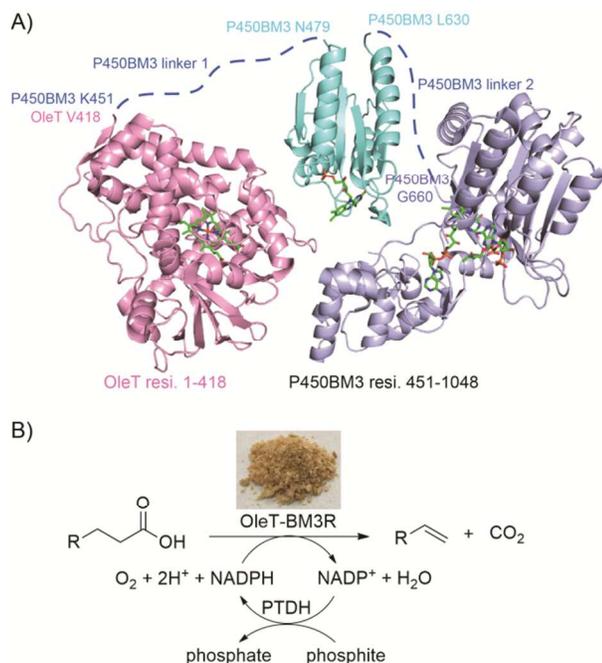


Figure 1. (A) Structural schematic of OleT-BM₃R shown as the cartoon representation. OleT-BM₃R is composed of OleT_{JE} (residues 1-418, PDB code 4L54, pink), FMN binding domain (residues 479-630 of P450BM₃, PDB code 1BVY, cyan) and FAD/NADPH-binding domain (residues 660-1048 of P450BM₃, PDB code 4DQL, blue) of P450BM₃. The cofactors, heme, FMN, FAD and NADP⁺ are displayed as sticks. (B) Oxidative decarboxylation of carboxylic acids to olefins by OleT-BM₃R (pictured in lyophilized form) that uses O₂ with NADPH recycling. PTDH = phosphite dehydrogenase.

oxidant and NADPH as the electron donor, converts structurally diversified carboxylic acids to their corresponding olefins under mild aqueous conditions. This stable fusion protein reduces the effort required for protein production and purification and simplifies the reaction system, making it more suitable for synthetic and industrial application.

To start, we chose CYP_{102A1} (P450BM₃) from *Bacillus megaterium* as the template for the construction of the self-sufficient OleT_{JE} fusion protein. The heme domain of P450BM₃ is fused with the reductase domain (BM₃R) as a single polypeptide, establishing an efficient electron transfer to the heme iron that is not contingent on encountering a discrete redox partner.^{40,41} Powered by this efficient electron transfer chain, P450BM₃'s hydroxylase activity is >1000 fold higher than those of other P450 fatty acid hydroxylases.⁴² Superimposition of their crystal structures revealed a similar structural fold between OleT_{JE}³⁰ and the heme domain of P450BM₃⁴³ (RMSD=3.6 Å) (Figure S1). The replacement of the heme domain of P450BM₃ with OleT_{JE} (residue 1-418) resulted in a soluble protein, named OleT-BM₃R (Figure 1A). Preliminary experiments showed that the fusion protein OleT-BM₃R exhibited high decarboxylation activity towards stearic

Table 1. The oxidative decarboxylation of various saturated FAs by OleT-BM₃R with NADPH recycling^a

Entry	Substrate	Product	Yield [%] ^b
1	CH ₃ (CH ₂) ₁₈ CO ₂ H (1a)	CH ₃ (CH ₂) ₁₆ CH=CH ₂ (2a)	70
2	CH ₃ (CH ₂) ₁₆ CO ₂ H (1b)	CH ₃ (CH ₂) ₁₄ CH=CH ₂ (2b)	73
3	CH ₃ (CH ₂) ₁₄ CO ₂ H (1c)	CH ₃ (CH ₂) ₁₂ CH=CH ₂ (2c)	60
4	CH ₃ (CH ₂) ₁₂ CO ₂ H (1d)	CH ₃ (CH ₂) ₁₀ CH=CH ₂ (2d)	52
5	CH ₃ (CH ₂) ₁₀ CO ₂ H (1e)	CH ₃ (CH ₂) ₈ CH=CH ₂ (2e)	46
6	CH ₃ (CH ₂) ₉ CO ₂ H (1f)	CH ₃ (CH ₂) ₇ CH=CH ₂ (2f)	58
7	CH ₃ (CH ₂) ₈ CO ₂ H (1g)	CH ₃ (CH ₂) ₆ CH=CH ₂ (2g)	70
8	CH ₃ (CH ₂) ₇ CO ₂ H (1h)	CH ₃ (CH ₂) ₅ CH=CH ₂ (2h)	47
9	CH ₃ (CH ₂) ₆ CO ₂ H (1i)	CH ₃ (CH ₂) ₄ CH=CH ₂ (2i)	40
10	CH ₃ (CH ₂) ₅ CO ₂ H (1j)	CH ₃ (CH ₂) ₃ CH=CH ₂ (2j)	70
11	CH ₃ (CH ₂) ₄ CO ₂ H (1k)	CH ₃ (CH ₂) ₃ CH=CH ₂ (2k)	68
12	CH ₃ (CH ₂) ₃ CO ₂ H (1l)	CH ₃ CH ₂ CH=CH ₂ (2l)	11
13	CH ₃ (CH ₂) ₂ CO ₂ H (1m)	CH ₃ CH=CH ₂ (2m)	14

^aReaction conditions: purified OleT-BM₃R (3 μM), substrate (1 mM), catalase (100 U mL⁻¹), NADPH (200 μM), PTDH (2 μM), sodium phosphite (0.01 M), 1 mL scale, rt, 12 h. ^bdetermined by GC or headspace GC (see the supporting information).

acid (FA C₁₈:o) in the presence of oxygen and NADPH at ambient temperature [turnover number (TON) = 2520, turnover frequency (TOF) = 472 h⁻¹, NADPH coupling efficiency = 61%]. In contrast, when the same reaction was performed with OleT_{JE} and free BM₃R, the TOF and NADPH coupling efficiency were only 65 h⁻¹ and 29%, respectively. These results prove an enhanced electron transfer from BM₃R to OleT in our fusion protein. Compared with other redox systems for the decarboxylation of FAs with OleT_{JE}, this self-sufficient enzyme showed advantages in turnover number, rate of reaction, and NADPH coupling efficiency (Table S1). When this reaction was coupled to a phosphite dehydrogenase (PTDH) based NADPH regeneration system,⁴⁴⁻⁴⁶ the overall decarboxylation activity (TON = 2167) was maintained. Isolated BM₃R was susceptible to temperature-induced activity loss (inactivation rate = 0.22 min⁻¹ at 30 °C),⁴⁷ but the OleT-BM₃R fusion protein was stable at 37 °C for 24 h (Figure S2). Moreover, purified OleT-BM₃R can be lyophilized (Figure 1B) and conveniently stored at -20 °C for months without significant activity loss. These results demonstrate that OleT_{JE} is able to complement BM₃R in

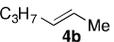
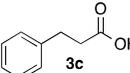
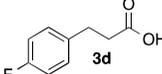
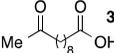
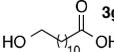
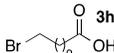
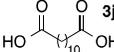
P450BM₃, together forming a structurally stable and functionally active entity.

With the engineered OleT-BM₃R in hand, we proceeded to explore the reactivity and scope of the oxidative decarboxylation of FAs (Table 1). Gratifyingly, both long- and medium-chain saturated FAs **1a-k** successfully participated in this transformation, delivering the desired LAOs **2a-k** up to 73% yield (entries 1-11). Short-chain fatty acids **1l-m** also worked albeit with meager yields (entries 12-13). These results indicate good compatibility between OleT and BM₃R in the fusion protein. Notably, utilizing naturally occurring even-numbered saturated FAs afforded prohibitively expensive odd-numbered terminal olefins, such as **2a** and **2b**, which are inaccessible through the oligomerization of ethylene. Meanwhile, α - and β -hydroxylations occurred alongside the desired decarboxylation process. Formation of the nonvolatile α/β -hydroxylated products was also dependent on chain length (Table S2). The catalytic cycle of OleT_{JE} involves abstraction of a C α /C β hydrogen atom of the carboxylic acid by the iron (IV)-oxo heme π -cation radical intermediate (Compound I), followed by a competition between the \cdot OH rebound and carbon-carbon scission that delivers the corresponding hydroxylation and decarboxylation products.^{30,48-50} This mechanism guarantees the exclusive formation of α -alkenes from the decarboxylation pathway. As a result, this mild enzymatic approach displays advantages over synthetic methods, which employ transition-metal catalysts under harsh conditions and thus partially form the thermodynamically stable internal alkenes.

To extend the synthetic application of this enzymatic transformation, we further explored its substrate scope and functional group tolerance (Table 2). To our delight, both cyclic and acyclic α -branched carboxylic acids **3a-b** underwent the expected reaction, providing the corresponding olefins in reasonable yields (entries 1-2). Similarly, phenylpropanoic acids **3c-d** smoothly delivered the respective vinylbenzenes **4c-d** with improved yields (entries 3-4) and only traces of hydroxylation products were formed in these reactions (Table S2). Functional groups like olefins and ketones were tolerated and a free-hydroxyl moiety was also untouched during the reaction (entries 5-7). Remarkably, sensitive groups such as bromo and aldehyde, which are vulnerable under conditions using transition-metal catalysts, survived due to our mild enzymatic conditions (entries 8-9). When diacid **3j** was examined, the anticipated oxidative decarboxylation occurred on both acids (entry 10). Interestingly, only terminal diene was formed. Although no ω -alkenoic acid intermediate was found, its α/β -hydroxy derivatives were detected (Table S2), indicating that the dicarboxylic acid underwent tandem decarboxylation via ω -alkenoic acid intermediate.

To testify its potential for preparative application, this enzymatic transformation was performed in a concentrated manner with stearic acid as a substrate (Table 3). Simply increasing the substrate concentration from 1 mM to 10 mM led to a slightly diminished yield but dramatically improved product titer and volumetric productivity

Table 2. Further substrate scope study.^a

Entry	Substrate	Product	Yield [%] ^b
1			30
2			30
3			78
4			44
5			71
6			48
7			42
8			40
9 ^c			11 ^d
10			42

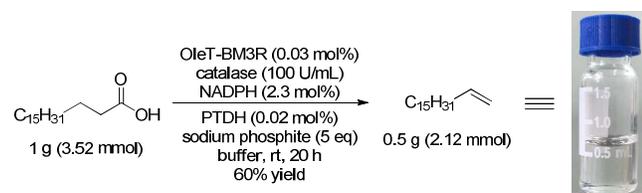
^aReaction conditions: see conditions in Table 1. ^bdetermined by headspace GC. ^csubstrate (10 mM), sodium phosphite (0.05 M), 30 mL scale, 20 h. ^disolated yield.

(entry 1 vs 2). Compared to the previous OleT/CamAB/FDH cascade system,³⁷ our newly developed OleT-BM₃R/PTDH coupled system offers higher product titer (1.40 g L⁻¹ vs. 0.93 g L⁻¹) and volumetric productivity (116.7 mg L⁻¹ h⁻¹ vs. 42.5 mg L⁻¹ h⁻¹), presumably because of the greater TOF (472 h⁻¹ vs. 53 h⁻¹) and NADPH coupling efficiency (61% vs. 25%). Meanwhile, simultaneously elevating the concentrations of both catalyst and substrate further benefited the product titer and volumetric productivity at the cost of the conversion (entries 3-4). The product titer of our OleT-BM₃R/PTDH system could reach as high as 2.51 g L⁻¹ while the volumetric productivity climbed up to 209.2 mg L⁻¹ h⁻¹ at a low catalyst loading (~0.02 mol%) (entry 4). Notably, cell-free lysate allowed the production of 1.26 g L⁻¹ (5.28 mM, 53% yield) 1-heptadecene from 10 mM stearic acid without the

Table 3. Oxidative decarboxylation of stearic acid at various substrate concentrations by OleT-BM₃R with NADPH recycling.^a

Entry	OleT-BM ₃ R [μM]	Substrate [mM]	Yield [%]	Titer [g L ⁻¹]	Volumetric productivity [mg L ⁻¹ h ⁻¹]
1	3	1	73	0.17	14.5
2	3	10	59	1.40	116.7
3	9	20	40	1.91	159.2
4	9	40	26	2.51	209.2
5	5b	10	53	1.26	105.0
6	3c	10	54	1.29	107.5

^aReaction conditions: OleT-BM₃R (indicated), catalase (100 U mL⁻¹), NADPH (200 μM), PTDH (2 μM), sodium phosphite (0.05 M), 1 mL scale, rt, 12 h; ^bCell-free lysate without NADPH supplement. ^cThe activity of lyophilized OleT-BM₃R was tested after 2-month storage at -20 °C.

**Figure 2.** Gram scale enzymatic oxidative decarboxylation of stearic acid.

need for protein purification (entry 5). This cell-free lysate system is economically desirable as no additional NADPH is required. Additionally, lyophilized OleT-BM₃R exhibited comparable reactivity after two-month storage at -20 °C (entry 6). Moreover, our enzymatic system proved scalable, and 1g of stearic acid was successfully converted to 1-heptadecene in 60% yield (Figure 2). As the obtained product is particularly expensive, our enzymatic reaction provides this LAO not only through a sustainable approach, but also in a cost-effective fashion. Overall, the outstanding stability and scalability of this self-sufficient biocatalyst demonstrates encouraging potential for preparation of LAOs.

In summary, we engineered a catalytically self-sufficient fusion protein, OleT-BM₃R, for oxidative decarboxylation of FAs without the need for auxiliary redox partners. The simplified reaction could be set up with less effort in protein production and purification and performed under mild conditions in aqueous solution using O₂ as the oxidant and NADPH as the electron donor. It could also use inexpensive sodium phosphite as the final electron source when coupled with a PTDH based NADPH regeneration system. A similar structure between OleT_{JE} and the heme domain of P450BM₃, a good compat-

ibility of OleT_{JE} and BM₃R, and the structural elements in P450BM₃ that facilitate domain-domain interactions and electron transfer processes altogether contribute to the boosted decarboxylation reactivity and broad substrate scope of the fusion protein. The stability and scalability of this enzyme make it a useful biocatalytic platform for the sustainable synthesis of LAOs. In addition, high ferricyanide and cytochrome c reduction rates (9850 min⁻¹ and 3330 min⁻¹, respectively) were determined for this fusion protein, indicating that it was powered by the rapid electron transfer in BM₃R.⁵¹ But a comparable reaction rate of P450BM₃-catalyzed fatty acid hydroxylation (usually TOF > 400 min⁻¹) was not observed.⁴² Thus, further engineering this fusion protein for enhanced interaction and electron transfer between the FMN binding domain and OleT, which might additionally benefit its catalytic performance, is worth future exploration. Meanwhile, production of LAOs in microbial platforms with this biocompatible catalyst is also underway in our laboratories.

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Author Contributions

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ASSOCIATED CONTENT

Supporting Information includes Experimental Procedures GC/Headspace GC results and ¹H Spectra data. This material is available free of charge via the Internet at <http://pubs.acs.org>.

ABBREVIATIONS

LAOs, linear α-olefins; FAs, fatty acids; TON, turnover number; TOF, turnover frequency; PTDH, phosphite dehydrogenase; NADP, nicotinamide adenine dinucleotide phosphate.

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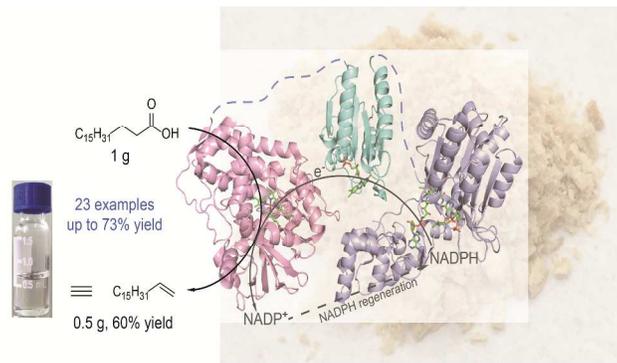
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