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Unnatural Biosynthesis by an Engineered Microorganism with Heterologously Expressed Natural Enzymes and an Artificial Metalloenzyme

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Abstract

Synthetic biology enables microbial hosts to produce complex molecules that are otherwise produced by organisms that are rare or difficult to cultivate, but the structures of these molecules are limited to those formed by chemical reactions catalyzed by natural enzymes. The integration of artificial metalloenzymes (ArMs) that catalyze unnatural reactions into metabolic networks could broaden the cache of molecules produced biosynthetically by microorganisms. We report an engineered microbial cell expressing a heterologous biosynthetic pathway, which contains both natural enzymes and ArMs, that produces an unnatural product with high diastereoselectivity. To create this hybrid biosynthetic organism, we engineered *Escherichia coli* (*E. coli*) with a heterologous terpene biosynthetic pathway and an ArM containing an iridium-porphyrin complex that was transported into the cell with a heterologous transport system. We improved the diastereoselectivity and product titer of the unnatural product by evolving the ArM and selecting

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Author contributions: J.F.H., D.S.C., A.M., and J.D.K. conceived the project and all authors participated in designing experiments. J.H., Z.L., and B.L.B. conducted all experiments and collected data for the project. All authors interpreted the data and wrote the manuscript.

Declarations

Competing interests: JDK has a financial interest in Amyris, Lygos, Demetrix, Maple Bio, Napigen, Apertor Pharma, Ansa Biotechnologies, Berkeley Yeast, and Zero Acre Farms.

the appropriate gene induction and cultivation conditions. This work shows that synthetic biology and synthetic chemistry can produce, together with natural and artificial enzymes in whole cells, molecules that were previously inaccessible to nature.

Keywords

artificial metalloenzymes; biosynthesis; engineered microorganism

Introduction

Molecules produced by natural biosynthetic pathways are the inspiration or the actual matter in many products, ranging from medicines and agrochemicals to commodity chemicals and biofuels. Although many molecules with structures that are the envy of chemists are produced by these natural or engineered biosynthetic pathways, the structures remain limited by the available chemical reactions catalyzed by natural enzymes. If artificial enzymes could be incorporated into such biosynthetic pathways, then unnatural products could be produced from a combination of nature's reactions and chemists' imaginations.

Recently, artificial metalloenzymes (ArMs) have been created that catalyze a series of unnatural reactions²⁻¹³. However, most of these reactions do not occur on the types of molecules that could be produced by biosynthesis. In contrast, an iridium-containing analog of a heme enzyme catalyzes the cyclopropanation of unconjugated, hindered, terminal alkenes that are present in natural products, such as terpenes⁴. This reactivity provides the potential to create an artificial biosynthetic pathway that combines natural pathways manipulated by synthetic biology with the catalysis created by ArMs.

The assembly of artificial metalloenzymes has advanced from *in vitro* reconstitution of purified proteins with organometallic complexes to assembly of ArMs in the periplasm^{13,14} or on the outer-membrane^{15,16} of living cells. However, methods to assemble ArMs in more complex intracellular environments are lacking^{17,18}. We hypothesized that artificial cytochrome P450s containing synthetic heme derivatives could be generated in the cytoplasm by employing the natural heme transport machinery¹⁹ and the affinity of hemoproteins to heme derivatives. Thus, artificial cytochrome P450s are potential ArMs for conducting unnatural biotransformations *in vivo* in a fashion that enables the assembly of artificial biosynthetic pathways.

Terpenes are the largest and most structurally diverse family of natural products²⁰. Their core skeletons are assembled in one step and modified by a suite of tailoring enzymes including cytochrome P450s²¹. While native P450s typically catalyze oxidations of terpenes, extensively engineered P450s, such as P450_{BM3}²², and artificial P450s substituted with abiotic metalloporphyrins, could catalyze unnatural transformations^{3-5, 23,24} that would introduce an artificial dimension to the diversification of terpene structures. Here, we report the combination of limonene biosynthesis by a series of heterologously expressed natural enzymes and the catalysis by an artificial metalloenzyme Ir-CYP119 (CYP119 containing Ir(Me)MPIX as cofactor; CYP119, a P450 enzyme from *Sulfolobus solfataricus*; MPIX, mesoporphyrin IX) to produce a terpenoid that was previously inaccessible in a microbial

host. This unnatural combination of synthetic biology and synthetic chemistry mimics the production of a terpene and subsequent conversion of this hydrocarbon to a terpenoid by P450 enzymes during natural biosynthesis.

Our overall strategy for production of the unnatural terpenoid is depicted in Fig. 1. The artificial metalloenzyme is assembled in *E. coli* with the aid of a cofactor transport system and overexpression of the apo-enzyme. In concert, the same *E. coli* is engineered with the pathway for production of a natural product that serves as the substrate for the artificial metalloenzyme. This combination of natural and unnatural reactions in the same cell would produce an unnatural derivative of a classic natural product.

Results And Discussion

Assembling the ArM

in vivo The first major challenge facing this integration of an artificial metalloenzyme with a cellular biosynthetic process is to assemble such enzymes in the cytoplasm²⁵. Previously, we reported the *in vitro* construction of Ir-CYP119 by incorporation of Ir(Me)MPIX into the heme binding site of the purified apo-form of CYP119. To construct this artificial enzyme in a cell, the cofactor must enter into the cytoplasm from the medium through the cell membranes. Metal-substituted heme derivatives have been reported to be transported across cell membranes when the outer-membrane receptor ChuA is overexpressed^{26–28}. Thus, we first considered whether the co-expression of such a heme transporter and CYP119 would enable the artificial cofactor Ir(Me)MPIX to be transported into *E. coli* and incorporated into the apo-CYP119 mutant in the cytoplasm.

To test the feasibility of assembling Ir-CYP119 and catalyzing a reaction in *E. coli*, we studied the cyclopropanation of the monoterpene (–)-carvone. We used the diastereoselectivity of the reaction as a parameter to distinguish any background reaction or reaction catalyzed by free cofactor from the reaction catalyzed by the assembled artificial metalloenzyme. During our previous study, Ir-CYP119 generated from *in vitro* reconstitution of Ir(Me)MPIX and purified apo-CYP119 mutants catalyzed the cyclopropanation of (–)-carvone with EDA (ethyl diazoacetate) to give an 8 : 1 : 1 : 1 ratio of diastereomeric products. The major diastereomer of this reaction is a minor diastereomer of the 1 : 1 : 3 : 3 ratio of diastereomers formed from the analogous reaction catalyzed by the free iridium porphyrin complex⁴. Thus, a reaction that forms a mixture of diastereomers resembling the ratio produced by the pure ArM *in vitro* would denote reaction by the assembled ArM in the cell.

To this end, *E. coli* BL21(DE3) cells co-expressing the heme transporter ChuA and the CYP119 mutant were supplemented with 10 μM Ir(Me)MPIX, and the cell pellets containing *in vivo*-assembled Ir-CYP119 were directly employed for catalyzing the reaction of (–)-carvone with EDA after re-suspension in reaction buffer. However, the diastereomeric ratio (dr) of the whole-cell reaction was 1.7 : 1.0 : 2.7 : 3.3 (Supplementary Fig. 1a), indicating that the majority of the product of the whole-cell system was formed from a reaction catalyzed by the free iridium porphyrin complex, rather than by an assembled ArM.

In addition, no product was detected when *E. coli* cultures were incubated in the absence of Ir(Me)MPIX (Supplementary Fig. 1a).

To reduce interference from the background reactivity of the free iridium porphyrin, we evaluated the reaction of (–)-carvone with EDA catalyzed by cells grown in the presence of Ir(Me)MPIX in concentrations ranging from 10 μM to 0.02 μM. These experiments showed that the dr of the cyclopropane product was much higher (23 : 3.5 : 1.0 : 1.2) when the concentration of Ir(Me)MPIX was 0.02 μM (Fig. 2a). We surmised that the lower concentrations of Ir(Me)MPIX reduced the amount of product formed by a background reaction catalyzed by the cofactor unbound to CYP119. Control experiments showed that the dr was low for products formed in *E. coli* expressing ChuA without co-expression of CYP119 (Supplementary Fig. 1b). Under the conditions of this control experiment, the cyclopropanation would be catalyzed by the free iridium cofactor.

Next, we sought to combine the Ir-CYP119 with a natural biosynthetic pathway to produce an unnatural terpenoid. The analysis of Ir(Me)MPIX uptake for cyclopropanation in whole cells described above was conducted with carvone because we had shown previously that cyclopropanation of this terpenoid occurred in high yield and high diastereoselectivity⁴. However, the biosynthesis of (–)-limonene, which has the same carbon skeleton as (–)-carvone, occurs with much higher titer^{29,30}, so we sought to assess the potential to combine heterologous biosynthesis with artificial metalloenzymes by generating unnatural derivatives of limonene.

Initial attempts to express a heterologous biosynthetic pathway to form limonene in concert with expression of CYP119 regulated by a T7 promoter caused the titer of limonene to decrease drastically from 188 mg/L without co-expression of CYP119 to 28 mg/L with co-expression (Supplementary Fig. 2a). To restore the production of limonene, we expressed CYP119 with a weaker promoter. However, when the reaction of exogenously added (–)-carvone with EDA was conducted with cells co-expressing ChuA and CYP119 with the weaker *lacUV5* promoter, the cyclopropane was formed with low diastereoselectivity, again implying that assembly of the artificial metalloenzyme was inefficient (Supplementary Fig. 3). Thus, we sought conditions that would assemble the artificial metalloenzyme more efficiently at low expression levels of CYP119 to generate product with high diastereoselectivity.

To increase the efficiency of the cellular uptake of Ir(Me)MPIX, we replaced the ChuA transporter with a heme transport system encoded by the *hug* operon from *Plesiomonas shigelloides*, which has been used less frequently than ChuA, but has been shown previously to increase expression yields of hemoglobin to unusually high levels with heme supplementation^{31,32}. Reactions of (–)-carvone with EDA catalyzed by *E. coli* cells expressing the HUG transport system, in concert with CYP119 (expressed from the *lacUV5* promoter), formed the cyclopropane product with a high diastereoselectivity of 82% (the % diastereoselectivity in this case is the percentage of the major diastereomer in the 24.0 : 3.1 : 1.0 : 1.2 ratio of diastereomeric products) under conditions that are identical to those of the reactions with cells expressing the ChuA transporter (Fig. 2b).

To elucidate the origin of the higher efficiency of the assembly of ArMs in cells expressing the HUG transport system, the distribution of iridium was analyzed by ICP-MS (inductively coupled plasma mass spectrometry) (Fig. 2c). In cells harboring CYP119 in the presence of 0.1 μ M Ir(Me)MPIX, 65% of the iridium was located in the cytoplasm of the cells expressing the HUG transport system, while only 10% of the iridium was located in the cytoplasm of the cells expressing no transporter. Further, the cytoplasmic fraction isolated from the cells co-expressing the HUG transport system catalyzed the cyclopropanation of (–)-carvone with a diastereoselectivity of 80%, and the yield of cyclopropane was more than 13 times higher than that of the reaction catalyzed by the cytoplasmic fraction from cells expressing no transporter (Supplementary Fig. 4). These results indicate that the HUG system significantly enhances the uptake of Ir(Me)MPIX into the cytoplasm.

Cyclopropyl Limonene Biosynthesis

Having achieved the diastereoselective cyclopropanation of (–)-carvone with EDA in cells expressing CYP119, even at a relatively low level, we sought to integrate such reactivity with the biosynthesis of limonene from a simple sugar (Fig. 3a). Again, we assessed whether the expression of the artificial metalloenzyme would affect the production of limonene. The production of limonene was unaffected by the expression level of CYP119 achieved with the pJHA135 plasmid (Fig. 3b). Moreover, the limonene titer was unaffected by the presence of 0.1 μ M Ir(Me)MPIX (Supplementary Fig. 2b). Finally, both Ir(Me)MPIX and cyclopropyl limonene did not inhibit cell growth within the concentration range relevant for our study (Supplementary Fig. 5).

The cyclopropanation of limonene added exogenously to the whole-cell system catalyzed by *E. coli* expressing the *hug* operon and CYP119 in the M9-rich medium was then evaluated. These cells produced the cyclopropyl limonene with a diastereoselectivity of 52% (1.0 : 3.3 : 1.1 : 1.0 dr) (Fig. 3c). This ratio of diastereomers is distinct from that of the reaction catalyzed by free Ir(Me)MPIX (1.0 : 1.2 : 3.5 : 4.2 dr) and the diastereoselectivity was fortuitously higher than that of the reaction catalyzed by the same Ir-CYP119 mutant *in vitro* (1.0 : 2.4 : 2.3 : 2.9 dr) (Fig. 3c). We suggest that the higher diastereoselectivity in the whole-cell system results from a greater stability of the enzyme *in vivo* than *in vitro*.

To fully combine terpene production and artificial catalysis within the same cell, gene expression for limonene biosynthesis and CYP119 was induced concurrently with addition of Ir(Me)MPIX (Fig. 3d). After overnight incubation, the synthesis of cyclopropyl limonene was initiated by addition of EDA. Under these conditions, 250 μ g/L of cyclopropyl limonene was produced after 4.5 h with diastereoselectivity (43%, 1.0 : 3.2 : 1.8 : 1.5 dr) that was within 10% of that of the whole-cell reaction with exogenously added limonene. This similarity in diastereoselectivity indicates that the cyclopropyl limonene is formed from the combination of the heterologous biosynthesis of limonene and cyclopropanation catalyzed by the ArM. To assess the possibility that formation of the cyclopropane was catalyzed by the Ir-CYP119 released from dead cells, the cells were separated from the culture by centrifugation. The catalytic activity of the supernatant and of resuspended cells was evaluated by the reaction of (–)-carvone and EDA. The formation of cyclopropyl carvone with resuspended cells occurred with high diastereoselectivity and a yield more than 10

times that of the reaction conducted with the supernatant (Supplementary Fig. 6). These data show that the product was produced primarily by intact cells.

Increasing diastereoselectivity and titer

To increase the diastereoselectivity and titer of the cyclopropyl limonene from the artificial pathway, we evolved the CYP119 mutant. We constructed a small library of CYP119 mutants containing additional changes to the amino acids around the binding site of the cofactor and tested the diastereoselectivity of these mutants for the cyclopropanation of limonene in whole cells (Supplementary Fig. 7). After three rounds of screening, mutant P/R256W/V254A (P designates the parent) catalyzed the cyclopropanation of limonene with a higher diastereoselectivity of 72% (1.0 : 9.6 : 1.7 : 1.1 dr) and a higher yield (Fig. 4a,b). This evolved mutant was then introduced into the *E. coli* strain containing the limonene pathway and *hug* operon to perform artificial production of cyclopropyl limonene. With the procedures previously developed (Fig. 3d), the titer and diastereoselectivity of the products formed from this new pathway were 265 $\mu\text{g/L}$ and 59% (1.0 : 7.0 : 2.3 : 1.5 dr) respectively (Supplementary Fig. 8).

The conversion efficiency of limonene by natural P450s in heterologous hosts is known to be low^{30,33}, and the titer of cyclopropyl limonene is limited, in part, by the volatility of limonene. The titer of limonene is just 2 mg/L when the cells are grown without an organic overlay, but is 145 mg/L when the cells are grown with a dodecane overlay (Supplementary Fig. 9). However, an overlay is used only if limonene is the final product; it would be incompatible with our system in which we must preserve limonene intracellularly as a substrate for the ArM. Thus, we sought to optimize the titer of cyclopropyl limonene by alternate methods. To do so, we investigated the potential toxicity of EDA to *E. coli* in the process of producing cyclopropyl limonene and investigated the effect of adding EDA portion-wise (Supplementary Fig. 10). We found that addition of EDA at an initial concentration of 2 mM and refeeding EDA three times over 10 h led to a titer of cyclopropyl limonene that was almost 3 times its initial value (from 258 $\mu\text{g/L}$ to 760 $\mu\text{g/L}$) with little change in diastereoselectivity (54%, 1.0 : 6.4 : 2.6 : 1.9 dr) (Fig. 4c). Additional increases in product titer would likely be gained by further optimization of the protocol for feeding EDA, investigation of additional enzyme mutants, and manipulating the expression levels of proteins in the pathway.

Conclusions

Production of an unnatural product using a microorganism engineered with a heterologous natural product biosynthetic pathway, an ArM, and a heterologous cofactor transport system demonstrates the feasibility of creating organic molecules by artificial biosynthesis comprising synthetic biology and synthetic chemistry. One can envision this combination of biosynthesis and artificial metalloenzymes being applied to engineered pathways that produce unusual core structures, with enzymes that catalyze functionalization of C-H bonds to install abiotic groups at positions where P450s might operate naturally, or with the artificial metalloenzyme operating in the middle of a pathway followed by diversification of the abiotic product with evolved downstream enzymes. One can even envision creating

reagents, such as diazoesters, biosynthetically based on known pathways to these species³⁴. Such goals could be achieved in *E. coli* or in a range of organisms that produce diverse classes of natural products, such as polyketides or various alkaloids, thus opening a realm of biosynthesis beyond that limited by the scope of natural reactions.

Methods

General procedure for the artificial biosynthesis of cyclopropyl limonene

BL21(DE3) cells were transformed with a plasmid (pJBEI6410) coding for enzymes in limonene biosynthetic pathway and a plasmid (pJHA135 or pJHA170) encoding both the heme transport system HUG and the corresponding CYP119 mutant. Individual colonies from freshly transformed plates (incubated at 30 °C) were inoculated into 2 mL M9-rich medium supplemented with 100 µg/mL carbenicillin and 25 µg/mL kanamycin. The cultures were grown overnight at 30 °C, 200 rpm for 18 h. In a 50 mL glass culture tube, 10 mL of M9-rich medium was inoculated with 100 µL of overnight starter culture and incubated at 37 °C, 200 rpm to an OD₆₀₀ of ~1. The cultures were then induced by adding 50 µM IPTG, at which time 2 µL of a 0.5 mM solution of Ir(Me)MPIX in DMSO was added to give a final Ir(Me)MPIX concentration of 0.1 µM. Upon induction, the incubation temperature was reduced to 30 °C, and expression was allowed to continue for 18 h at 200 rpm. To the 10 mL culture was added 21.2 µL EDA dissolved in 100 µL ethanol to give a final EDA concentration of 20 mM and a final ethanol concentration of 1% (v/v), and the mixture was further incubated at 30 °C, 200 rpm for 4.5 h. To analyze the titer of cyclopropyl limonene, aliquots of cell culture (1.4 mL) were removed from the tube and extracted with an equal volume of ethyl acetate. The organic layer (1 mL) was concentrated to 90 µL with a Speedvac, and to this concentrate was added 10 µL of ethyl acetate containing 1 g/L (-)-carvone (carvone used as internal standard). The final solution was analyzed by GC-MS.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Data availability:

All data are available in the main text or the Supplementary Information.

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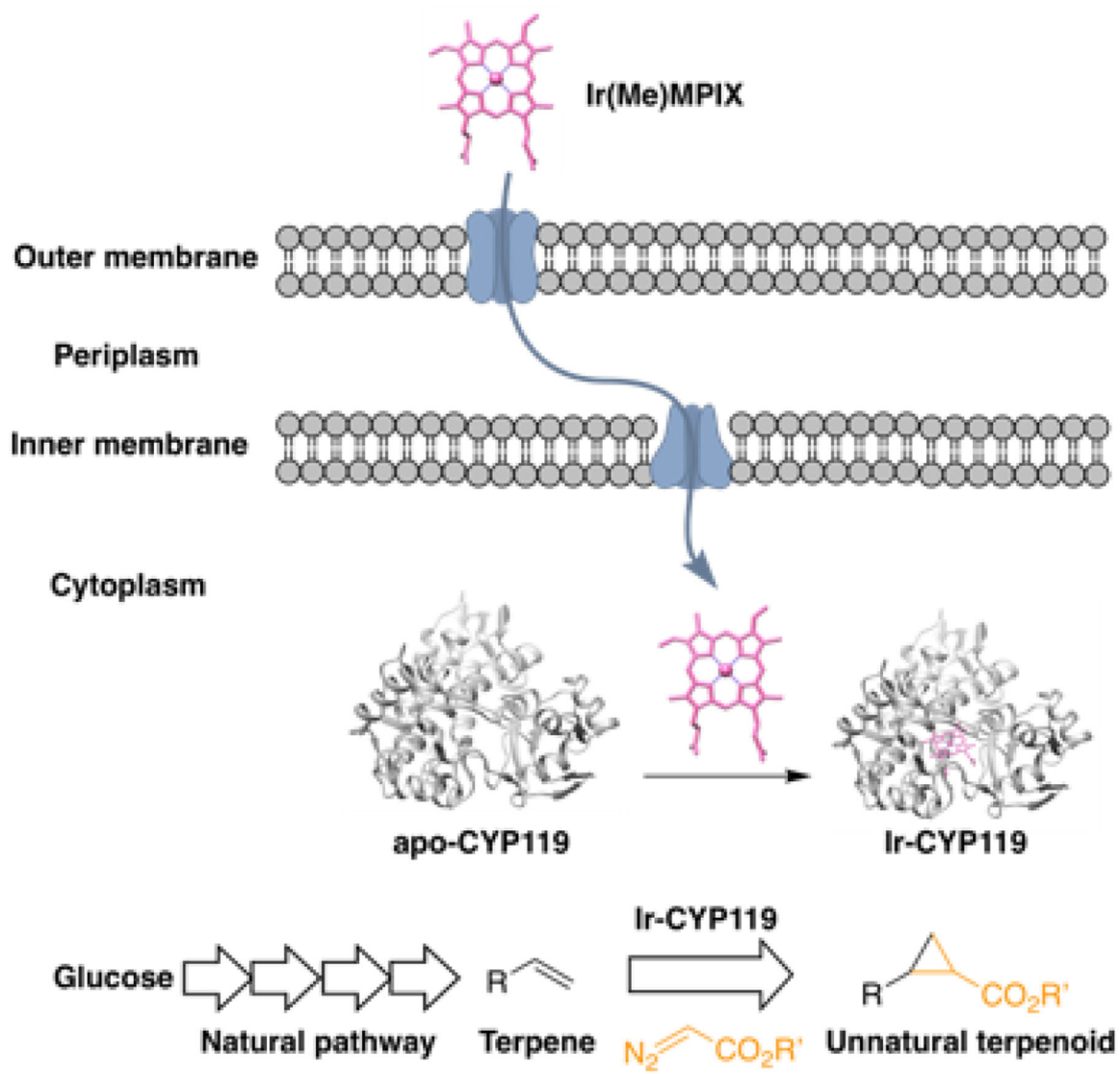
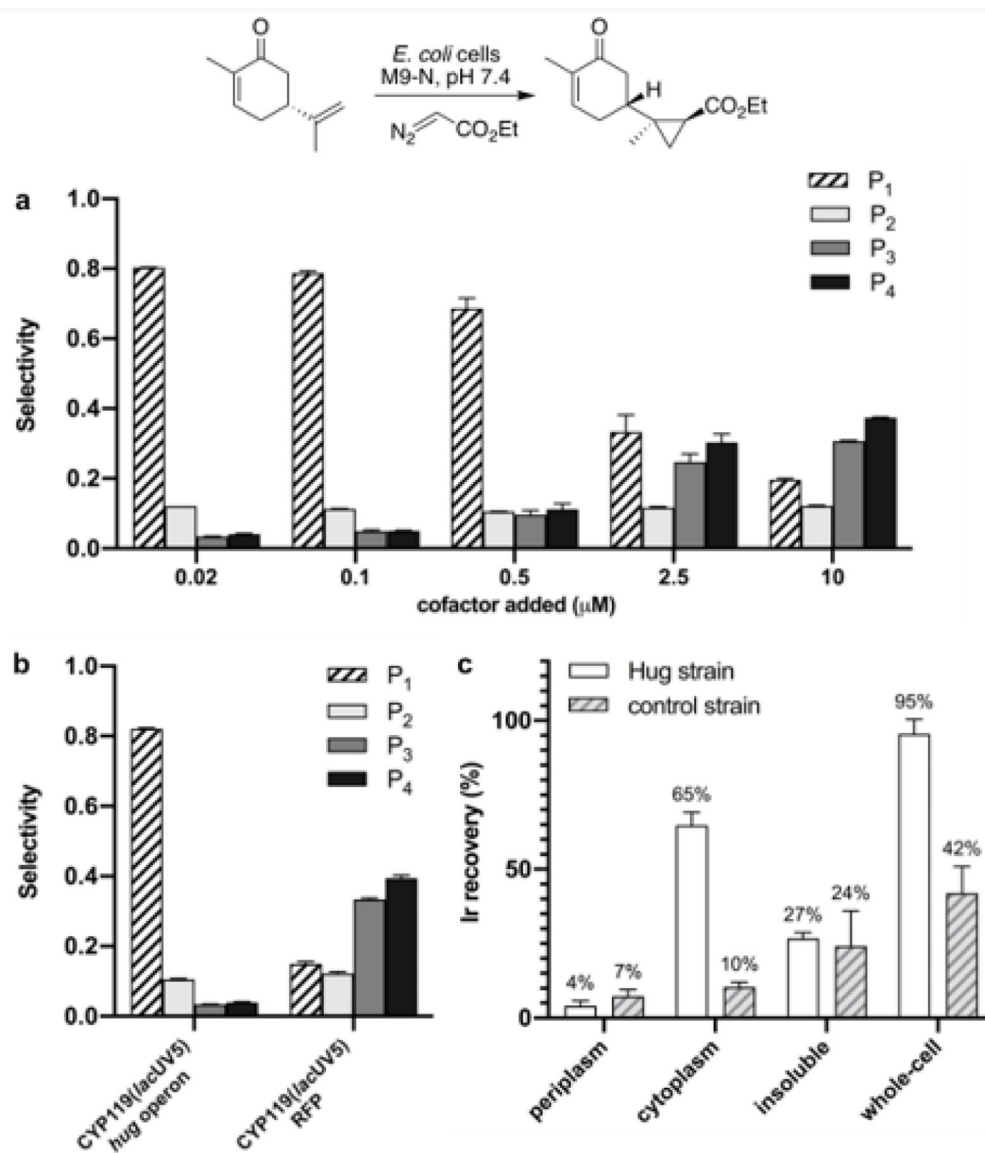


Figure 1.
A schematic representation of a reaction sequence to produce an unnatural terpenoid by combining a heterologously expressed natural biosynthetic pathway and an ArM in *E. coli*.

**Figure 2.**

The whole cells containing Ir-CYP119 catalyze the cyclopropanation of (-)-carvone with high diastereoselectivity. a, Decreasing the concentration of Ir(Me)MPIX increased the diastereoselectivity of whole-cell reactions. The identity of the major diastereomer has been reported previously⁴, the conformations of minor diastereomers are not known. b, *E. coli* cells co-expressing CYP119 (with lacUV5 promoter) and the hug operon could maintain the high diastereoselectivity under low expression level of CYP119. *E. coli* cells containing CYP119 (with lacUV5 promoter) and an RFP-coding plasmid are shown as control (RFP, Red Fluorescent Protein). Both strains were cultivated in the presence of 0.1 μM Ir(Me)MPIX. c, The percentages of added iridium (0.1 μM) that were recovered from the subcellular fractions of *E. coli* cells. Hug strain: *E. coli* co-expressing the HUG system and CYP119 (with lacUV5 promoter). Control strain: *E. coli* co-expressing RFP and CYP119 (with lacUV5 promoter). P1-P4 are the four diastereomeric products numbered in

the order of elution by GC (gas chromatography). Selectivity in the plots corresponds to the ratio of each diastereomer versus the total of the four diastereomeric products. All data are shown as the average from three biological replicates, with error bars indicating 1 standard deviation.

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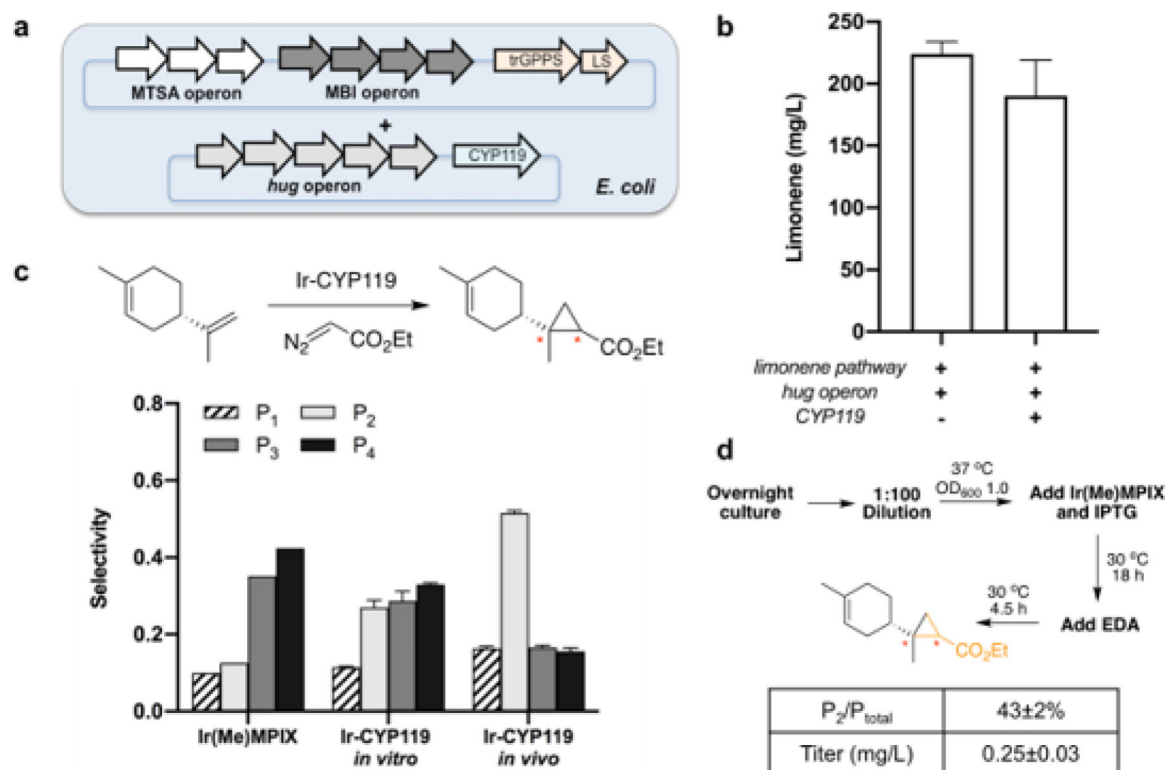
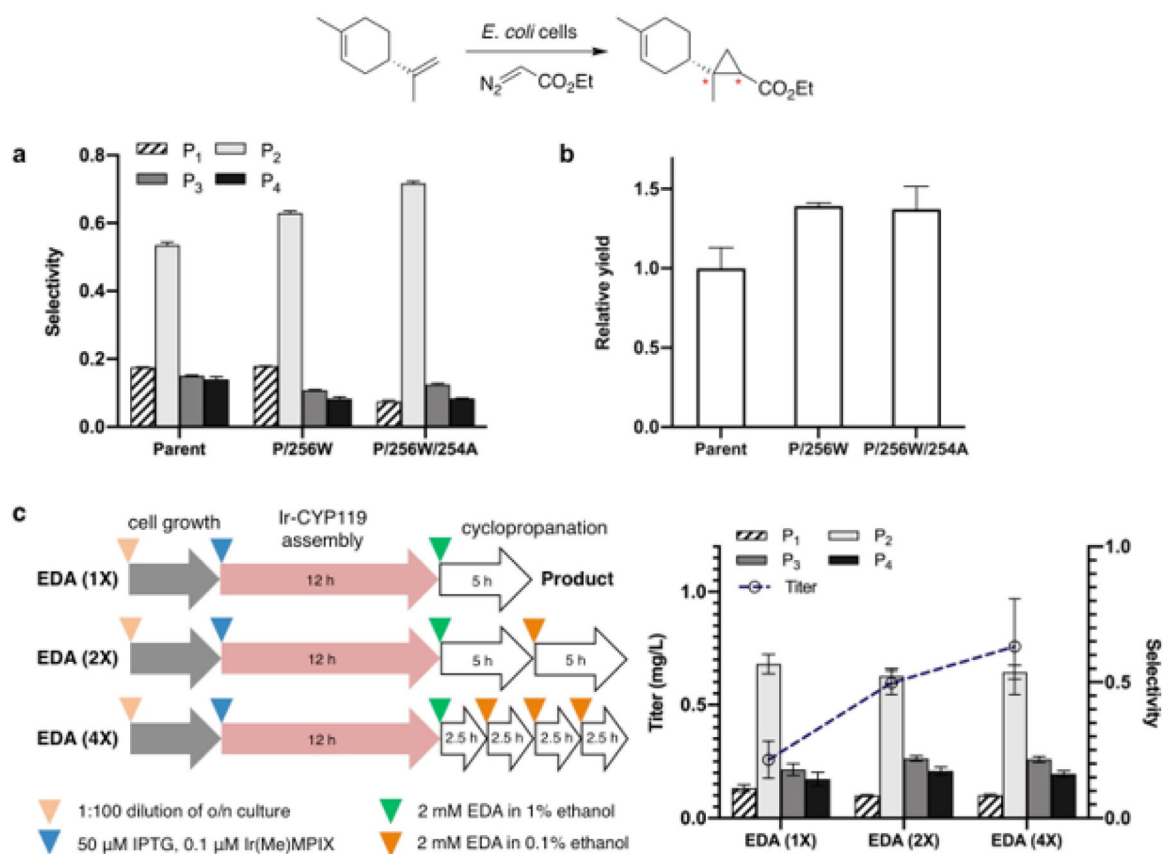


Figure 3.

Combining limonene biosynthesis with ArM. a, A schematic diagram of *E. coli* strain that combines the limonene biosynthetic pathway and the ArM that catalyzes cyclopropanation of limonene. The string of genes at the top represents the plasmid harboring the mevalonate-based limonene biosynthetic pathway (pJBEI6410)29. The bottom one represents the plasmid harboring the HUG system and CYP119 mutant (pJHA135). b, (-)-Limonene was produced with high titer in *E. coli* strains expressing the limonene pathway/hug operon (pJBEI6410+pJHA047) and limonene pathway/hug operon/CYP119 (pJBEI6410+pJHA135), respectively. c, *E. coli* cells expressing Ir-CYP119 catalyzed the cyclopropanation of exogenously added (-)-limonene with diastereoselectivity distinct from that of the free cofactor and enhanced over that of the reaction *in vitro*. d, The cyclopropyl limonene was produced diastereoselectively from glucose in *E. coli* expressing the limonene pathway/hug operon/CYP119. P1-P4 are the four diastereomeric products numbered in the order of elution by GC. The selectivity of P1-P4 corresponds to the ratio of each diastereomer versus the total of the four diastereomeric products. P_{total} is the sum of four diastereomeric products. All data from whole-cell reactions are shown as the average from three biological replicates, with error bars indicating 1 standard deviation.

**Figure 4.**

Increasing the diastereoselectivity and titer of cyclopropyl limonene by directed evolution and process optimization. a, Directed evolution of CYP119 mutants with higher diastereoselectivity. b, The evolved mutants also catalyzed the cyclopropanation in higher yields. The relative yield is the amount of cyclopropyl limonene formed from the whole-cell reaction catalyzed by evolved CYP119 mutants relative to that of the reaction with the parent mutant. c, The improved titer of cyclopropyl limonene with evolved CYP119 mutant (C317G, T213G, V254A, L155W, R256W) and batch-wise addition of EDA. The diagram shows different conditions used for the production of cyclopropyl limonene by the artificial biosynthetic pathway. The concentrations of reagents listed in the diagram are the final concentrations in culture media. All data are shown as the average from three biological replicates, with error bars indicating 1 standard deviation.