Direct Ligation of Carrier Protein-Bound Thioesters: A Versatile Method for the Characterization of Fatty Acid Tailoring Enzymes during Lipopeptide Biosynthesis

| Table S1. Oligonucleotide | pairs used for | or PCR ar | nplification | reactions. |
|---------------------------|----------------|------------|--------------|------------|
| | pulls used it | n i cit ui | Inpinioution | reactions. |

| Oligonucleotide Primers (5'-3') | Restriction Site | Expression vector | target gene |
|--|---------------------|----------------------|------------------|
| AAAAAA <u>GGATCC</u> ATGAGTACGGACCCCA AGTCGGTTG | BamHI | nOTev | ACP (Sco3249) |
| AAAAAA <u>AAGCTT</u> TCACGCCGCTTCCAGA CCCG | HindIII | pQTev | |
| AAAAAA <u>GAATTC</u> ACGCAACGCGAAGAAG AGCTGGCC | EcoRI | ET2 2a(+) | hxcO |
| AAAAAA <u>CTCGAG</u> CGGGCGTACTCCGGCC TGCA | XhoI | pET28a(+) | |
| AAAAAA <u>GAATTC</u> CCGAAGCTGCGGATCG CAGTCG | EcoRI | nET2 8a(+) | hcmO |
| AAAAAACTCG <u>CTCGAG</u> CGGCGGCGGCAG CGGTG | XhoI | pET28a(+) | ncmO |

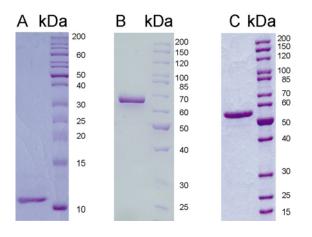


Figure S1. Coomassie stained SDS-PAGE of purified ACP (A, 11.5 kDa), HxcO (B, 66.8 kDa), and HcmO (B, 47.0 kDa).

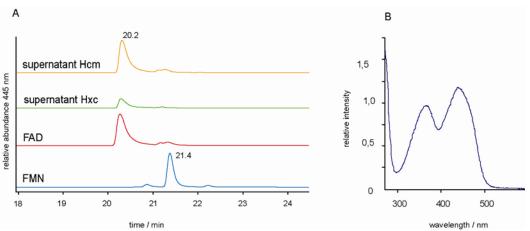


Figure S2. HPLC Analysis of HxcO and HcmO cofactor and UV-visible spectrum of HcmO. The UV-visible spectrum of HcmO highlights the λ_{max} at 377 and 450 nm.

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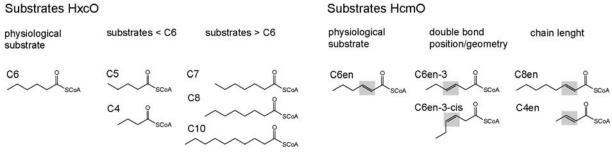


Figure S3. Acyl-CoA substrates used in this study.

| Table S2. MALDI-TOF | analysis of chem | ically synthesized fa | tty acid CoA derivatives. |
|---------------------|------------------|-----------------------|---------------------------|
| | | | |

| FA-CoA | c.m. [M+H] ⁺ | o.m. [M+H] ⁺ |
|-------------------|--------------------------------|--------------------------------|
| C4 | 838.1 | 838.2 |
| C5 | 852.2 | 852.2 |
| C6 | 866.2 | 866.1 |
| C7 | 880.2 | 880.2 |
| C8 | 894.2 | 894.1 |
| C10 | 922.2 | 922.2 |
| C6-EN2 | 864.2 | 864.1 |
| C8-EN2 | 892.2 | 892.2 |
| C4-EN2 (crotonyl) | 836.1 | 836.2 |
| C6-EN3 | 864.2 | 864.2 |
| C6-EN3cis | 864.2 | 864.2 |

Table S3. MALDI-TOF analysis of chemoenzymatically synthesized CDA derivatives.

| | c.m. [M+H] ⁺ | o.m. [M+H] ⁺ |
|-----------------|--------------------------------|--------------------------------|
| Hexanoyl-CDA | 1466.6 | 1466.7 |
| Hex-2-enoyl-CDA | 1464.5 | 1464.7 |

Table S4. [M+H]²⁺ mass fragments of fatty acid-*S*-Ppan-PCP (Lys⁵⁷-Arg⁷¹) subjected to MS².

| Chain length | Saturated fatty acid | Unsaturated fatty acid | Epoxidized fatty acid |
|--------------|----------------------|------------------------|-----------------------|
| C4 | 1032.5 | 1030.5 | 1038.5 |
| C5 | 1039.5 | 1037.5 | 1045.5 |
| C6 | 1046.5 | 1044.5 | 1052.5 |
| C7 | 1053.5 | 1051.5 | 1059.5 |
| C8 | 1060.5 | 1058.5 | 1068.5 |
| C10 | 1074.5 | 1072.5 | 1080.5 |

Direct Ligation of Carrier Protein-Bound Thioesters: A Versatile Method for the Characterization of Fatty Acid Tailoring Enzymes during Lipopeptide Biosynthesis Assays with acyl-CoA substrates

Reaction mixtures for the detection of HxcO oxidation/epoxidation products were prepared using 250 μ M hexanoyl-CoA substrate, 5-50 μ M HxcO and 100 μ M FAD in assay buffer. Reaction mixtures for the detection of HcmO epoxidation product were prepared using 250 μ M hexenoyl-CoA substrate, 10 μ M HxcO, 250 μ M FAD and 250 μ M NAD(P)H in assay buffer. Incubations were carried out at different temperatures and several periods of time.

Assays with chemoenzymatically synthesized CDA substrates

HxcO Assay: The reaction mixture contained 100 μ M hexanoyl-CDA, 20 μ M enzyme and 60 μ M FAD in a total volume of 75 μ L. After incubation at 25 °C for 30 min the reaction was quenched by the addition of 15 μ L formic acid and directly analyzed by LC-ESI-MS.

HcmO Assay: The reaction mixture contained 100 μ M hex-2-enoyl-CDA, 20 μ M enzyme, 60 μ M FAD, 100 μ M NAD(P)H in a total volume of 75 μ L. After incubation at 25 °C for 30 min the reaction was quenched by the addition of 15 μ L formic acid and directly analyzed by LC-ESI-MS.

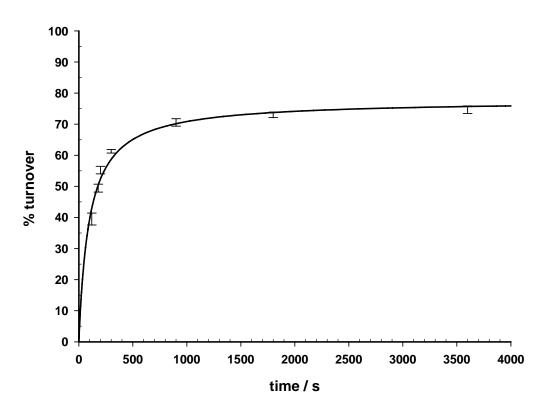


Figure S4. Kinetics of (2R,3S)-2,3-epoxyhexanoic-S-Ppan-ACP formation mediated by HxcO. Samples were trypsinized at different time points following initiation of the enzymatic reactions, and were then fractionated and analyzed by HPLC-MS². The percentage of HxcO reaction product tethered to the ACP is plotted as a function of time.