

Figure 2. FTMS monitoring of the loading of benzoate and salicylate on the ACP EncC. (A) ATP (10 mM)-dependent transfer of benzoic acid (10 mM) by EncN (2 μ M) on *holo*-EncC (20 μ M). (B) Sfp-catalyzed phosphopantetheinylation of *apo*-EncC (20 μ M) by benzoyl-CoA⁷ (~0.25 mM) generated in situ. (C) Same reaction conditions as panel A except that equimolar amounts of benzoic, salicylic, and 2,3-dihydroxybenzoic acids were added. (D) Same reaction conditions as panel A except that benzoic acid was omitted.

12 264.99 Da theor, Δ 0.57 amu), which is +16 Da larger than benzoate (Figure 2D). The HPLC levels of unreacted *holo*-EncC are the same in the benzoate species as in the salicylate species, so not only does EncN activate salicylate, it does so with ease. These ACP ligase data support our prior CoA ligase activity observation that EncN has broad substrate tolerance.⁷ We tested initially various aryl acids as primers in a minimal enterocin PKS enzyme assay harboring the KS $_{\alpha\beta}$ heterodimer EncAB, *holo*-EncC, EncD, EncN, and the *Streptomyces glaucescens* FabD (SgFabD) with the appropriate cofactors and the primer unit malonyl-CoA. The products of this enzymatic transformation when supplemented with the natural primer benzoic acid are wailupemycins D–G,³ which are formed via the spontaneous cyclization of the reduced nascent polyketide. Alternative incubation with *p*-fluorobenzoate and 2- and 3-thiophene carboxylates gave wailupemycin F and G analogues (1–6) (Figure 3), which were identical upon HPLC-MS comparison with authentic standards generated by *in vivo* mutasynthesis.⁷ With the knowledge that alternate starters could successfully be incorporated *in vitro*, the same enzymatic assay conditions were employed to probe whether substrates that were activated *in vitro* by EncN yet failed *in vivo*⁷ could be successfully extended *in vitro*. To our delight, five new sets of wailupemycin F and G derivatives were generated, including those derived from 3- and 4-hydroxybenzoates, 3,4-dihydroxybenzoate, *p*-chlorobenzoate, and nicotinate (7–16) (Figure 3). We had thus successfully synthesized *in vitro* 10 novel unnatural polyketides and a further six unnatural products that we had previously generated *in vivo* via mutasynthesis.

Upon the addition of the FAD-dependent favorskiiase enzyme EncM to the minimal enterocin synthase system, we generated a series of desmethyl-5-deoxyenterocin analogues. To confirm the syntheses of the predicted polyketides, HPLC-MS was again employed to analyze the resulting organic extracts generated from the enzymatic incubations. Desmethyl-5-deoxyenterocin analogues

primed with *p*-fluoro-, *p*-chloro-, and 3- and 4-hydroxybenzoates and 2- and 3-thiophenecarboxylates were detected in the respective organic enzyme assay extracts by LC-MS (17–22) (Figures 3 and 4). The extracts also contained the wailupemycin F and G analogues as expected. Further inspection of the fluoro and chloro products revealed additional analogues most likely corresponding to the C-3-epimers of 17 and 18 (Figure 3), respectively, which are related to the natural product 3-*epi*-5-deoxyenterocin, a minor metabolite of the wild-type bacterium.⁴ Rearranged analogues primed with 3,4-dihydroxybenzoate or nicotinate, on the other hand, were not evident by LC-MS even though the wailupemycin F and G analogues were indeed produced. Further incubation with cyclohex-1-enecarboxylate was unsuccessful, representing the only example where the *in vivo* mutasynthesis incorporation of a precursor⁷ was not mirrored *in vitro*.

We speculate that the inability to generate dihydroxybenzoyl- or nicotinoyl-derived desmethyl-5-deoxyenterocin analogues may be due to the instability of the enzyme-bound linear poly- β -keto intermediate or its incompatibility with the favorskiiase EncM. We observed that the *in vitro* production of polyketides across the board tended to favor the wailupemycin shunt products rather than the rearranged enterocin analogues, which was contrary to that observed with the natural primer benzoate, in which the natural enterocins were the major *in vitro* products.³ This may be attributed to the *in vitro* system being more flexible than the *in vivo* system and the absence of editing enzymes such as the hydrolase EncL (J.A.K., Q.C., B.S.M., unpublished observations).

In conclusion, we successfully directed the one-pot, total enzymatic synthesis of 24 wailupemycin and enterocin analogues, the majority of which are new chemical entities. Although the enterocin benzoate:CoA ligase EncN has broad *in vitro* substrate specificity toward other aryl acids, corresponding mutasynthesis experiments yielded only a small collection of unnatural analogues.⁷

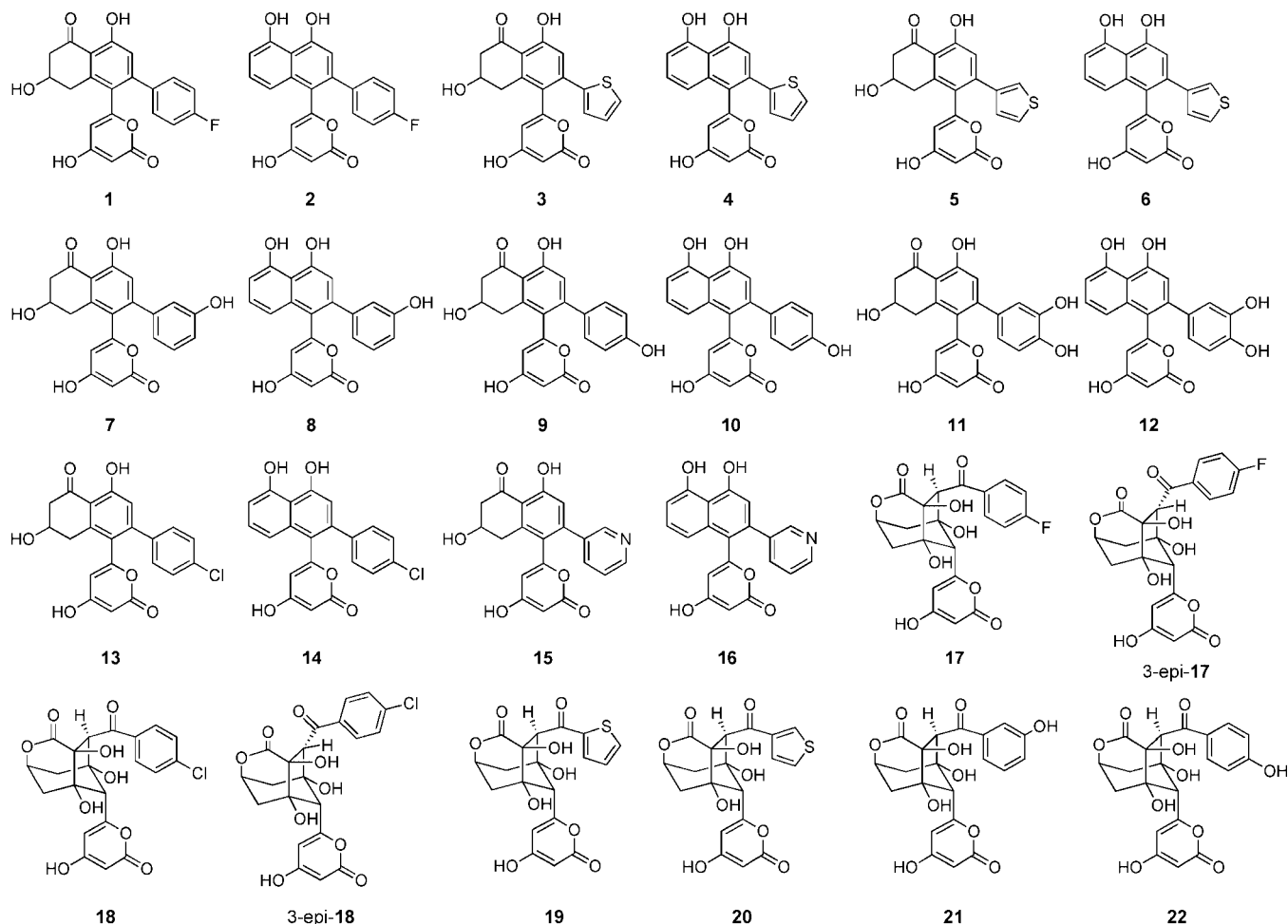


Figure 3. Analogues of wailupemycins F and G, desmethyl-5-deoxyenterocin, and 3-*epi*-desmethyl-5-deoxyenterocin generated *in vitro* using purified proteins.

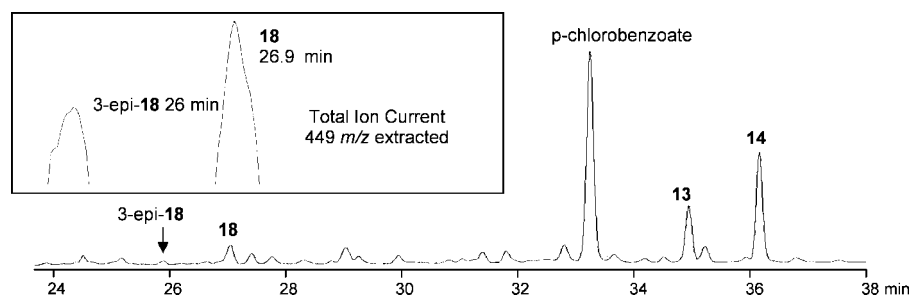


Figure 4. Representative reversed-phase HPLC chromatogram at 340 nm of an organic extract of the EncABCDMN/SgFabD enzymatic incubation of *p*-chlorobenzoate (1 mM) and malonyl-CoA (2.5 mM). Chlorinated products are marked and were analyzed by MS; remaining chromatographic peaks are rather associated with the enzyme/coenzyme mixture.

The *in vivo* experiment highlighted both the strengths and limitations of precursor-directed biosynthesis with mutant bacteria, as unnatural products were indeed produced, yet the biotransformations were not faithful in the majority of cases. Thus, *in vitro* total enzyme reconstitution provides an alternative approach to interrogate the dexterity of a biosynthetic pathway without the competing complications of whole cell transformations involving uptake, metabolism, toxicity, and transport. In the suite of experiments described here with alternative priming molecules, we effectively expanded the functional nature of the enterocin starter unit.

Experimental Section

General Experimental Procedures. Protein purification and enzyme assays were performed as described in ref 3. Extracts were analyzed using a Hewlett-Packard 1100 series HPLC system linked to a Hewlett-

Packard 1100 MSD mass spectrometer using atmospheric pressure ionization and operating in the positive-ion mode.

In Vitro Synthesis of Wailupemycin Analogues. The *in vitro* synthesis of wailupemycin analogues followed the published protocol in ref 3 except for the replacement of benzoic acid with 0.5 mM aryl acids. HPLC-MS analysis verified the production of **1** (33.0 min, m/z [MH⁺] 383.0), **2** (34.5 min, m/z [MH⁺] 365.0), **3** (31.5 min, m/z [MH⁺] 371.0), **4** (33.3 min, m/z [MH⁺] 353.0), **5** (31.3 min, m/z [MH⁺] 371.0), **6** (33.1 min, m/z [MH⁺] 353.0), **7** (28.6 min, m/z [MH⁺] 381.0), **8** (30.8 min, m/z [MH⁺] 363.0), **9** (27.9 min, m/z [MH⁺] 381.0), **10** (30.1 min, m/z [MH⁺] 363.0), **11** (26.4 min, m/z [MH - H₂O⁺] 379.0), **12** (29.4 min, m/z [MH⁺] 379.0), **13** (35.0 min, m/z [MH⁺] 399.0/401.1), **14** (36.2 min, m/z [MH⁺] 381.0/383.0), **15** (21.8 min, m/z [MH⁺] 366.1), and **16** (23.4 min, m/z [MH⁺] 348.0).

In Vitro Synthesis of Enterocin and Wailupemycin Analogues. The addition of *holo*-EncM (2 μM) to the reconstituted minimal

enc PKS preceded the addition of the aryl acid primers, per ref 3, to give **17** (24.0 min, m/z [MH⁺] 433.1), 3-epi-**17** (23.0 min, m/z [MH⁺] 433.1), **18** (26.9 min, m/z [MH⁺] 449.0/451.0), 3-epi-**18** (26.0 min, m/z [MH⁺] 449.0/451.0), **19** (20.2 min, m/z [MH⁺] 421.0), **20** (20.0 min, m/z [MH⁺] 421.0), **21** (19.9 min, m/z [MH⁺] 431.0), and **22** (19.2 min, m/z [MH⁺] 431.0).

Acknowledgment. This research was supported by NIH grants GM067725 to N.L.K. and AI47818 to B.S.M.

References and Notes

- (1) Kopp, F.; Marahiel, M. A. *Curr. Opin. Biotechnol.* **2007**, *18*, 513–520.
- (2) (a) Roessner, R. A.; Scott, A. I. *Chem. Biol.* **1996**, *3*, 325–330. (b) Fecik, R. A. *Nat. Chem. Biol.* **2007**, *9*, 531–532. (c) Sattely, E. S.; Fischbach, M. A.; Walsh, C. T. *Nat. Prod. Rep.* **2008**, *25*, 757–793.
- (3) Cheng, Q.; Xiang, L.; Izumikawa, M.; Meluzzi, D.; Moore, B. S. *Nat. Chem. Biol.* **2007**, *3*, 557–558.
- (4) (a) Sitachitta, N.; Gadepalli, M.; Davidson, B. S. *Tetrahedron* **1996**, *52*, 8073–8080. (b) Piel, J.; Hoang, K.; Moore, B. S. *J. Am. Chem. Soc.* **2000**, *122*, 5415–5416.
- (5) (a) Piel, J.; Hertweck, C.; Shipley, P.; Hunt, D. S.; Newman, M. S.; Moore, B. S. *Chem. Biol.* **2000**, *7*, 943–955. (b) Xiang, L.; Kalaitzis, J. A.; Nilsen, G.; Chen, L.; Moore, B. S. *Org. Lett.* **2002**, *4*, 957–960. (c) Hertweck, C.; Xiang, L.; Kalaitzis, J. A.; Cheng, Q.; Palzer, M.; Moore, B. S. *Chem. Biol.* **2004**, *11*, 461–468. (d) Xiang, L.; Kalaitzis, J. A.; Moore, B. S. *Proc. Natl. Acad. Sci. U.S.A.* **2004**, *101*, 15609–15614. (e) Izumikawa, M.; Cheng, Q.; Moore, B. S. *J. Am. Chem. Soc.* **2006**, *128*, 1428–1429.
- (6) (a) Xiang, L.; Moore, B. S. *J. Biol. Chem.* **2002**, *277*, 32505–32509. (b) Xiang, L.; Moore, B. S. *J. Bacteriol.* **2003**, *185*, 399–404.
- (7) Kalaitzis, J. A.; Izumikawa, M.; Xiang, L.; Hertweck, C.; Moore, B. S. *J. Am. Chem. Soc.* **2003**, *125*, 9290–9291.
- (8) (a) Bumpus, S. B.; Kelleher, N. L. *Curr. Opin. Chem. Biol.* **2008**, *12*, 475–482. (b) Dorrestein, P. C.; Kelleher, N. L. *Nat. Prod. Rep.* **2006**, *23*, 893–918.

NP800598T