

### Communication

# The Biosynthesis of Spinosyn in *Saccharopolyspora spinosa*: Synthesis of the Cross-Bridging Precursor and Identification of the Function of SpnJ

Hak Joong Kim, Rongson Pongdee, Qingquan Wu, Lin Hong, and Hung-wen Liu

J. Am. Chem. Soc., 2007, 129 (47), 14582-14584 • DOI: 10.1021/ja076580i

Downloaded from http://pubs.acs.org on February 9, 2009



## More About This Article

Additional resources and features associated with this article are available within the HTML version:

- Supporting Information
- Links to the 3 articles that cite this article, as of the time of this article download
- Access to high resolution figures
- Links to articles and content related to this article
- Copyright permission to reproduce figures and/or text from this article

View the Full Text HTML





#### Published on Web 11/07/2007

# The Biosynthesis of Spinosyn in *Saccharopolyspora spinosa*: Synthesis of the Cross-Bridging Precursor and Identification of the Function of SpnJ

Hak Joong Kim, Rongson Pongdee, Qingquan Wu, Lin Hong, and Hung-wen Liu\*

Division of Medicinal Chemistry, College of Pharmacy, and Department of Chemistry and Biochemistry, University of Texas at Austin, Austin, Texas 78712

Received September 6, 2007; E-mail: h.w.liu@mail.utexas.edu

Spinosyns A (1) and D (2) are polyketide-derived macrolides produced by *Saccharopolyspora spinosa*.<sup>1</sup> The combination of 1 and 2 serves as the active ingredient in a commercial insecticide, Naturalyte. This mixture exhibits excellent insecticidal activity with low mammalian toxicity and little detrimental environmental effects.<sup>2</sup> Structurally, the spinosyns consist of a 22-membered macrolactone ring fused to a perhydro-*as*-indacene core scaffold. In addition, they are glycosylated with tri-*O*-methylrhamnose and forosamine at C-9 and C-17, respectively. The aglycone portion of spinosyns is unusual among polyketide-derived secondary metabolites due to the presence of three intramolecular carbon–carbon bonds that constitute the *as*-indacene skeleton. The unusual nature of the spinosyn aglycone suggests an intriguing biosynthetic pathway which has generated much recent attention.<sup>3,4</sup>

The spinosyn (*spn*) biosynthetic gene cluster was cloned from *S. spinosa*.<sup>3</sup> Sequence analysis coupled with gene disruption experiments suggested that macrolactone **4**, which is derived from the linear polyketide precursor **3**, is a likely intermediate to form the *as*-indacene aglycone (**9**),<sup>3,4</sup> and the proteins encoded by *spnF*, *spnJ*, *spnL*, and *spnM* could be involved in the cross-bridging modifications. SpnJ exhibits sequence homology to a flavin-dependent dehydrogenase (55% identity, 68% similarity) from *Streptomyces steffisburgensis*,<sup>5</sup> and the SpnF, SpnL, and SpnM proteins show resemblance to methyltransferase (SpnF and SpnL) and lipase (SpnM),<sup>3</sup> but lack part of the feature residues/motifs conserved among members of each enzyme family. Hence, their biological functions are not apparent and their roles in the cross-bridging reactions cannot be established with certainty.

If **4** is indeed a precursor of the polycyclic aglycone and the cross-bridging reactions are all post-cyclization events, oxidation of the 15-hydroxyl group of **4** to a keto group  $(\mathbf{4} \rightarrow \mathbf{5})$  and deprotonation at C-14 to form the C-3/C-14 linkage  $(\mathbf{5} \rightarrow \mathbf{6} \rightarrow \mathbf{7})$  likely occur first (Scheme 1, route A). Subsequent deprotonation at C-14 of **7** and elimination of the 11-hydroxyl group to give **8** is expected to trigger the second cross-coupling reaction  $(\mathbf{8} \rightarrow \mathbf{9})$  to produce the *as*-indacene core. However, it is also possible that the initial oxidation of the 15-hydroxyl group takes place on the linear precursor **3** when it is still linked to the acyl carrier protein (ACP) associated with the polyketide synthase SpnE (Scheme 1, route B). The keto product **10** is next cleaved from ACP catalyzed by a thioesterase (TE) to give **5**, which then undergoes sequential cyclizations in a similar manner as described above. In this pathway, compound **4** is bypassed.

Although a [4 + 2] cyclization reaction has been speculated to be a key step in the biosynthesis of many natural products, few have been investigated at the molecular level.<sup>6</sup> In the case of the polyketide-derived lovastatin (14, Scheme 2), which contains an octahydronaphthalene core, in vitro study with purified polyketide synthase (PKS) showed that it is capable of converting a linear *N*-acetylcysteamine mimic of the appropriate enzyme-bound



polyketide intermediate (12) to the correct isomer of the octahydronaphthalene moiety (13).<sup>6b,f,g</sup> By analogy to 14, the crossbridging reactions in the spinosyn case could also take place when the polyketide intermediate 10 is still tethered to ACP. In this scenario, C-15 oxidation should occur on the polyketide intermedi-

Scheme 4



ate 3 to give 10, which would then undergo cyclization to form the tricyclic polyketide 11 (Scheme 1, route C). In this route, the intermediacies of 4-8 are no longer necessary.

Intrigued by the complexity of the tetracyclic nucleus of the spinosyns, especially the possible involvement of a Diels-Alder-type [4 + 2] cyclization  $(7 \rightarrow 8 \text{ or } 10 \rightarrow 11)$ , we decided to study the proposed cross-bridging reactions by first focusing on the initial oxidation step to distinguish the three possible pathways. Herein, we report the expression and purification of SpnJ, the chemical synthesis of 3 and 4, and the results showing that 4, but not 3, is a substrate for SpnJ. Our results clearly demonstrate that SpnJ is the predicted oxidase and is specific for C-15 oxidation of macrolactone 4. These findings provide compelling evidence establishing the early sequence of events in the cross-bridging reactions.

Outlined in Scheme 3 is our retrosynthetic analysis for the preparation of **3** and **4**, in which two main couplings were envisioned: a Julia-Kocienski olefination of sulfone **15** with aldehyde **16** to establish the *E*-alkene geometry at C-12/C-13, and a palladium-mediated Heck reaction to connect the C-5/C-6 carbon–carbon bond with **17**. Once the linear polyketide **3** is constructed, a Yamaguchi macrolactonization to form the ester linkage at C-1 will afford **4**.<sup>7</sup>

Our synthesis of the aryl sulfone 15, shown in Scheme 4, began with the Weinreb amide 18 derived from  $\delta$ -valerolactone.<sup>8</sup> Asymmetric addition<sup>9</sup> of diethyl zinc catalyzed by (-)-(1S,2R)-N,Ndibutylnorephedrine [(-)-DBNE] provided **19**, whose enantiomeric purity (92% ee) and absolute configuration were determined by NMR analysis of the corresponding (R)- and (S)-Mosher esters. Subsequent silvlation followed by reduction furnished 20. Aldol reaction<sup>10</sup> between **20** and Crimmins' oxazolidinethione auxiliary 21 gave the syn-aldol adduct 22. The configurations at C-16/C-17 were determined by application of the Zimmerman-Traxler transition state model as well as analysis of the  $J_{16,17}$  values, which are in accord with literature precedent.<sup>11</sup> Compound 23, derived from 22 via silvlation and reduction, was subjected to Ley oxidation and then Brown's (+)-Ipc<sub>2</sub>B(allyl) reagent<sup>12</sup> to yield 24 as a single stereoisomer. The relative configurations at C-15/C-17 were determined by conversion of 24 to the corresponding 1,3-acetonide followed by application of Rychnovsky's <sup>13</sup>C NMR method,<sup>13</sup>



providing proof for a 1,3-*syn* relationship. Silyl ether protection followed by a two-step Lemieux–Johnson oxidation of the terminal olefin and reduction of the resulting aldehyde gave **26**. It was crucial to conduct the oxidation of the terminal alkene under neutral pH as a nonbuffered reaction medium resulted in unwanted deprotection of the triethylsilyl ether at C-15 which required reprotection. Completion of the requisite sulfone **15** was achieved by conversion of **26** to aryl sulfide **27** under Mitsunobu conditions followed by oxidation employing hydrogen peroxide and ammonium molyb-date.<sup>14</sup>

Our synthesis of aldehyde 16 is depicted in Scheme 5. Asymmetric Brown allylation of dialdehyde 28, obtained from Dmannitol,<sup>15</sup> followed by benzoylation of the newly generated secondary alcohols and removal of the isopropylidene protecting group under acidic hydrolysis afforded diol 29 in excellent overall yield. Oxidative cleavage of the vicinal diol in 29 facilitated by lead tetraacetate followed by sodium borohydride reduction of the ensuing aldehyde afforded primary alcohol 30. Next, protection of the alcohol functionality in 30 as its PMB ether followed by exchange of the benzoate protecting groups for silyl ethers provided **31**. Following an oxidation of the terminal olefin in **31** using Jin's protocol,16 the resulting aldehyde was subjected to Takai olefination,<sup>17</sup> affording the desired (E)-vinyliodide **32** in good overall yield as a single stereoisomer. Oxidative removal of the PMB ether employing DDQ in aqueous dichloromethane and oxidation of the liberated hydroxy group with Dess-Martin periodinane then provided aldehyde 16.

With fragments 15 and 16 in hand, we assembled the presumed biosynthetic precursors 3 and 4 (Scheme 6). The use of KHMDS in THF provided the best yield and stereoselectivity in forming the C(12)-C(13) olefin by the Julia-Kocienski olefination<sup>18</sup> protocol. Completion of the triene moiety was accomplished by a Heck reaction involving 33 with ethyl dienoate 17<sup>19</sup> under ligand-free conditions,<sup>20</sup> affording **34** as a mixture of isomers (E/Z = 6/1 at C(4)-C(5)) that were separated by flash column chromatography. Chemoselective deprotection of the TES ether utilizing PPTS in ethanol followed by saponification furnished seco-acid 35. Acid 35 was then subject to Yamaguchi macrolactonization<sup>21</sup> followed by global deprotection of the silicon-based protecting groups employing HF•Pyr in cold ethanol for 4 days to give 4 in moderate yields.<sup>22</sup> The preparation of the linear substrate 3 was achieved in a similar manner. Due to the difficulty of preparing the ACP-bound linear polyketide chain, the intended linear substrate 3 was synthesized as the N-acetylcysteamine (NAC) thioester 3'. Since the corresponding N-acetylcysteamine (NAC)-linked thioesters have been commonly used in the biosynthetic studies as surrogates for ACP-bound acyl esters, 3' is expected to be an effective substitute for 3. As shown in Scheme 6, saponification of 34 afforded acid 36, which was then coupled with N-acetylcysteamine under Yamaguchi esterification conditions. Final deprotection using HF•Pyr complex led to the desired linear substrate 3'.22

Scheme 6



With sufficient quantities of 3' and 4 now available, we evaluated whether the linear polyketide thioester 3' or the macrolactone 4 is a substrate for SpnJ, which is predicted to be an oxidase. Since the functions of SpnF, SpnL, and SpnM are not obvious, it would be interesting to examine whether SpnJ is a multifunctional enzyme capable of catalyzing not only oxidation at C-15 but also deprotonation at C-14 to initiate the Michael addition to form a C-C bond between C-3 and C-14, or the 1,4-elimination of the 11-OH group setting the stage for a possible Diels-Alder-type [4 + 2]cycloaddition.

To characterize SpnJ, the spnJ gene was cloned into a pET28b-(+) vector and the resulting construct was expressed in E. coli BL21(DE3) cells.<sup>22</sup> After induction under 25 mM isopropyl  $\beta$ -Dthiogalactopyranoside (IPTG), the N-terminal His<sub>6</sub>-tagged SpnJ protein (60.6 kDa) was purified to near homogeneity using Ni-NTA resin.<sup>22</sup> The absorption spectrum of the purified protein is typical for that of a flavoprotein. To determine the function of SpnJ, SpnJ (20  $\mu$ M) was incubated with 2 mM 3' and 40  $\mu$ M FAD in Tris-HCl buffer (pH 8.0) at 29 °C. No new product was detected after overnight incubation. Because 3' cannot be processed by SpnJ, it is reasonable to conclude that 3 is not a substrate for SpnJ.

In contrast, TLC and reverse-phase HPLC analysis of the incubation mixture containing SpnJ (20  $\mu$ M), 2 mM 4, and 40  $\mu$ M FAD in Tris-HCl buffer (pH 8.0) revealed the time-dependent formation of a new product (nearly quantitative conversion after 4 h at 29 °C, see Figure 1), which was identified by <sup>1</sup>H and <sup>13</sup>C NMR spectroscopy as well as HR-CIMS analysis as the C-15 ketone derivative 5.22 These results clearly established that SpnJ is a flavin oxidase and 4 is the substrate for SpnJ. The fact that no trace of cross-bridging products was detected indicated that SpnJ is only a monofunctional oxidase. These results provide strong support for the conversion of 4 to 5 as the first step in the post-cyclization reactions (vide supra) and set the stage for the study of the putative Diels-Alderase activity.

In summary, the experiments described herein detail a convergent synthesis of 4 postulated to be involved in spinosyn biosynthesis. More importantly, we have validated the catalytic function of SpnJ and established 4 as the precursor of the tricyclic nucleus of spinosyns. Future work will encompass characterization of the



Figure 1. (1) TLC and (2) HPLC analysis of incubation mixture without SpnJ (A) or with SpnJ (B)<sup>21</sup> (TLC: silica gel,  $CH_2Cl_2/MeOH = 93:7$ ; HPLC: C-18, 2% NH<sub>4</sub>OAc/CH<sub>3</sub>CN = 70:30 to 20:80 over 120 min).

remaining enzymes (SpnF, SpnL, and SpnM) believed to be involved in the cross-bridging event as well as their biological roles leading to an understanding of the mechanism of formation for the spinosyn aglycone.

Acknowledgment. This work was supported by grants from the National Institutes of Health (GM35906 and GM54346).

Supporting Information Available: Experimental procedures (PDF). This material is available free of charge via the Internet at http:// pubs.acs.org.

#### References

- Kirst, H. A.; Michel, K. H.; Martin, J. W.; Creemer, L. C.; Chio, E. H.; Yao, R. C.; Nakatsukasa, W. M.; Boeek, L. D.; Occolowitz, J. L.; Paschal, J. W.; Deeter, J. B.; Jones, N. D.; Thompson, G. D. Tetrahedron Lett. 1991, 32, 4839–4842.
- (a) Sparks, T. C.; Crouse, G. D.; Durst, G. Pest. Manag. Sci. 2001, 57, (b) Spins, (b) Salgado, V. L.; Sparks, T. C. In *Comprehensive Molecular Insect Science*; Iatrou, K., Gill, S. S., Gilbert, L. I., Eds.; Pergamon: Oxford, 2005; Vol. 6, pp 137–173.
- Waldron, C.; Matsushima, P.; Rosteck, P. R., Jr.; Broughton, M. C.; Turner, J.; Madduri, K.; Crawford, K. P.; Merlo, D. J.; Baltz, R. H. Chem. Biol. 2001, 8, 487-499.
- Martin, C. J.; Timoney, M. C.; Sheridan, R. M.; Kendrew, S. G.; Wilkinson, B.; Staunton, J. C.; Leadlay, P. F. Org. Biomol. Chem. 2003, (4)1. 4144-4147
- (5) Gullon, S.; Olano, C.; Abdelfattah, M. S.; Brana, A. F.; Rohr, J.; Mendez, C.; Salas, J. A. *Appl. Environ. Microbiol.* **2006**, *72*, 4172–4183. (6) (a) Solanapyrone Synthase: Katayama, K.; Kobayashi, T.; Oikawa, H.;
- Honma, M.; Ichihara, A. Biochim. Biophys. Acta 1998, 1384, 387-395. (b) Lovastatin biosynthesis: Auclair, K.; Sutherland, A.; Kennedy, J.; Witter, D. J.; Van den Heever, J. P.; Hutchinson, C. R.; Vederas, J. C. J. *Am. Chem. Soc.* **2000**, *122*, 11519–11520. Macrophomate synthase: (c) Ose, T.; Watanabe, K.; Mie, T.; Honma, M.; Watanabe, H.; Yao, M.; Ose, I., Watahabe, K., Mie, T., Hollina, M., Watahabe, H., Tao, M.,
   Oikawa, H.; Tanaka, I. *Nature* 2003, *422*, 185–189. (d) Guimaraes, C.
   R.; Udier-Blagovic, M.; Jorgensen, W. L. J. Am. Chem. Soc. 2005, *127*, 3577–3588. (e) Serafimov, J. M.; Westfeld, T.; Meier, B. H.; Hilvert, D. J. Am. Chem. Soc. 2007, *129*, 9580–9581. (f) Review: Oikawa, H. Bull.
   Chem. Soc. Jpn. 2005, *78*, 537–554. (g) Stocking, E. M.; Williams, R. M. Angew. Chem., Int. Ed. 2003, 42, 3078-3115.
- The configurations at C-9, -11, -15,<sup>4</sup> and -17 were assigned based on the fact that the four ketoreductase (KR) domains in the *spn* PKS assembly are all type A KRs (Caffrey, P. *ChemBioChem* **2003**, 4, 654–657).<sup>22</sup>
- (8) Molander, G. A.; McWilliams, J. C.; Noll, B. C. J. Am. Chem. Soc. 1997, Soai, K.; Yokoyama, S.; Hayasaka, T.; Ebihara, K. Chem. Lett. 1988, 843-846. (9)
- (10) Crimmins, M. T.; King, B. W.; Tabet, E. A.; Chaudhary, K. J. Org. Chem.
- 2001, 66, 894-902. (11) Heathcock, C. H.; Pirrung, M. C.; Sohn, J. E. J. Org. Chem. 1979, 44,
- 4294-4299 (12) Brown, H. C.; Randad, R. S.; Bhat, K. S.; Zaidlewicz, M.; Racherla, U. S. J. Am. Chem. Soc. **1990**, *112*, 2389–2392.
- (13) Rychnovsky, S. D.; Rogers, B. N.; Richardson, T. I. Acc. Chem. Res. **1998**, *31*, 9–17.
- (14) Paquette, L. A.; Chang, S. K. Org. Lett. 2005, 7, 3111-3114.
- (15) Saniere, M.; Lemerrer, Y.; Barbe, B.; Koscielniak, T.; Depezay, J. C. *Tetrahedron* **1989**, 45, 7317–28.
- (16) Yu, W.; Mei, Y.; Kang, Y.; Hua, Z.; Jin, Z. Org. Lett. 2004, 6, 3217-3219
- (17) Evans, D. A.; Black, W. C. J. Am. Chem. Soc. 1993, 115, 4497-4513. (18) Blakemore, P. R.; Cole, W. J.; Kocienski, P. J.; Morley, A. Synlett 1998,
- 26 28
- (19) Rodriguez, J.; Waegell, B. Synthesis 1988, 534-535.
- (20) Jeffery, T. *Tetrahedron Lett.* 1985, 26, 2667–2670.
  (21) Inanaga, J.; Hirata, K.; Saeki, H.; Katsuki, T.; Yamaguchi, M. *Bull. Chem. Soc. Jpn.* 1979, 52, 1989–1993.
- (22)See Supporting Information for spectral data and experimental details. JA076580I