

The in Vitro Characterization of the Iterative Avermectin Glycosyltransferase AveBI Reveals Reaction Reversibility and Sugar Nucleotide Flexibility

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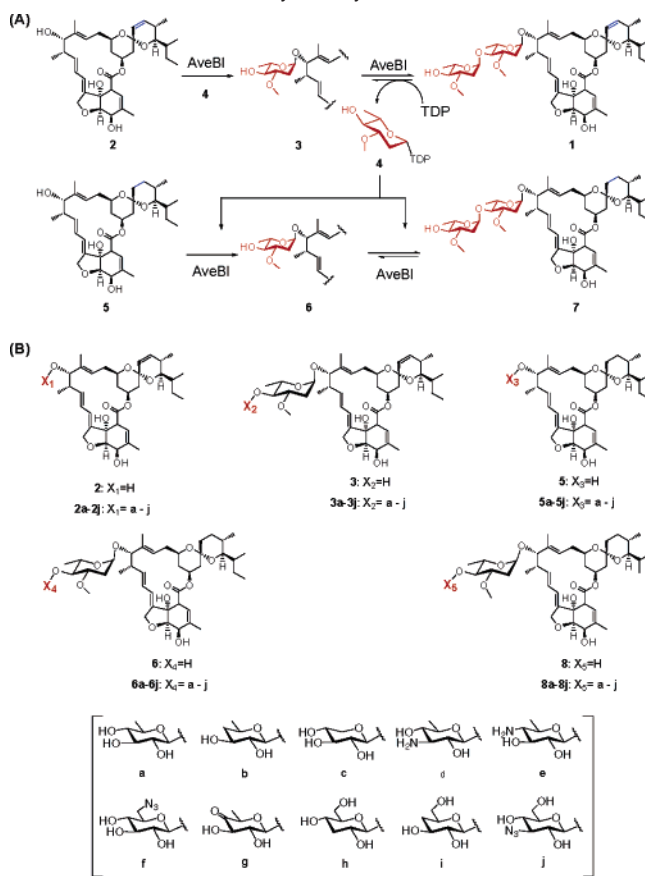
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Avermectins (AVMs, e.g., Scheme 1, **1**) are 16-membered macrocyclic lactones produced by *Streptomyces avermitilis*. The avermectins, and the related C₂₂–C₂₃-reduced ivermectin (IVM, e.g., Scheme 1, **7**), target the γ -aminobutyric acid (GABA)-related chloride ion channels unique to nematodes, insects, ticks, and arachnids, with little or no mammalian toxicity.¹ The widespread commercial use of these remarkable anthelmintic agents began ~25 years ago as veterinary antiparasitic agents and has more recently expanded to clinical applications for the control of onchocerciasis, stongyloidiasis, and lymphatic filariasis. From a biosynthetic perspective, the AVMs are one of only a few known natural products postulated to derive from iterative glycosylation.² Specifically, a single glycosyltransferase (GT) is required for the attachment of the AVM oleandrosyl-disaccharide (AveBI), proposed to proceed in a stepwise, tandem manner (Scheme 1A). Evidence in support of iterative glycosylation includes the existence of a single glycosyltransferase gene (*aveBI*) within the AVM gene locus,³ *S. avermitilis* crude extract studies suggestive of TDP-oleandrose (Scheme 1A, **4**) as an immediate precursor to the AVM oleandrose moiety,⁴ and the production of a variety of glycosylated AVMs via in vivo pathway engineering.⁵ Herein we describe, the first definitive in vitro biochemical verification of AveBI-catalyzed tandem glycosylation. Furthermore, consistent with the recent illumination of the reversibility of natural product GT-catalyzed reactions,⁶ this study reveals the AveBI-catalyzed reaction to also be reversible, the utility of which is demonstrated by generating 50 AVM variants.

The *aveBI* gene was amplified from pWHM473⁵ and assessed in several expression systems. However, the functional expression of *aveBI* was only achieved in *S. lividans* TK64 by the use of expression vectors pPWW49 and pPWW50.⁷ The N-His₆-AveBI fusion was subsequently purified to greater than 90% purity from this recombinant strain via affinity purification (Figure S1)⁸ and used directly for these studies. Aglycons **2**, **3**, **5**, **6**, and **8** (Scheme 1B) were prepared for this study via selective acid-mediated hydrolysis of AVM B1a (**1**) and IVM (**7**).⁹ Given the lack of availability of TDP- β -L-oleandrose,¹⁰ we opted to first examine the reversibility of the AveBI reaction using commercially available **1** and TDP based upon the recent precedent of the reversibility of natural product glycosyltransferase-catalyzed reactions.⁶

RP-HPLC analysis of an in vitro assay containing 100 μ M **1**, 2 mM TDP, and 12 μ M AveBI revealed the formation of **3** from **1** (30%, Scheme 1A and Figure S2), while **1** remained unchanged in control assays lacking either TDP or AveBI.^{11a} This notable requirement of TDP is inconsistent with a hydrolytic event to provide free sugar and alternatively implicates a corresponding AveBI-catalyzed production of TDP- β -L-oleandrose (**4**). To confirm this hypothesis and assess whether AveBI was capable of catalyzing an “aglycone exchange” reaction,⁶ a reaction containing 100 μ M **1**, 100 μ M **5**, 2 mM TDP, and 12 μ M AveBI was subsequently analyzed. Examination of this reaction revealed the production of

Scheme 1. (A) Tandem Sugar Assembly by AveBI-Catalyzed Aglycone-Exchange Reaction; (B) Library of AVM Analogues Constructed via AveBI-Catalyzed Glycorandomization



3 (63%) from **1** and the subsequent transfer of oleandrose to **5**, to provide **6** (28%) and trace amounts of **7** (7%) (Scheme 1A and Figure S2).^{11b} Consistent with recent studies,⁶ this facile TDP-dependent aglycone exchange supports the in situ intermediacy of TDP- β -L-oleandrose (**4**). Cumulatively, these studies unequivocally establish AveBI as the GT responsible for the stepwise tandem assembly of the AVM oleandrosyl disaccharide and reveal the AveBI-catalyzed reaction to be readily reversible and amenable to aglycone exchange strategies.⁶

The AveBI sugar nucleotide specificity was subsequently probed with 22 NDP-sugars (generated chemically or chemoenzymatically, Figure S3, Supporting Information).¹² As a representative example,¹³ IVM aglycone (**5**) with TDP-6-deoxy- α -D-glucose led to a new product (99% conversion, Figure 1A), the LC–MS of which was consistent with the anticipated product **5a** (Scheme 1B). Substitution of TDP-6-deoxy- α -D-glucose with UDP-6-deoxy- α -D-glucose in the same assay gave **5a** in only 10% yield, indicating a preference for

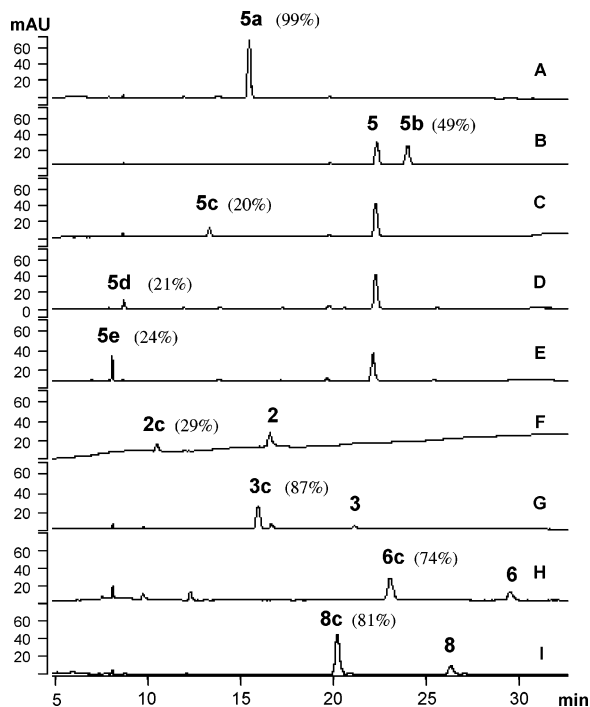


Figure 1. RP-HPLC analysis of representative AveBI reactions. Panels A–E depicted the formation of glycosides **5a**–**5e** in AveBI reactions with **5** as an acceptor. Panels F–I represented the attachment of xylose to aglycons **2**, **3**, **5**, and **8** to form **2c**, **3c**, **5c**, and **8c** by AveBI, respectively. Conversion rates for each reaction were indicated in parentheses. Assay and HPLC conditions are available in Supporting Information.

TDP-sugars. Further AveBI–IVM assays revealed that nine additional TDP-sugar substrates were converted to their corresponding IVM glycosides **5b**–**5j** (Scheme 1B). In a similar fashion, the same set sugars were transferred to aglycons **2**, **3**, **6**, and **8**, producing glycosides **2a**–**2j**, **3a**–**3j**, **6a**–**6j**, and **8a**–**8j** (Scheme 1, Figure 1), respectively. The conversion rates for **a**–**e** glycosides ranged from 18% to 99% while only trace production (1–10%) of **f**–**j** glycosides was observed, with the exception of **6h** (25%) and **6g** (19%). All products were confirmed by LC–MS (Supporting Information, Tables S1 and S2), and controls lacking AveBI or sugar nucleotide gave no reaction. Consistent with the previous *in vivo* studies,⁵ tandem additions of D-configured sugars to aglycone **2** and **5**, or trisaccharide AVM derivatives, were not observed in this study. While this study suggests AveBI to be particularly tolerant of C-6 and/or C-4 sugar modifications, the attachment of unnatural sugar appendages appears to inhibit subsequent disaccharide formation.

In summary, this study is noteworthy for a number of reasons. First, this work provides direct biochemical evidence of the AveBI-catalyzed tandem sugar addition within AVM biosynthesis. Second, this study greatly extends the repertoire of known AveBI D-sugar nucleotide substrates and provides a rapid one-pot strategy for the generation of 50 differentially glycosylated AVMs. Third, in contrast to the *in vitro* macrolide GT studies to date,¹⁴ this study reveals AveBI does not require an “auxiliary/activator” protein for activity. Finally, this study demonstrates the recently established “sugar/aglycone exchange” strategies, based upon the reversibility of GT-catalyzed reactions,⁶ are also applicable to macrolides.

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Supporting Information Available: Experimental procedures and compound characterization data. This material is available free of charge via the Internet at <http://pubs.acs.org>.

References

- (1) For recent reviews see: (a) Geary, T. G. *Trends Parasitol.* **2005**, *21*, 530–532. (b) Omura, S.; Crump, A. *Nat. Rev. Microbiol.* **2004**, *2*, 984–989. (c) Dourmishev, A. L.; Dourmishev, L. A.; Schwartz, R. A. *Int. J. Dermatol.* **2005**, *44*, 981–988. (d) Ikeda, H.; Omura, S. *Chem. Rev.* **1997**, *97*, 2591–2610. (e) Yoon, Y. J.; Kim, E. S.; Hwang, Y. S.; Choi, C. Y. *Appl. Microbiol. Biotechnol.* **2004**, *63*, 626–634.
- (2) Some other potential examples include members of the angucycline, aureolic acid and orthosomycin families (e.g., see Luzhetskyy, A.; Fedoryshyn, M.; Durr, C.; Taguchi, T.; Novikov, V.; Bechtold, A. *Chem. Biol.* **2005**, *12*, 725–729.).
- (3) Ikeda, H.; Nonomiya, T.; Usami, M.; Ohta, T.; Omura, S. *Proc. Natl. Acad. Sci. U.S.A.* **1999**, *96*, 9509–9514.
- (4) Schulman, M. D.; Acton, S. L.; Valentino, D. L.; Arison, B. H. *J. Biol. Chem.* **1990**, *265*, 16965–16970.
- (5) (a) Wohler, S.; Lomovskaya, N.; Kulowski, K.; Fonstein, L.; Occi, J. L.; Gewain, K. M.; MacNeil, D. J.; Hutchinson, C. R. *Chem. Biol.* **2001**, *8*, 681–700. (b) Liao, J.; Lomovskaya, N.; Fonstein, L.; Wohler, S.; Hutchinson, C. R.; Thorson, J. S. Unpublished.
- (6) (a) Minami, K.; Kakinuma, K.; Eguchi, T. *Tetrahedron Lett.* **2005**, *46*, 6187. (b) Zhang, C.; Griffith, B. R.; Fu, Q.; Albermann, C.; Fu, X.; Lee, I.-K.; Li, L.; Thorson, J. S. *Science* **2006**, *313*, 1291.
- (7) (a) Doumith, M.; Weingarten, P.; Wehmeier, U. F.; Salah-Bey, K.; Benhamou, B.; Capdevila, C.; Michel, J. M.; Piepersberg, W.; Raynal, M. C. *Mol. Gen. Genet.* **2000**, *264*, 477–485. (b) Zhang, C. S.; Stratmann, A.; Block, O.; Bruckner, R.; Podeschwa, M.; Altenbach, H. J.; Wehmeier, U. F.; Piepersberg, W. *J. Biol. Chem.* **2002**, *277*, 22853–22862.
- (8) The sequence-confirmed AveBI PCR product was inserted into vector pPWW50 to give expression plasmid pCAM4.10, which was introduced into *Streptomyces lividans* TK64. The cells expressing N-(His)₆-AveBI were resuspended in 30 ml of buffer A (20 mM NaH₂PO₄, pH 7.5, 500 mM NaCl, 10 mM imidazole) supplemented with 1 mg/ml of lysozyme. Cells were lysed by three rounds of French-press (1200 psi). The supernatant was loaded onto a HisTrap HT column (1 ml) and the N-(His)₆-tagged AveBI eluted with a linear gradient of imidazole (10–500 mM) in buffer A via FPLC. After desalting through PD-10 column the purified AveBI was stored in the buffer containing 25 mM Tris-HCl (pH 8.0), 100 mM NaCl, and 10 % glycerol. This purified recombinant AveBI was 50–80% active in the absence of exogenous Mg²⁺.
- (9) (a) Wei, G. H.; Du, Y. G.; Linhardt, R. J. *Tetrahedron Lett.* **2004**, *45*, 6895–6898. (b) Wei, L.; Wei, G.; Zhang, H.; Wang, P. G.; Du, Y. *Carbohydr. Res.* **2005**, *340*, 1583–1590.
- (10) The synthesis of TDP- β -L-oleandrose has not been reported, and the chemoenzymatic synthesis of the closest related sugar nucleotide (TDP- β -L-olivose, which lacks the sugar 3'-OMe) required six linear steps with an overall reported yield of 20% (Amann, S.; Drager, G.; Rupprath, C.; Kirschning, A.; Elling, L. *Carbohydr. Res.* **2001**, *335*, 23–32.).
- (11) (a) Generally, AveBI assays were performed in a total volume of 100 μ L in Tris-HCl buffer (50 mM, pH 8.0) containing 2 mM MgCl₂. Reversibility of AveBI reaction was assayed by co-incubation of 100 μ M avermectin B1a (**1**) or ivermectin (**7**) and 2 mM TDP with 12 μ M AveBI at 30 °C overnight. (b) The AveBI-catalyzed aglycone exchange reaction was assayed by co-incubation of 100 μ M **1**, 100 μ M **5**, and 2 mM TDP with 12 μ M AveBI at 30 °C overnight.
- (12) The sugar nucleotides for this study were generated as previously described. (a) Barton, W. A.; Biggins, J. B.; Jiang, J.; Thorson, J. S.; Nikolov, D. B. *Proc. Natl. Acad. Sci. U.S.A.* **2002**, *99*, 13397–13402. (b) Jiang, J.; Biggins, J. B.; Thorson, J. S. *Angew. Chem., Int. Ed.* **2001**, *40*, 1502–1505. (c) Barton, W. A.; Lesniak, J.; Biggins, J. B.; Jeffrey, P. D.; Jiang, J.; Rajashankar, K. R.; Thorson, J. S.; Nikolov, D. B. *Nat. Struct. Biol.* **2001**, *8*, 545–551. (d) Jiang, J.; Biggins, J. B.; Thorson, J. S. *J. Am. Chem. Soc.* **2000**, *122*, 6803–6804. (e) Albermann, C.; Soriano, A.; Jiang, J.; Vollmer, H.; Biggins, J. B.; Barton, W. A.; Lesniak, J.; Nikolov, D. B.; Thorson, J. S. *Org. Lett.* **2003**, *5*, 933–936. (f) Fu, X.; Albermann, C.; Jiang, J.; Liao, J.; Zhang, C.; Thorson, J. S. *Nat. Biotechnol.* **2003**, *21*, 1467–1469. (g) Borisova, S. A.; Zhang, C.; Takahashi, H.; Zhang, H.; Wong, A. W.; Thorson, J. S.; Liu, H. W. *Angew. Chem., Int. Ed.* **2006**, *45*, 2748–2753.
- (13) A typical reaction to assess sugar nucleotide specificity contained 50 μ M aglycon (**1**–**3**, **5**–**8**), approximately 300 μ M TDP-sugar and 12 μ M AveBI incubated at 30 °C overnight.
- (14) (a) Borisova, S. A.; Zhao, L.; Melancon, I. C.; Kao, C. L.; Liu, H. W. *J. Am. Chem. Soc.* **2004**, *126*, 6534–6535. (b) Yuan, Y.; Chung, H. S.; Leimkuhler, C.; Walsh, C. T.; Kahne, D.; Walker, S. *J. Am. Chem. Soc.* **2005**, *127*, 14128–14129.

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