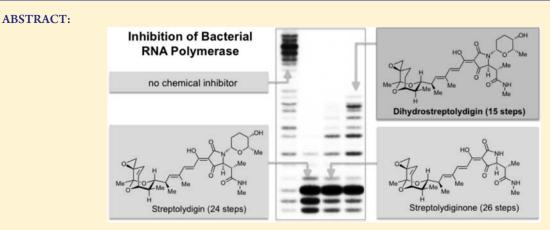


## **Chemical Synthesis Enables Biochemical and Antibacterial Evaluation** of Streptolydigin Antibiotics

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## Supporting Information



Inhibition of bacterial transcription represents an effective and clinically validated anti-infective chemotherapeutic strategy. We describe the evolution of our approach to the streptolydigin class of antibiotics that target bacterial RNA polymerases (RNAPs). This effort resulted in the synthesis and biological evaluation of streptolydigin, streptolydiginone, streptolic acid, and a series of new streptolydigin-based agents. Subsequent biochemical evaluation of RNAP inhibition demonstrated that the presence of both streptolic acid and tetramic acid subunits was required for activity of this class of antibiotics. In addition, we identified 10,11dihydrostreptolydigin as a new RNAP-targeting agent, which was assembled with high synthetic efficiency of 15 steps in the longest linear sequence. Dihydrostreptolydigin inhibited three representative bacterial RNAPs and displayed in vitro antibacterial activity against S. salivarius. The overall increase in synthetic efficiency combined with substantial antibacterial activity of this fully synthetic antibiotic demonstrates the power of organic synthesis in enabling design and comprehensive in vitro pharmacological evaluation of new chemical agents that target bacterial transcription.

## ■ INTRODUCTION

Cellular RNA polymerase (RNAP) is a multisubunit enzyme that catalyzes the synthesis of RNA from the corresponding DNA template. Despite overall structural and functional similarities, bacterial RNAP sequences are substantially different from those of the corresponding eukaryotic enzymes, which explains why rifamycin antibiotics can be used to selectively block bacterial RNAPs, while having no effect on eukaryotic RNAPs.<sup>1</sup> As a result of potent RNAP inhibition, rifamycins display broad-spectrum antibacterial activity.<sup>2</sup> Three semisynthetic derivatives of rifamycin A, including rifampin, rifapentine, and rifabutin, are currently in clinical use for treatment of Mycobacterial infections, including tuberculosis and leprosy.<sup>3</sup> Despite the high potency, low toxicity, and broad antibacterial spectrum of rifamycins, pathogens develop resistance to this class of antibiotics at a relatively high rate by

substitution of the amino-acid residues in the rifamycin binding site of bacterial RNAP.<sup>4</sup> Rapid onset of bacterial resistance is the primary reason why current use of rifamycins is restricted to combinations with other drugs, such as isoniazid, or to clinical emergencies. Thus, there is a significant need for the development of new antibiotics that target bacterial RNAPs by different biochemical mechanisms and display broad-spectrum antibacterial activity. Several other classes of natural products have been shown to inhibit bacterial RNAPs by binding to alternative regions of this multisubunit protein, which typically yields notable antibiotic activity. Such compounds were found to be effective against rifamycin-resistant RNAPs and strains.<sup>5</sup>

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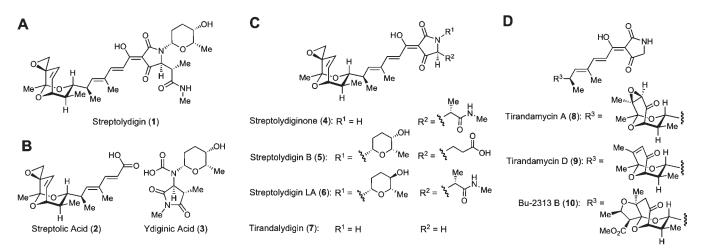


Figure 1. Structures of dienoyl tetramic acids antibiotics and degradation fragments. (A) Structure of streptolydigin (1), the parent member of this antibiotic family, which was isolated from *S. lydicus*. (B) Structures of streptolic acid (2) and ydiginic acid (3), which were obtained by chemical degradation of streptolydigin (1). (C) Structures of several derivatives of streptolydigin, which were biosynthetically produced in re-engineered *S. lydicus* strains, and structure of tirandalydigin (7), which was isolated from *S. tirandis*. (D) Structures of two representative tirandamycins (8 and 9), which were isolated from *S. tirandis* and marine-derived *Streptolyces* species, as well as Bu-2313B (10), which was isolated from oligosporic actinomycete strain, no. E864-61.

Streptolydigin (1, Figure 1A) is a dienoyl tetramic acid antibiotic,<sup>6</sup> which elicits its antibacterial activity by inhibiting initiation, elongation, and pyrophosphorylation steps of bacterial RNAP.<sup>7</sup> High-resolution X-ray crystallographic characterization of the streptolydigin-RNAP complex revealed a unique biochemical mechanism of RNAP inhibition.8 Streptolydigin (1) traps the bridge-helix of the RNAP in a straight conformation and induces opening of the trigger-loop of the enzyme. As a result, streptolydigin (1) stabilizes the catalytically inactive substratebound transcription intermediate and blocks structural isomerization of RNAP into a fully active state, which requires conformational changes of both the bridge-helix and the triggerloop moieties.8 The streptolydigin-binding region is located 20 Å away from the rifamycins binding site.<sup>9</sup> As a result of this unique biochemical mechanism of RNAP inhibition and a distinct binding site, streptolydigin (1) and rifamycins exhibit only minimal cross-resistance.<sup>8b,10</sup>

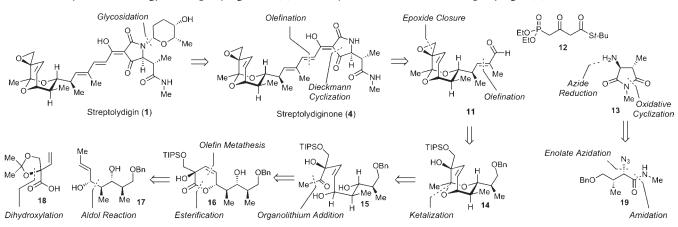
The structure of streptolydigin (1) features an epoxide-containing bicyclic ketal connected by a polyene spacer to a highly substituted, glycosylated acyl tetramic acid. Structure elucidation of streptolydigin (1) entailed initial oxidative degradation of the natural product into two simplified subunits, streptolic acid (2)and ydiginic acid (3), which derived from the bicyclic ketal fragment and the tetramic acid subunit of the natural product, respectively (Figure 1B).<sup>11</sup> Complete stereochemical assignment of streptolic acid (2) was ultimately secured by X-ray crystal-lographic analysis.<sup>12</sup> Re-engineering of streptolydigin biosynthesis in S. lydicus enabled recent production of several new antibiotics shown in Figure 1C, including streptolydiginone (4),<sup>13</sup> which represents a streptolydigin aglycone, as well as streptolyidigin B  $(5)^{14}$  and streptolydigin LA (6).<sup>13</sup> Following the initial report on isolation of streptolydigin in 1956,6 several other members of the dienoyl tetramic acid antibiotic family have been identified, including tirandalydigin (7),<sup>15</sup> tirandamycins (i.e., 8 and 9),<sup>16</sup> Bu-2312B (10),<sup>17</sup> and nocamycins, which were found to be structurally analogous to 10.18 The bicyclic ketal subunit of tirandalydigin (7) is identical to that of streptolydigin (Figure 1C). However, the tetramic acid moiety of this

metabolite lacks L-rhodinose and the amide containing sidechain. Tirandamycins (8 and 9) and Bu-2312B (10) possess the same unsubstituted acyl tetramic acid subunit of tirandalydigin (7), but differ in the substitution of the ketal moiety (Figure 1D). Despite isolation and biosynthetic production of a number of structurally homologous dienoyl tetramic acid antibiotics over the years, streptolydigin (1) features the most elaborate structure and the highest antimicrobial activity recorded within this class.

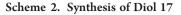
Evaluation of the antibiotic activity of streptolydigin (1) against a broad panel of microbial strains using standard broth dilution testing revealed that this natural product elicited notable activity against a number of Gram-positive organisms.<sup>6a</sup> Inhibition of several representative *Clostridium* and *Streptococcus* species by 1 was particularly potent with minimum inhibitory concentrations (MICs) as low as 0.04  $\mu$ g/mL. In addition, inhibition of *Mycobacterium* strains, including *M. tuberculosis*, was also observed.<sup>6a</sup> Streptolydigin (1) was shown to be generally more potent than tirandalydigin (7) and tirandamycin A (8) in a panel of several anaerobic and aerobic gram-positive bacteria.<sup>19</sup> While streptolydiginone (4) and streptolydigin B (5) inhibited growth of *S. albus*, they were found to be less active as compared to streptolydigin (1) and streptolydigin LA (6).<sup>13,14</sup>

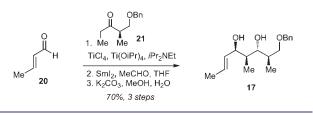
Streptolydigin (1) was also evaluated in a series of in vivo studies conducted in mice.<sup>6c</sup> The maximum tolerated doses of 1 were 1800 mg/kg/day when administered orally and 500 mg/kg/day when administered subcutaneously.<sup>6c</sup> Importantly, administration of streptolydigin (1) at subcutaneous doses of 100–330 mg/kg/day protected animals against infection with *S. hemolyticus, S. pneumoniae,* and *P. multocida.*<sup>6c</sup> Similarly, Bu-2313B (10) was effective in protecting mice against *P. fragilis* and *C. perfingens* when administered by both oral and subcutaneous routes.<sup>17a</sup>

While several total syntheses of tirandamycin A  $(8)^{20}$  have been developed over the years and the preparation of two advanced fragments of streptolydigin (1) has been described,<sup>21</sup> development of viable synthetic approaches to streptolydigin (1) and closely related tirandalydigin (7) remained challenging in part due to the presence of a highly labile alkenyl epoxide



#### Scheme 1. Synthetic Strategy to Streptolydiginone (4) and Projected Conversion to Streptolydigin (1)





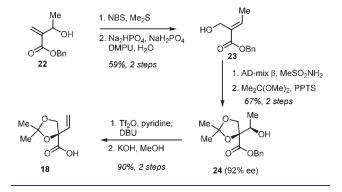
fragment, which is absent in tirandamycin A (8). Indeed, while a recent synthetic effort enabled assembly of a protected tirandalydigin (7), the final acid-mediated deprotection step was not successful presumably due to the instability of the epoxide moiety.<sup>22</sup>

Following our initial communication of the synthesis of streptolydigin (1),<sup>23</sup> which afforded this natural product with a longest linear sequence of 24 steps, we describe here the evolution of our synthetic approach to this important class of RNAPtargeting antibiotics. This effort resulted in the assembly and biological evaluation of a series of streptolydigin-based agents, which demonstrated that the presence of both streptolic acid and tetramic acid subunits was required for activity of streptolydigin antibiotics. In addition, we identified a new RNAP-targeting synthetic antibiotic, 10,11-dihydrostreptolydigin, which was assembled with the high synthetic efficiency of 15 steps in the longest linear sequence. Dihydrostreptolydigin inhibited three representative bacterial RNAPs and displayed in vitro antibacterial activity against S. salivarius. The overall increase in synthetic efficiency combined with substantial antibacterial efficacy of this synthetic congener of streptolydigin demonstrates the power of organic synthesis in enabling design and comprehensive in vitro pharmacological evaluation of new antibiotics that target bacterial transcription.

#### RESULTS AND DISCUSSION

Syntheses of Streptolic Acid, Methyl Streptolate, and Streptolydiginone. Guided by the logic of the streptolydigin biogenesis, which entails final-stage glycosylation of streptolydiginone (4) with L-rhodinose,<sup>13</sup> we devised our initial synthetic strategy shown in Scheme 1. Streptolydiginone (4) would derive

Scheme 3. Synthesis of Carboxylic Acid 18

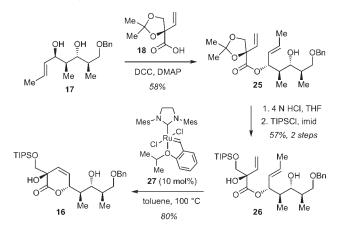


from three simplified fragments, including aldehyde 11, diethylphosphono-3-oxobutanthioate 12, and imide 13. The assembly process would entail the Horner-Wadsworth-Emmons olefination of aldehyde 11 with phosphonate 12, followed by N-acylation of imide 13 and Dieckmann cyclization under the protocol developed by Rinehart.<sup>24</sup> The imide 13 would derive from azide 19, which would be in turn assembled by the Evans diastereoselective enolate azidation.<sup>25</sup> The assembly of aldehyde 11 would entail late-stage installation of the epoxide moiety starting with appropriately protected vicial diol 14 containing the requisite bicyclic core of the streptolic acid, which would be derived from the corresponding triol 15. To control the cisgeometry of alkene 15, we envisioned that this acyclic product would be prepared starting from lactone 16 by a controlled carbonyl addition. The synthesis of lactone 16 would entail chemoselective acylation of diol 17 with carboxylic acid 18, followed by a ring-closing metathesis.

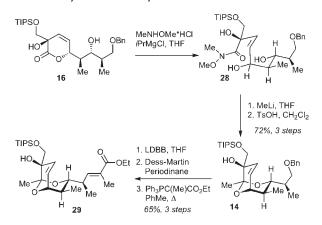
Preparation of diol 17 (Scheme 2) commenced with the aldol reaction of crotonaldehyde (20) with titanium enolate derived from benzyloxyketone  $21^{26}$  to give the corresponding *syn* aldol product with excellent diastereoselectivity (dr >95:5). Subsequent anti-selective Evans—Tishchenko reduction<sup>27</sup> and saponification of the resulting acetate delivered the target diol 17 in 70% yield for three steps as a single diastereomer.

The synthesis of carboxylic acid **18** is shown in Scheme 3. The sequence began from hydroxy enoate **22**, which was readily prepared by Baylis—Hillman reaction of benzyl acrylate with acetaldehyde. Alcohol transposition in **22** was achieved via its initial conversion into the corresponding allylic bromide, which underwent hydrolysis in aqueous buffer to give primary alcohol **23**. Subjection of **23** to Sharpless dihydroxylation,<sup>28</sup> followed by chemoselective protection of the corresponding triol as the acetonide, delivered secondary alcohol **24** in 92% ee. Conversion of alcohol **24** into the corresponding triflate, followed by elimination with DBU and hydrolysis of the benzyl ester, gave the requisite carboxylic acid **18** containing a terminal alkene.

Scheme 4. Synthesis of Lactone 16



Scheme 5. Synthesis of Bicyclic Ester 29

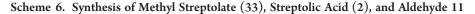


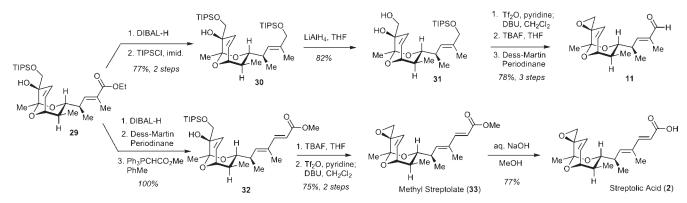
Chemoselective acylation of alcohol 17 with acid 18 was achieved in the presence of dicyclohexyl carbodiimide (DCC) and dimethylaminopyridine (DMAP) to give ester 25 (Scheme 4). Subjection of 25 directly to a series of Ru-based alkene metathesis catalysts afforded only low yields of the desired unsaturated lactone, presumably due to considerable steric bulk presented by the acetonide moiety. Acid-promoted removal of acetonide (4 N aqueous HCl in THF), followed by protection of the primary alcohol as a tri-isoprolylsilyl (TIPS) ether, afforded diene 26. Ring-closure of 26 to lactone 16 proved to be superior to that using the acetonide 25. The optimum efficiency was achieved using Ru catalyst  $27^{29}$  to give lactone 16 in 80% yield.

We next examined the conversion of lactone 16 to the requisite bicyclic ketal. While direct conversion of 16 to the corresponding methyl ketone using methylithium proved unsuccessful, this transformation could be efficiently achieved in a two-step fashion involving initial formation of the Weinreb amide 28, followed by methyllithium addition (Scheme 5). Subsequent acid-catalyzed intramolecular acetalization afforded bicyclic ketal 14 in 72% yield for three steps. Reductive deprotection of benzyl ether, followed by Dess—Martin oxidation of the resulting alcohol and Wittig olefination, furnished ester 29, a key structural subunit for the assembly of streptolydigin antibiotics.

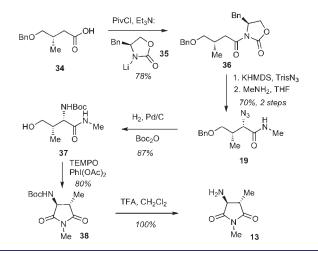
Subsequent elaboration of **29** required installation of the exocyclic epoxide and conversion of the ester to the aldehyde moiety. The optimized sequence for this functional group transformation began with 1,2-reduction of unsaturated ester **29** using DIBAL-H, followed by protection of the resulting allylic alcohol as a bis-TIPS ether **30** (Scheme 6). Chemoselective removal of the TIPS ether adjacent to the unprotected tertiary alcohol was efficiently achieved by subjecting **30** to LiAlH<sub>4</sub>, presumably due to the participation of the neighboring alkoxide in the desilylation step. The resulting diol **31** was converted to the epoxide via intermediacy of the corresponding primary triflate. Finally, deprotection of the silyl ether and Dess–Martin oxidation efficiently afforded aldehyde **11**.

We also examined the conversion of ester 29 to streptolic acid (2). While streptolic acid represents a product of chemical degradation of streptolydigin, the importance of this synthetic effort was 2-fold. First, this conversion would enable us to verify the correct structure of ester 29 by correlating analytical data of the synthetically produced streptolic acid (2) with those present in the literature. Second, the synthesis of compounds containing the fully functionalized bicyclic ketal core of streptolydigin, which was connected to the dienoyl moiety, would enable us

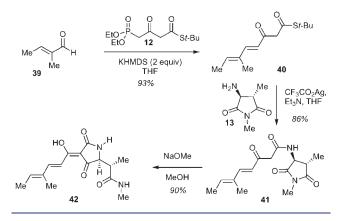




## Scheme 7. Synthesis of Imide 13



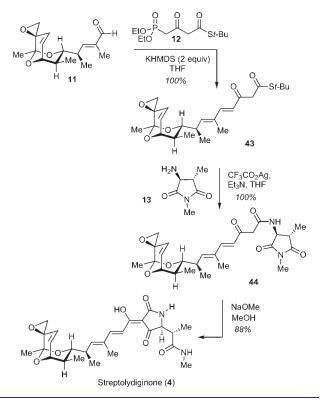
Scheme 8. Synthesis of 4-Methyl-2,4-hexadienoyl Tetramic Acid (42)



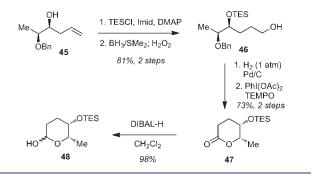
to test the importance of the streptolic acid subunit for biological activity of the parent natural product (vide infra). This transformation began with the reduction of ester **29** with DIBAL-H, followed by the Dess–Martin oxidation of the resulting alcohol and Wittig olefination to give chain-extended ester **32** with excellent efficiency (Scheme 6). Fluoride-mediated desilylation of **32**, followed by conversion of the resulting diol to an epoxide via the intermediacy of the corresponding primary triflate, gave methyl streptolate (**33**). Subjection of **33** to basic hydrolysis afforded streptolic acid (**2**), which proved to be identical to the corresponding degradation product of streptolydigin, thus securing the chemical structure of all prior synthetic intermediates.<sup>11,21c</sup>

Having secured synthetic access to the streptolic acid moiety, we turned our attention to the construction of the tetramic acid fragment, which would arise from aspartimide **13** (Scheme 7). The assembly process began with the conversion of known acid **34** to the corresponding imide **36** by addition of lithium oxazolidinone **35** to the mixed anhydride derived from **34**. The diastereoselective azide transfer was performed according to the Evans protocol,<sup>25</sup> which entailed generation of the potassium enolate, followed by treatment with 2,4,6-tri-isopropylbenzenesulfonyl azide (TrisN<sub>3</sub>). The resulting

## Scheme 9. Synthesis of Streptolydiginone (4)



Scheme 10. Synthesis of Protected L-Rhodinose (48)



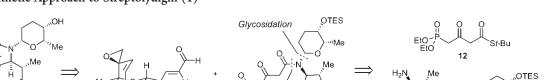
imide was treated with methylamine to give amide **19**. Reduction of azide **19** under a hydrogen atmosphere with simultaneous hydrogenolysis of the benzyl ether, followed by in situ Bocprotection of the resulting amine, furnished carbamate **37**. Subjection of amide **37** to a TEMPO-mediated oxidative cyclization afforded the corresponding aspartimide **38**, which was treated with trifluoroacetic acid (TFA) to enable Boc-deprotection to give primary amine **13**.

We next examined the assembly of the simplified methyl hexadienoyl tetramic acid **42**, which corresponded to the structure of streptolydiginone (**4**) with a fully truncated bicyclic ketal (Scheme 8). This sequence was designed to model the final stages of the streptolydiginone synthesis and would also enable us to test the importance of the tetramic acid subunit of this agent on its biological activity (vide infra). The sequence began with the Horner–Wadsworth–Emmons olefination of unsaturated

Olefination

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49

EtO'

11

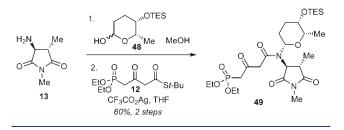
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Scheme 12. Synthesis of Phosphonate 49

Dieckmann

Cyclization

Streptolydigin (1)



aldehyde **39** with known phosphonate  $12^{30}$  to give keto thioester **40**. Treatment of **40** with silver(I) trifluoroacetate in the presence of amine **13** delivered ketoamide **41**, which underwent Dieckmann cyclization<sup>12</sup> upon exposure to methanolic sodium methoxide to give tetramic acid **42** in 90% yield.

The stage was now set for completing the assembly of streptolyidiginone (4). To this end, we subjected aldehyde 11 to the Horner-Wadsworth-Emmons olefination with phosphonate  $12^{30}$  (Scheme 9). This transformation efficiently delivered the desired keto thioester 43. Subsequent treatment of 43 with silver(I) trifluoroacetate in the presence of amine 13 resulted in the formation of ketoamide 44. Under such conditions, both steps proceeded with quantitative efficiency without any detectable degradation of the labile epoxide moiety. The next step entailed the Dieckmann cyclization of 44 using a methanolic solution of sodium methoxide, according to the Rinehart protocol,  $^{12}$  to deliver streptolydiginone (4) in 88% yield and a longest linear sequence of 26 steps. The structure of 4 was confirmed by comparison of the 500 MHz <sup>1</sup>H NMR and 125 MHz <sup>13</sup>C NMR spectral data for the sodium salt of synthetic streptolydiginone with the corresponding NMR spectra for this aglycon, which was isolated by the Salas group from the re-engineered S. lydicus strain with suppressed glycosylation of streptolydigin.<sup>13</sup>

We next examined the glycosylation of streptolydiginone (4). The synthesis of the L-rhodinose moiety of streptolydigin is depicted in Scheme 10. Protection of the known homoallylic alcohol  $45^{30}$  as a triethylsilyl (TES) ether, followed by hydroboration-oxidation of the alkene, gave alcohol 46. Hydrogenolysis of the benzyl ether, followed by PhI(OAc)<sub>2</sub>-mediated oxidative cyclization of the resulting diol, afforded lactone 47. Reduction of 47 with DIBAL-H delivered the desired lactol 48.

Our exploratory studies revealed that simple acyl tetramic acids could be efficiently glycosylated with protected L-rhodinose derivatives by treatment of the corresponding lactol or glycal with tetramic acid in choloroform at 20 °C in the presence of a catalytic amount of *p*-TsOH. However, glycosylation of the streptolydiginone's tetramic acid moiety could not be achieved under the same reaction conditions presumably due to the increased steric bulk of the branched alkyl substituent in the direct proximity to the reactive N-H bond of the tetramic acid.

Me 13

Synthesis of Streptolydigin. Because of the inability to glycosylate streptolydiginone (4), we examined an alternative synthetic approach to streptolydigin (1), which entailed installation of the glycoside moiety prior to the formation of the final tetramic acid (Scheme 11). The key disconnection of streptolydigin (1) was designed to unite two advanced fragments, including bicyclic aldehyde 11 and phosphonate 49. The latter would be assembled by N-glycosylation and N-acylation starting with three fragments 12, 13, and 48, which were previously prepared in our route to streptolydiginone (4). It is noteworthy that phosphonates similar to 49 were prepared by both the Boeckman and the Schlessinger laboratories.<sup>21a,b</sup> However, no attempts to use such fragments for elaboration of streptolydigin have been reported. The main challenge was to ensure that the projected assembly process would be compatible with the highly labile terminal epoxide moiety of aldehyde 11.

Following a series of exploratory studies, we devised a simple, two-step sequence for the assembly of *N*-glycosyl phosphonate **49** (Scheme 12). Direct treatment of amine **13** with TES-protected rhodinose **48** afforded the desired *N*-glycosylated product as a single requisite diastereomer. Subsequent Ag-promoted *N*-acylation of this glycosyl amine with thioester **12**<sup>31</sup> afforded phosphonate **49** in 60% yield for two steps.

After significant optimization, we ultimately developed a oneflask protocol for synthesis of streptolydigin (1), which entailed treatment of the phosphonate 49 with 3 equiv of t-BuOK, followed by the addition of aldehyde 11 and a rapid acidic workup with 1 M aqueous solution of HCl at 0 °C (Figure 2A). This onepot process entailed several consecutive steps, including initial deprotonation of 49 to produce enolate I (Figure 2B), Dieckmann cyclization to give tetramic acid II, subsequent abstraction of two acidic protons to give III, followed by the Horner-Wadsworth-Emmons olefination and final neutralization with concomitant acid-mediated TES deprotection to furnish 1. Synthetic streptolydigin (1) proved to be identical in all respects to the authentic sample of the natural product.<sup>32</sup> This sequence enabled the assembly of this natural product with a longest linear sequence of 24 steps and a higher level of convergency as compared to our initial route, which afforded streptolydiginone (4) in 26 chemical steps.

RNAP Inhibitory Activity of Streptolydigin, Streptolydiginone, Streptolic Acid, and Tetramic Acid. Our next objective was to examine the inhibition of bacterial RNAP by streptolydigin (1), streptolydiginone (4), as well as other compounds prepared during our initial synthetic studies. While antibacterial activity of several members of this antibiotic family has been reported, the RNAP inhibitory profile has only been studied for streptolydigin (1)<sup>7,8</sup> and tirandamycin A (8).<sup>32</sup> These studies

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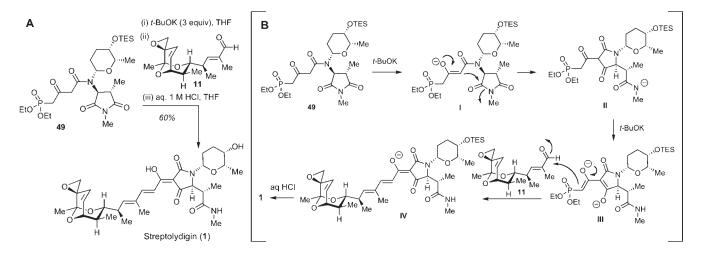
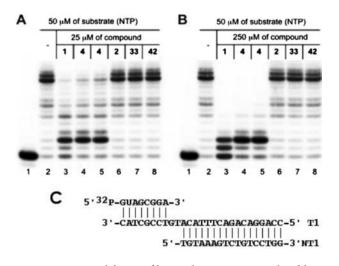


Figure 2. Completion of the streptolydigin synthesis. (A) Optimized one-flask protocol for conversion of phosphonate 49 and aldehyde 11 to streptolydigin (1). (B) Mechanism of streptolydigin formation, which entails initial Dieckmann cyclization to give II, Horner–Wadsworth–Emmons olefination of aldehyde 11 with III, followed by subsequent acidification of IV with concomitant removal of the TES group to give 1.



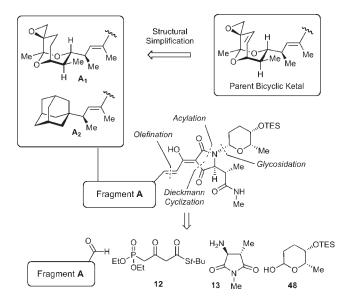
**Figure 3.** In vitro inhibition of bacterial transcription catalyzed by *T. aquaticus* RNAP. Artificial transcription elongation complexes containing radioactively labeled 8-nt RNA primer were assembled on nucleicacid scaffolds (C) using *T. aquaticus* RNAP core enzyme. Where indicated, reaction mixtures were supplemented with 25 (A) or 250  $\mu$ M (B) of small-molecule inhibitors, and transcription was initiated by the addition of NTPs. Reaction products were resolved by denaturing polyacrylamide gel electrophoresis and revealed by autoradiography.

demonstrated that tirandamycin A was a substantially less potent bacterial RNAP inhibitor.<sup>33</sup> The antimicrobial data of dienoyl tetramic acid antibiotics have been obtained using only a small number of compounds, isolated from various microbial producers, which provided only limited structure—activity relationship data. While notable success in re-engineering streptolydigin biosynthesis has been achieved, this approach has thus far provided only a few compounds with modified glycoside and tetramic acid subunits of the parent natural product.<sup>13,14</sup>

Our study was aimed at evaluating the importance of two major structural subunits of streptolydigin (1), including streptolic acid and the tetramic acid fragments. To this end, we examined inhibition of a well-characterized *Thermus aquaticus* RNAP<sup>34</sup> by following in vitro transcription in the presence of

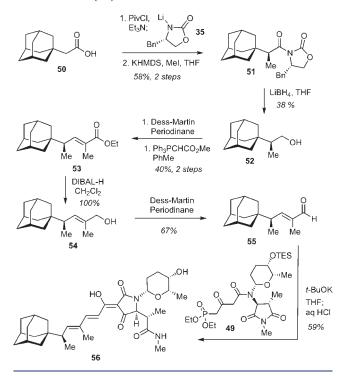
50  $\mu$ M of nucleotide triphosphates (NTPs) and two inhibitor concentrations (25 and 250 µM, Figure 3A and B, respectively). The assay consisted of assembling T. aquaticus RNAP complexes with nucleic acid scaffolds that mimic the structure of nucleic acids in the transcription elongation complex (Figure 3C). It was previously shown that artificial transcription elongation complexes assembled on nucleic acid scaffolds are functional<sup>35</sup> and can be used to monitor the effects of various inhibitors of transcription elongation.<sup>36</sup> To monitor RNAP-catalyzed elongation of the 8-nucleotide-long RNA component of the scaffold, it was radioactively labeled at the 5' end, allowing its visualization after denaturing gel-electrophoresis (lane 1). The addition of NTPs led to the appearance of a runoff product, which resulted from RNAP-catalyzed extension of the nascent transcript until the end of the template (lane 2). The addition of synthetic streptolydigin (1, 25  $\mu$ M) inhibited the appearance of runoff transcript (Figure 3A, lane 3), an expected result for a transcription elongation inhibitor. A shorter transcript that has undergone only two cycles of nucleotide addition was formed. At 250  $\mu$ M of inhibitor, the amount of the extended transcript decreased, and a significant amount of the initial nascent transcript remained intact (a product of a single nucleotide addition was also observed, Figure 3B, lane 3). The same pattern of transcription elongation inhibition was observed when an authentic sample of 1, which was isolated from *S. lydicus*, was used (data not shown). We also found that streptolydiginone (4), both as a free tetramic acid (lane 4) and as a corresponding sodium salt (lane 5), was effective at inhibiting T. aquaticus RNAP. At the highest concentration tested, the observed RNAP inhibitory activity of 4 was visibly lower than that of streptolydigin (compare lanes 3 with lanes 4 and 5, Figure 3B). This observation is correlated with weaker antibacterial activity of streptolydiginone (4) against S. albus, which was reported previously.<sup>13</sup>

We were unable to observe any RNAP inhibition by either the streptolic acid (2, lane 6) or the corresponding methyl ester (33, lane 7). Furthermore, the truncated tetramic acid subunit of streptolydiginone, which is represented by compound 42, was also inactive at both concentrations tested (lane 8). These results demonstrate that the presence of both structural subunits of streptolydigin is essential for activity of this class of antibiotics.

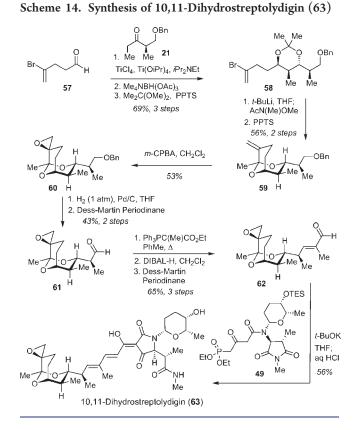


**Figure 4.** General strategy to simplified analogues of streptolydigin. The synthetic approach would entail a sequential condensation of four reaction partners, including aldehyde-containing fragment **A**, phosphonate **12**, imide **13**, and protected L-rhodinose **48**. The projected assembly process entails *N*-glycosidation, *N*-acylation, Dieckman cyclization, olefination, and final deprotection.

# Scheme 13. Synthesis of Adamantane-Substituted Dienoyl Tetramic Acid (56)



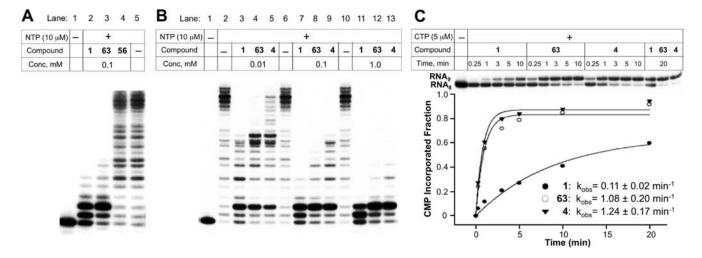
This conclusion is also in agreement with previous crystallographic studies of streptolydigin—RNAP complexes, which established that both streptolic and tetramic fragments of the natural product interacted with the bridge-helix and the triggerloop regions of the protein, respectively.<sup>8</sup> Thus, removal of each



of the individual structural subunits of streptolydigin would result in a substantial decrease of the binding efficiency and loss of inhibitory activity.

Rationally Simplified Streptolydigin Analogues: Design, Synthesis, and RNAP Inhibition. Our initial biochemical analysis revealed that truncation of either streptolic or tetramic acid subunits of streptolydigin resulted in complete loss of RNAP inhibitory activity. We envisioned that additional modifications could be introduced into the streptolic subunit of the natural product provided that the fully functionalized tetramic acid fragment remains intact (Figure 4). Specifically, we planned to remove the C(10)-C(11) alkene moiety in the bicyclic ketal core of the natural product (fragment  $A_1$ , Figure 4). This structural alteration was expected to considerably increase the efficiency of our assembly process (vide infra). In addition, we intended to substitute the entire bicyclic ketal moiety with an adamantyl fragment, which was expected to preserve the overall shape and most of the physicochemical properties of this relatively hydrophobic subunit (fragment A2, Figure 4). Each new compound would derive from a highly convergent sequence that would entail a condensation of four building blocks following our previously established synthetic sequences (Figure 4). Because of structural simplification of fragment A, we envisioned that such derivatives would be assembled with significantly higher synthetic efficiency as compared to our original routes to streptolydigin (1) and streptolydiginone (4), which required 24 and 26 steps, respectively.

The synthesis of the adamantane-bearing derivative began with asymmetric Evans alkylation<sup>37</sup> of chiral imide derived from the commercially available carboxylic acid **50** (Scheme 13). This process delivered the requisite alkylation product in good



**Figure 5.** In vitro inhibition of transcription catalyzed by *T. aquaticus* RNAP by streptolydigin derivatives. (A) A comparison of RNAP inhibitory activity of streptolydigin (1), 10,11-dihydrostreptolydigin (63), and adamantane derivative 56 using 10  $\mu$ M NTP and 0.1 mM of each inhibitor. (B) Dose-dependent RNAP inhibition by streptolydigin (1), streptolydiginone (4), and 10,11-dihydrostreptolydigin (63). (C) Quantitative assessment of inhibitory effects of streptolydigin (1), streptolydiginone (4), and 10,11-dihydrostreptolydigin (63) on the rate of CMP incorporation into nascent 8-nt starting RNA.

efficiency and excellent diastereomeric purity. Subsequent elaboration entailed reduction of imide **51** with LiBH<sub>4</sub>, followed by Dess—Martin oxidation of alcohol **52**, and Wittig olefination of the resulting aldehyde to give unsaturated ester **53**. Reduction of **53** with DIBAL-H, followed by Dess—Martin oxidation of the allylic alcohol **54**, afforded aldehyde **55**. Treatment of aldehyde **55** with phosphonate **49** in the presence of excess of *t*-BuOK using our previously developed one-flask protocol for sequential olefination-cyclization-deprotection afforded requisite adamantane-containing tetramic acid **56** in 59% yield (13-step longest linear sequence from commercially available precursors).

Construction of 10,11-dihydrostreptolydigin (63) commenced with a Ti-mediated aldol reaction between ketone 21 and aldehyde 57, followed by an anti-reduction of the corresponding hydroxyketone to afford diol, which was protected as acetonide 58 (Scheme 14). Bromine-lithium exchange and subsequent acylation gave the expected enone, which was treated with PPTS in methanol to generate bicyclic ketal 59. Concaveface epoxidation of alkene 59 delivered desired epoxide 60. Hydrogenolysis of 60, followed by Dess-Martin oxidation of the resulting alcohol, afforded aldehyde 61. Subjection of 61 to a Wittig olefination furnished an unsaturated ester, which was subjected to reduction with DIBAL-H, followed by Dess-Martin oxidation to deliver aldehyde 62. Final assembly of 10,11-dihydrostreptolydigin (63) was performed under the conditions developed for the preparation of streptolydigin. Treatment of ketoamide 49 with excess potassium tert-butoxide, followed by addition of aldehyde 62 and acidic workup, afforded 10,11-dihydrostreptolydigin (63). Removal of the C(10)-C(11) alkene moiety enabled significant increase in the overall synthetic efficiency of this route (15-step longest linear sequence) as compared to our original approach to streptolydigin, which required 24 synthetic steps.

Having completed the syntheses of compounds **56** and **63**, we examined their ability to inhibit transcription catalyzed by *T. aquaticus* RNAP (Figure 5) using a nucleic acid scaffold similar to the one presented in Figure 3C but having a longer downstream (transcribed) segment. Streptolydigin (1) was included as a

positive control (lane 2, Figure 5A). We found that 10,11dihydrostreptolydigin (63) inhibited *T. aquaticus* RNAP (lane 3, Figure 5A) very similarly to streptolydigin (1), with transcript elongation blocked after incorporation of two nucleotides into the growing RNA strand. This observation suggested that removal of the C(10)-C(11) alkene preserved a substantial amount of the parent activity from the parent natural product 1 (lane 2, Figure 5A). Interestingly, we did not observe any inhibition of RNAP by adamantane-containing analogue 56 at 100  $\mu$ M (lane 4, Figure 5A) or 1 mM (data not shown) concentrations. This finding indicated that the adamantane core of 56 did not provide a viable replacement for a bicyclic ketal moiety of streptolydigin (1).

We next compared the relative efficacy of T. aquaticus RNAP inhibition by three compounds, which were shown to elicit activity, including streptolydigin (1), streptolydiginone (4), and 10,11-dihydrostreptolydigin (63). The results are presented in Figure 5B. At the two highest concentrations of streptolydigin (1) (0.1 and 1.0 mM), the transcription was blocked either completely or following the addition of one or two nucleotides (Figure 5B, lanes 7 and 11). At the same concentrations of streptolydiginone (4) and 10,11-dihydrostreptolydigin (63), inhibition occurred almost exclusively after incorporation of two nucleotides (Figure 5B, lanes 8, 9, 12, and 13). These results once again confirmed that 10,11-dihydrostreptolydigin (63) and streptolydiginone (4) were effective at inhibiting transcription in vitro. This conclusion was further supported by an experiment conducted in the presence of subsaturating  $(10 \,\mu\text{M})$  concentrations of inhibitors (Figure 5B, lanes 3, 4, and 5). Similar results were obtained in experiments using a closely related T. thermophilus RNAP, as well as RNAPs from M. smegmatis (data not shown). RNAPs from Gram-negative organisms such as Francisella tularensis and E. coli required much higher concentrations of inhibitors to achieve the same extent of inhibition (data not shown), confirming earlier observations that RNAPs from Gramnegative bacteria are significantly more resistant to streptolydigin than their counterparts from Gram-positive organisms.

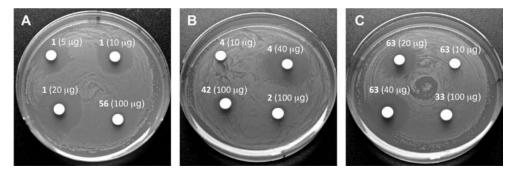


Figure 6. Antimicrobial activity of synthetic streptolydigin antibiotics against *S. salivarius*. Each compound was tested according to the general protocol described in the Experimental Procedures. Most potent compounds, including 1, 4, and 63, were tested at several concentrations shown. All other compounds were tested at a concentration of 100  $\mu$ g/disk.

We also followed the rate of single nucleotide incorporation catalyzed by T. aquaticus RNAP in the presence of streptolydigin (1), streptolydiginone (4), and 10,11-dihydrostreptolydigin (63). This study provided a quantitative basis for assessing the difference in inhibitory activities of each of the three compounds tested (Figure 5C). In this assay, transcription complexes containing radioactively labeled nascent transcript were supplemented with CTP, which was necessary to extend the transcript by only one nucleotide. The concentration of CTP (5  $\mu$ M) and the reaction temperature (25 °C) were kept low to slow the RNAP catalysis rate, which enabled following the CMP addition in real time in the presence of inibitors. In the absence of inhibitors, the reaction was complete (full conversion of 8-nt starting RNA to 9-nt product) at the first time point sampled (15 s, data not shown). In the presence of inhibitors, the reaction rate was slowed considerably, with full extension achieved at 5 min for streptolydiginone (4) and 10,11-dihydrostreptolydigin (63), which appeared to inhibit nucleotide addition with similar efficiency. Streptolydigin (1) inhibition was more potent, with nascent transcript extension reaction being  $\sim$  50% complete after 20 min of incubation. The data obtained were used to derive the apparent rates of RNAP-catalyzed single nucleotide extension, which were found to be 0.11, 1.08, and 1.24  $\min^{-1}$  in the presence of streptolydigin (1), 10,11-dihydrostreptolydigin (63), and streptolydiginone (4), respectively. Overall, the results of in vitro transcription assays establish that removal of either the glycoside moiety of streptolydigin or the C(10)-C(11) alkene preserved RNAP inhibitory activity of the resulting compounds, while weakening their in vitro potency by about 10-fold.

Antimicrobial Activity against S. salivarius. We also examined antimicrobial activity of all streptolydigin-based compounds prepared in our study. Originally, streptolydigin (1) was found to be effective in inhibiting several Streptococcus species, including S. faecalis, S. hemolyticus, S. lactis, S. mitis, S. viridans, and S. pneumoniae with MICs in the range of 0.19–3.12  $\mu$ g/mL.<sup>6a</sup> Therefore, we selected S. salivarius, a representative Streptococcus strain, which typically colonizes human oral and nasopharyngeal epithelia and only becomes harmful to the host if the immune status is altered or there is a loss of control of epithelial cell sensing and discriminatory systems.<sup>38</sup> Our main obective was to use this model, normally nonpathogenic organism to correlate the initially observed RNAP inhibitory activity of all compounds with their antibacterial action. The results of this study, which employed a standard disk-diffusion protocol, are shown in Figure 6. As expected, streptolydigin (1) inhibited growth of S. salivarius in a dose-dependent manner at concentrations of 5, 10,

and 20  $\mu$ g per disk (Figure 6A). Streptolydiginone (4) was also found to elicit antibacterial activity, albeit not as potent as that of streptolydigin (1). While substantial growth inhibition was observed at  $40 \mu g$  of 4 per disk, no effect on bacterial proliferation occurred at a lower dose of 10  $\mu$ g per disk (Figure 6B). Evaluation of 10,11-dihydrostreptolydigine (63) revealed dosedependent antibacterial action at concentrations of 20 and 40  $\mu$ g per disk (Figure 6C). Similar to streptolydiginone (4), no antibacterial activity was observed with 63 at 10  $\mu$ g per disk (Figure 6C). Overall, the observed antibacterial activities of 1, 4, and 63 were in excellent correlation with their RNAP inhibitory profiles. In addition, streptolic acid (2) and methyl streptolate (33) were found to be completely inactive even at a high concentration of 100  $\mu$ g/disk (Figure 6B and C). The adamantane-containing derivative 56 and tetramic acid 42 elicited only weak antibacterial action at 100  $\mu$ g/disk (Figure 6A and B) and no activity at lower concentrations tested (data not shown). The lack of activity of 2, 33, 42, and 56 was also in excellent agreement with the inability of such compounds to inhibit T. aquaticus RNAP. This study demonstrated once again that the presence of both streptolic and tetramic subunits of streptolydigin is required for its antibiotic activity and established that RNAP inhibition of streptolydigin-based compounds was well correlated with their antibacterial profile.

## CONCLUSIONS

We described the development of a general synthetic strategy to streptolydigin antibiotics, as well as their comprehensive biochemical and antimicrobial evaluation. Our initial studies resulted in the total synthesis of streptolydiginone (4) and streptolydigin (1) and enabled the assembly of a series of simplified derivatives of the parent natural products, which were designed to assess the importance of the individual streptolic and tetramic acid subunits. Indeed, the presence of both major fragments of streptolydigin was found to be essential for its activity. This finding was in agreement with previous X-ray crystallographic characterization of streptolydigin-RNAP complexes. In addition, our effort enabled a 15-step synthesis of 10,11-dihydrostreptolydigin, which inhibited three bacterial RNAPs and elicited antimicrobial activity against S. salivarius. The ability to access efficiently simplified analogues of streptolydigin that elicit substantial antibacterial activity demonstrates the power of our general synthetic approach, which could be employed for subsequent rational design and

the development of new streptolydigin-based antibiotics with enhanced potency and improved pharmacological profile.

#### EXPERIMENTAL PROCEDURES

Streptolic Acid (2). A solution of methyl streptolate 33 (13.0 mg, 0.039 mmol) in a mixture of methanol (1 mL) and water (0.1 mL) was treated with sodium hydroxide (16.0 mg, 0.39 mmol). The reaction mixture was left overnight at room temperature, concentrated in vacuo, and treated with aqueous HCl (0.4 mL of 1 N solution). The resulting solution was extracted twice with ethyl acetate. Combined organic extracts were washed with brine, dried over MgSO<sub>4</sub>, and concentrated in vacuo. Preparative TLC (development with ethyl acetate:methanol 20:1) afforded 10.0 mg (77% yield) of streptolic acid (2).  $[\alpha]_{D}^{24}$  = +135.0 (c = 0.7, EtOH); <sup>1</sup>H NMR (500 MHz, CDCl<sub>3</sub>)  $\delta$  0.69 (d, 3H, J =7.0 Hz), 1.04 (d, 3H, J = 7.0 Hz), 1.23 (s, 3H), 1.80 (s, 3H), 1.94 (m, 1H), 2.73 (m, 1H), 2.81 (d, 1H, J = 5.0 Hz), 2.98 (d, 1H, J = 5.0 Hz), 3.63 (d, 1H, J = 10.5 Hz), 4.35 (t, 1H, J = 4.5 Hz), 5.62 (d, 1H, J = 10.0 Hz), 5.81 (d, 1H, J = 15.5 Hz), 6.13 (d, 1H, J = 10.0 Hz), 6.34 (dd, 1H, J = 10.0, 5.0 Hz), 7.44 (d, 1H, J = 15.5 Hz); <sup>13</sup>C NMR (125 MHz, CDCl<sub>3</sub>)  $\delta$ 12.2, 12.5, 17.1, 22.2, 33.7, 35.0, 50.5, 55.0, 71.4, 76.1, 98.8, 115.1, 130.5, 132.6, 133.8, 143.3, 152.0, 172.5; HRMS (ESI) calculated for C<sub>18</sub>H<sub>25</sub>O<sub>5</sub>  $[M + H]^+$  321.1702, found 321.1702.

Streptolydiginone (4). A stirred solution of ketoamide 44 (16.0 mg, 0.033 mmol) in methanol (0.7 mL) was treated with sodium methoxide (0.33 mL of 0.5 M solution in methanol, 0.165 mmol) at 0 °C. The reaction mixture was stirred overnight at room temperature and concentrated in vacuo. The residue was dissolved in water (1 mL), treated with aqueous HCl (0.17 mL of 1 N solution) at 0 °C, and extracted twice with ethyl acetate. The combined organic extracts were washed with water, brine, then dried over MgSO4 and concentrated in vacuo to afford 14.0 mg (88% yield) of streptolydiginone (4).  $[\alpha]^{25}_{D}$  = +24.9 (c = 0.3, CHCl<sub>3</sub>); <sup>1</sup>H NMR (500 MHz, CDCl<sub>3</sub>)  $\delta$  0.70 (m, 3H), 1.05 (m, 6H), 1.23 (s, 3H), 1.90 (s, 3H), 1.93 (m, 1H), 2.73-2.91 (m, 6H), 2.98 (d, 1H, J = 5.0 Hz), 3.65 (d, 1H, J = 10.5 Hz), 4.10 (s, 1H), 4.35 (t, 1H, J = 4.5 Hz), 5.63 (d, 1H, J = 10.1 Hz), 5.67 (s, 1H), 6.26 (d, 1H, J = 9.9 Hz), 6.34 (dd, 1H, J = 10.1, 4.7 Hz), 7.10 (d, 1H, J = 15.6 Hz), 7.58 (d, 1H, J = 15.6 Hz); <sup>13</sup>C NMR (125 MHz, CDCl<sub>3</sub>)  $\delta$  11.4, 12.2, 12.5, 17.1, 22.2, 26.4, 34.1, 35.2, 40.9, 50.5, 55.0, 62.9, 71.4, 76.1, 98.9, 100.3, 116.2, 130.6, 133.8, 133.9, 146.0, 150.3, 175.0, 175.5, 175.7, 194.0; HRMS (ESI) calculated for  $C_{26}H_{35}N_2O_7 [M + H]^+$  487.2444, found 487.2442. A portion of the product was treated with dilute aqueous NaHCO<sub>3</sub>, concentrated in vacuo, and azeotroped three times with chloroform. The residue was extracted with chloroform, and the solution phase was concentrated in vacuo and azeotroped with chloroform to afford a sodium salt of streptolydiginone. <sup>1</sup>H NMR (500 MHz,  $(CD_3)_2SO$   $\delta$  0.65 (d, 3H, J = 6.9 Hz), 0.78 (d, 3H, J = 6.9 Hz), 0.98 (d, 3H, J = 6.8 Hz), 1.08 (s, 3H), 1.78 (s, 3H), 1.78 (m, 1H), 2.57 (d, 3H, *J* = 4.2 Hz), 2.66 (q, 1H, *J* = 6.9 Hz), 2.75 (m, 1H), 2.89 (d, 1H, *J* = 5.0 Hz), 2.94 (d, 1H, J = 5.0 Hz), 3.58 (m, 2H), 4.10 (s, 1H), 4.33 (t, 1H, J = 4.6 Hz), 5.62 (d, 1H, J = 10.1 Hz), 5.76 (d, 1H, J = 9.7 Hz), 6.08 (s, 1H), 6.39 (dd, 1H, J = 10.0, 4.6 Hz), 7.01 (d, 1H, J = 15.6 Hz), 7.67 (d, 1H, J = 15.6 Hz), 7.84 (q, 1H, J = 4.3 Hz); <sup>13</sup>C NMR (125 MHz, (CD<sub>3</sub>)<sub>2</sub>SO)  $\delta$ 10.5, 12.2, 12.6, 17.3, 22.3, 25.6, 32.9, 34.7, 40.3, 49.8, 54.8, 60.4, 70.4, 75.9, 98.2, 102.0, 126.8, 130.2, 133.6, 134.0, 137.5, 141.1, 175.0, 176.4, 181.4, 193.9.

**Streptolydigin (1).** A stirred solution of phosphonate **49** (30.0 mg, 0.05 mmol) in THF (0.4 mL) was treated with potassium *tert*-butoxide (0.15 mL of 1 M solution in THF, 0.15 mmol) at 0 °C. After 40 min, aldehyde **11** (7.0 mg, 0.0252 mmol) was added in one portion. The reaction was warmed to room temperature and left overnight. THF (0.5 mL) was added, and the reaction was quenched with aqueous HCI (0.18 mL of 1 M solution, 0.18 mmol, slight excess with respect to potassium *tert*-butoxide) at 0 °C. After 1 h, ethyl acetate and water were

added. Phases were separated, and the organic phase was washed three times with water and once with brine, dried over MgSO4, and concentrated in vacuo. Purification by flash chromatography on silica gel (chloroform, followed by the elution with chloroform:methanol 100:3) afforded 9.0 mg (60% yield) of streptolydigin (1). <sup>1</sup>H NMR (500 MHz,  $CDCl_3$ )  $\delta$  0.70 (d, 3H, J = 6.9 Hz), 1.05 (d, 3H, J = 6.9 Hz), 1.11 (m, 6H), 1.23 (s, 3H), 1.77 (m, 1H), 1.90 (s, 3H), 1.94 (m, 1H), 2.07 (d, 1H, *J* = 13.3 Hz), 2.50 (q, 1H, *J* = 11.7 Hz), 2.78 (m, 1H), 2.81 (d, 1H, *J* = 5 Hz), 2.89 (d, 3H, J = 4.4 Hz), 2.98 (d, 1H, J = 5 Hz), 3.05 (q, 1H, J = 6.9 Hz), 3.41 (s, 1H), 3.65 (m, 2H), 4.35 (t, 1H, J = 4.6 Hz), 4.86 (s, 1H), 5.59 (d, 1H, J = 11.2 Hz), 5.62 (d, 1H, J = 10.1 Hz), 5.85 (m, 1H), 6.25 (d, 1H, J = 9.9 Hz), 6.34 (dd, 1H, J = 10.1, 4.8 Hz), 7.16 (d, 1H, J = 15.6 Hz), 7.57 (d, 1H, J = 15.6 Hz); <sup>13</sup>C NMR (125 MHz, CDCl<sub>3</sub>)  $\delta$  10.1, 12.2, 12.5, 17.1, 17.2, 21.2, 22.2, 26.8, 30.2, 34.1, 35.2, 42.1, 50.5, 55.0, 62.8, 66.3, 71.4, 76.1, 76.5, 78.9, 98.9, 99.7, 116.2, 130.5, 133.8, 134.0, 146.1, 150.5, 173.7, 175.0, 175.2, 193.9; HRMS (ESI) calculated for  $C_{32}H_{45}N_2O_9[M + H]^+$  601.3125, found 601.3121. The optical rotation of synthetic sample of **41** was determined to be  $[\alpha]^{22}_{D} = -16.2$  (*c* = 0.7, CHCl<sub>3</sub>), which was in good agreement with that of an authentic sample of streptolydigin obtained from ChemCon GmbH.

Adamantane-Containing Tetramic Acid (56). A stirred solution of phosphonate 49 (23.0 mg, 0.039 mmol) in THF (0.3 mL) was treated with potassium tert-butoxide (0.12 mL of 1 M solution in THF, 0.12 mmol) at 0 °C. The resulting solution was stirred 30 min and transferred into a solution of aldehyde 55 (6.0 mg, 0.0259 mmol) in THF (0.1 mL) at 0 °C. The reaction was warmed to room temperature and left overnight. THF (0.4 mL) was added, and the reaction was quenched with aqueous HCl (0.15 mL of 1 M solution, 0.15 mmol, slight excess with respect to potassium tert-butoxide) at 0 °C. After 1 h, ethyl acetate and water were added. Phases were separated, and the organic phase was washed three times with water and once with brine, dried over MgSO<sub>4</sub>, and concentrated in vacuo. Purification by flash chromatography on silica gel (chloroform, followed by the elution with chloroform: methanol 100:3) afforded 8.5 mg (59% yield) of 56.  $[\alpha]^{25}_{D} = -96.4$  $(c = 0.8, \text{CHCl}_3);$  <sup>1</sup>H NMR (500 MHz, CDCl<sub>3</sub>)  $\delta$  0.90 (d, 3H, J = 7.0 Hz), 1.10 (m, 6H), 1.46 (d, 3H, J = 11.5 Hz), 1.54 (d, 3H, J = 11.5 Hz), 1.60 (d, 3H, J = 12.0 Hz), 1.67 (d, 3H, J = 12.0 Hz), 1.77 (m, 1H), 1.88 (s, 3H), 1.95 (s, 3H), 2.04 (m, 1H), 2.23 (m, 1H), 2.50 (q, 1H, J = 12.0 Hz), 2.89 (d, 3H, J = 4.5 Hz), 3.05 (q, 1H, J = 7.0 Hz), 3.40 (s, 1H), 3.64 (m, 1H), 4.86 (s, 1H), 5.57 (d, 1H, J = 11.0 Hz), 5.85 (s, 1H), 6.05 (d, 1H, *J* = 10.5 Hz), 7.12 (d, 1H, *J* = 15.5 Hz), 7.55 (d, 1H, *J* = 15.5 Hz); <sup>13</sup>C NMR (125 MHz, CDCl<sub>3</sub>) δ 10.1, 12.5, 13.3, 17.2, 21.2, 26.8, 28.7, 30.2, 36.0, 37.2, 39.8, 42.1, 44.0, 62.8, 66.3, 76.5, 78.9, 99.5, 115.5, 133.6, 150.6, 151.2, 173.8, 175.1, 175.4, 193.8.

10,11-Dihydrostreptolydigin (63). A stirred solution of phosphonate 49 (23.6 mg, 0.04 mmol) in THF (0.3 mL) was treated with potassium tert-butoxide (0.12 mL of 1 M solution in THF, 0.12 mmol) at 0 °C. The resulting solution was stirred 30 min and transferred into a solution of aldehyde 62 (7.5 mg, 0.0268 mmol) in THF (0.1 mL) at 0 °C. The reaction was warmed to room temperature and left overnight. THF (0.4 mL) was added, and the reaction was quenched with aqueous HCl (0.15 mL of 1 M solution, 0.15 mmol, slight excess with respect to potassium tert-butoxide) at 0 °C. After 1 h, ethyl acetate and water were added. Phases were separated, and the organic phase was washed three times with water and once with brine, dried over MgSO4, and concentrated in vacuo. Purification by flash chromatography on silica gel (chloroform, followed by the elution with chloroform:methanol 100:3) afforded 9.0 mg (56% yield) of 10,11-dihydrostreptolydigin (63).  $[\alpha]_{D}^{25} = -53.3 \ (c = 0.9, \text{CHCl}_3); {}^{1}\text{H NMR} \ (500 \text{ MHz}, \text{CDCl}_3)$  $\delta$  0.76 (d, 3H, J = 7.0 Hz), 1.07–1.12 (m, 9H), 1.22 (s, 3H), 1.73–1.82 (m, 4H), 1.90 (s, 3H), 1.93 (m, 1H), 2.04 (m, 1H), 2.14 (m, 1H), 2.51 (q, 1H, J = 12.0 Hz), 2.64 (d, 1H, J = 5.0 Hz), 2.81 (d, 1H, J = 5.0 Hz), 2.84 (m, 1H), 2.89 (d, 3H, J = 4.5 Hz), 3.05 (q, 1H, J = 7.0 Hz), 3.40 (s, 1H), 3.65 (m, 1H), 3.82 (d, 1H, J = 11.0 Hz), 3.98 (m, 1H), 4.86 (s, 1H), 5.59 (d, 1H, *J* = 11.0 Hz), 5.85 (d, 1H, *J* = 4.0 Hz), 6.24 (d, 1H, *J* = 10.0 Hz), 7.15 (d, 1H, *J* = 15.5 Hz), 7.56 (d, 1H, *J* = 16.0 Hz); <sup>13</sup>C NMR (125 MHz, CDCl<sub>3</sub>)  $\delta$  10.1, 12.3, 12.7, 17.2, 21.0, 21.2, 22.5, 26.8, 27.2, 27.4, 30.2, 30.4, 35.1, 35.3, 42.1, 51.3, 58.5, 62.8, 66.3, 70.3, 76.5, 78.9, 98.2, 99.7, 116.1, 134.3, 146.3, 150.6, 173.7, 175.0.

**RNAP and Scaffolds.** For in vitro transcription experiments, recombinant *T. aquaticus* core RNAP was purified as described earlier.<sup>34</sup> To assemble a previously described<sup>35</sup> nucleic acid scaffold used in Figure 3 experiment, the following oligonucleotide strands were used: RNA (5'-GUAGCGGA-3'), template DNA (3'-CATCGCCTGTACA-TTTCAGACAGGACC-5'), and nontemplate DNA (5'-TGTAAAGT-CTGTCCTGG-3'). To assemble a scaffold with longer transcribed part used in Figure 5, the same RNA was combined with 3'-CATCGCCTG-TACATTTCAGACAGGACCAGGTGTTGGG-5' template DNA and 5'-TGTAAAGTCTGTCCTGGTCCACAACCC-3' nontemplate DNA. To visualize transcription products, RNA was <sup>32</sup>P-labeled at the 5'- end with T4 polynucleotide kinase before scaffold assembly.

Transcription Assays. Transcription elongation complexes were reconstituted by combining 200 nM RNAP and 50 nM nucleic acids scaffold in  $10 \,\mu$ L of transcription buffer (25 mM Tris-HCl, 50 mM NaCl, 5 mM MgCl<sub>2</sub>, 0.5 mM DTT, pH 7.9) and incubation at 37 °C for 10 min. Following the addition of streptolydigin derivatives and a 10 min incubation at 37 °C, transcription was initiated by the addition of NTPs at concentrations indicated in the figures. After a 3 min incubation at 37 °C, reactions were terminated by the addition of 1 volume of formamide loading buffer. After being heated at 100 °C for 1 min, samples were separated by 20% PAGE with 7 M Uurea, followed by PhosphorImager analysis. For single-nuceotide addition experiments, the concentration of CTP was 5  $\mu$ M, and reaction incubation temperature was 25 °C. Reaction incubation times varied and are indicated in Figure 5C. For data analysis, the results of single-nucleotide addition experiments were fitted to a single exponent described by equation:  $P = A^*(1 - \exp(-kt))$ , where *P* is the fraction of the 9-nt product, *A* is the fraction of the product at  $t \rightarrow \infty$ , and *k* and *t* are the rate and time of the transcription reaction. The fitting was performed by nonlinear regression using the program SigmaPlot (Version 8.0).

**Bacterial Strains and Media.** *Streptococcus salivarius* Andrewes and Horder was obtained from ATCC (9758) and was grown in Bacto Tryptic Soy Broth Medium at  $37 \,^{\circ}$ C.

**Antibacterial Testing.** The antibiotic activity of all compounds was analyzed via disk diffusion assay. Several drops of *S. salivarius* culture were spread in a Petri dish containing Difco Tryptic Soy Agar. Paper disks of 5 mm diameter were used. Each disk was treated with a solution of each individual compound dissolved in ethanol  $(2-15 \,\mu\text{L}$  depending on the sample). Following evaporation of the ethanol, the disks were placed on top of agar. Plates were incubated at 37 °C for 24 h.

## ASSOCIATED CONTENT

**Supporting Information.** Detailed experimental procedures as well as analytical and spectral characterization data for all new compounds. This material is available free of charge via the Internet at http://pubs.acs.org.

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