

Cite this: *Chem. Sci.*, 2022, 13, 12776

All publication charges for this article have been paid for by the Royal Society of Chemistry

Ten-step asymmetric total syntheses of potent antibiotics anthracimycin and anthracimycin B†

Peilin Tian,^a Wenkang Ye,^b Xiayan Zhang,^a Yi Tong,^a Pei-Yuan Qian^{ID} *^{bc} and Rongbiao Tong^{ID} *^{ac}

The increase in antibiotic resistance calls for the development of novel antibiotics with new molecular structures and new modes of action. However, in the past few decades only a few novel antibiotics have been discovered and progressed into clinically used drugs. The discovery of a potent anthracimycin antibiotic represents a major advance in the field of antibiotics. Anthracimycin is a structurally novel macrolide natural product with an excellent biological activity profile: (i) potent *in vitro* antibacterial activity (MIC 0.03–1.0 $\mu\text{g mL}^{-1}$) against many methicillin-resistant *Staphylococcus aureus* (MRSA) strains, *Bacillus anthracis* (anthrax), and *Mycobacterium tuberculosis*; (ii) low toxicity to human cells ($\text{IC}_{50} > 30 \mu\text{M}$); (iii) a novel mechanism of action (inhibiting DNA/RNA synthesis). While the first total synthesis of anthracimycin was elegantly accomplished by Brimble *et al.* with 20 steps, we report a 10-step asymmetric total synthesis of anthracimycin and anthracimycin B (first total synthesis). Our convergent strategy features (i) one-pot sequential Mukaiyama vinylogous aldol/intramolecular Diels–Alder reaction to construct *trans*-decalin with high yield and excellent *endo/exo* selectivity and (ii) Z-selective ring-closing metathesis to forge the 14-membered ring. *In vitro* antibacterial evaluation suggested that our synthetic samples exhibited similar antibacterial potency to the naturally occurring anthracimycins against Gram-positive strains. Our short and reliable synthetic route provides a supply of anthracimycins for further in-depth studies and allows medicinal chemists to prepare a library of analogues for establishing structure–activity relationships.

Received 9th September 2022
Accepted 14th October 2022

DOI: 10.1039/d2sc05049h

rsc.li/chemical-science

Introduction

The alarming prevalence of bacteria resistant to multiple antibiotics raises a global concern on the availability of new antibiotics effective against such “superbugs” that are resistant to most or all known antibiotics.^{1,2} The search for new antibiotics with different strategies including bioassay-guided isolation of new antibacterial natural products, genome mining,^{3–8} and chemical synthesis/modifications of existing antibiotics^{9,10} continues to play an important role in addressing antibiotic resistance.^{11–14} However, only a few structurally novel antibacterial compounds have been discovered in the past five decades, and anthracimycin¹⁵ was one of these new antibiotics with a novel molecular structure and high antibacterial potency. Anthracimycin was discovered in 2013 by Fenical and co-worker

from marine microorganism *Streptomyces* species (CNH365) to be a structurally novel potent antibiotic against Gram-positive bacteria (MIC 0.03–0.5 $\mu\text{g mL}^{-1}$) including notorious methicillin-resistant *Staphylococcus aureus* (MRSA)¹⁶ and the bacterium *Bacillus anthracis* that has been used as a bioterrorism weapon to cause human-infectious anthrax (Fig. 1). In 2018, Reyes *et al.*¹⁷ identified 2-demethyl anthracimycin (anthracimycin B) from cultures of the deep-sea actinomycete *Streptomyces cyaneofuscantus* M-169 and found that it exhibited similar but less potent antibacterial activity against Gram-positive bacteria. Another anthracimycin analogue (anthracimycin BII-2619) was also identified in 2018 by Sirota *et al.*¹⁸ through a genome mining strategy to display Gram-positive antibacterial activity with lower mammalian cytotoxicity compared to that of anthracimycin. In 2020, Fukuda¹⁹ and co-workers isolated 2-epi-anthracimycin along with anthracimycin and evaluated its cytotoxicity against Jurkat cells ($\text{IC}_{50} = 50.5 \mu\text{M}$). While most macrolide antibiotics bind to the 50S ribosomal subunit to inhibit protein biosynthesis,²⁰ it was suggested that anthracimycin exerts its bactericidal effect through inhibiting DNA and/or RNA synthesis in the absence of DNA intercalation.²¹ The high antibacterial potency, new molecular structure, and novel mechanism of action make anthracimycin an ideal target for chemical synthesis and development of novel antibiotics.

^aDepartment of Chemistry, The Hong Kong University of Science and Technology, Clearwater Bay, Kowloon, Hong Kong, China. E-mail: rtong@ust.hk; Fax: +86 23581594; Tel: +86 23587357

^bDepartment of Ocean Science, The Hong Kong University of Science and Technology, Clear Water Bay, Kowloon, Hong Kong, China. E-mail: boqianpy@ust.hk

^cThe Southern Marine Science and Engineering Guangdong Laboratory (Guangzhou), Nansha, Guangzhou 511458, China

† Electronic supplementary information (ESI) available. CCDC 2175241 and 2194803. For ESI and crystallographic data in CIF or other electronic format see DOI: <https://doi.org/10.1039/d2sc05049h>



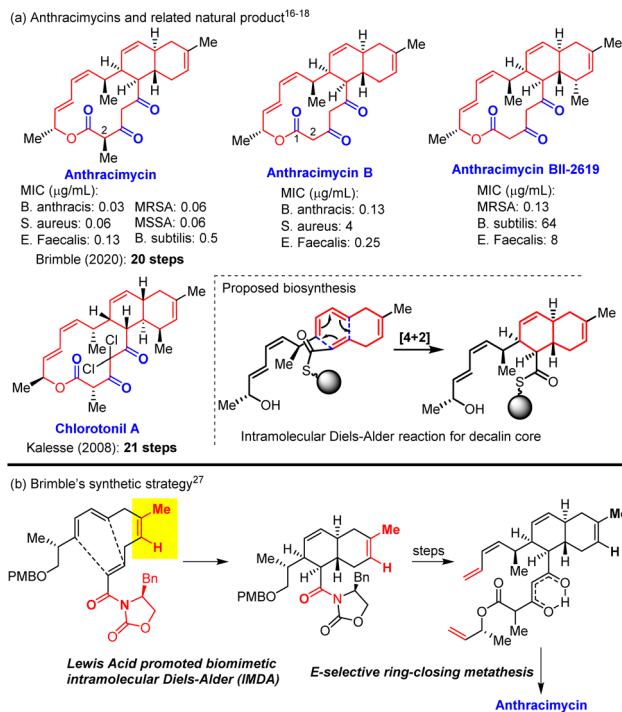


Fig. 1 Anthracimycin natural products.

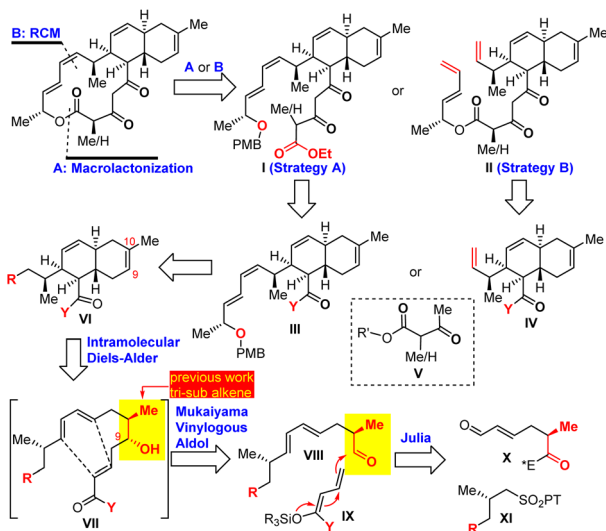
In 2019, two PhD theses (J. L. Freeman^{22,23} and G. Lodovici²⁴) disclosed detailed synthetic studies towards anthracimycin and presented a significant synthetic challenge of using the biomimetic intramolecular Diels–Alder (IMDA) reaction of the tetraene substrate (Scheme 1b)^{25,26} to construct the decalin core. Further optimization of Freeman's thesis work enabled Brimble²⁷ and co-workers to achieve the first total synthesis of anthracimycin in 20 steps with 0.4 mg pure anthracimycin. It was noted that Kalesse *et al.*²⁸ accomplished the total synthesis of structurally related chlorotoniil A²⁹ in 21 steps by using the

halogen-directed IMDA reaction and macrolactonization as the key steps. In this paper, we report a 10-step asymmetric total synthesis of anthracimycin and anthracimycin B (the first total synthesis) with substantially improved overall yield (>3% overall yield, 25 mg and 14 mg obtained, respectively).

Results and discussion

The key synthetic challenge of anthracimycin in the previous synthetic studies^{22–24,27} was the inefficient construction of the *trans*-decalin core through the biomimetic intramolecular Diels–Alder (IMDA) reaction of the tetraene substrate. To address this challenge, we proposed an IMDA reaction of an “unusual” triene substrate VII with a C9 secondary alcohol (Scheme 1), which served (1) as the precursor of the trisubstituted *Z*- Δ alkene, (2) to enhance the *endo/exo* selectivity and diastereoselectivity, and (3) to improve the convergency of the synthesis because the presence of the alcohol would allow the fragment union through C–C bond formation with many well-established transformations (*i.e.*, aldol, allylation, Grignard addition, *etc.*). With this central ideal in mind, our retrosynthetic analysis was conceived as depicted in Scheme 1. The 14-membered macrocycle would be forged in the last step by either macrolactonization²⁸ of I or *Z*-selective ring-closing metathesis (RCM) of II. Notably, *Z*-selective RCM might be advantageous since the preparation of the more challenging *Z*-configured alkene substrate at an early stage of synthesis was eliminated. Both I and II can be obtained by condensation of 1,3-dicarbonyl V with III and IV, respectively. The designed triene VII substrate for the IMDA reaction could be easily synthesized by Julia–Kocienski olefination³⁰ of aldehyde X and tetrazole XI followed by the Mukaiyama vinylogous aldol (MVA)^{31–35} reaction of aldehyde VIII and *O,O*-silyl ketene acetal IX.³⁶

Our synthesis consisted of two stages: *trans*-decalin synthesis and macrolactone synthesis as depicted in Scheme 2. We started with the preparation of known compound 2 (ref. 37 and 38) in 86% yield by Evans asymmetric alkylation³⁹ of commercially available (*S*)-4-benzyl-3-propionyl-2-oxazolidinone (1). Olefin cross metathesis of 2 with crotonaldehyde delivered 3 as an exclusive *E*-configuration in 85% yield. Julia–Kocienski olefination of *N*-phenyl tetrazole sulfone 4a (ref. 22 and 40) with aldehyde 3 followed by Dess–Martin periodinane (DMP) oxidation provided (*E,E*)-diene 5a (*E/Z* > 10 : 1), which was subjected to our planned Lewis acid-promoted Mukaiyama vinylogous aldol (MVA) reaction with the known silyl ketene acetal 6.^{36,41} Triarylborane [B(C₆F₅)₃]⁴² was found to effectively promote MVA to provide triene 7 in 71% yield with a 6 : 1 diastereomeric ratio. The thermal IMDA reaction of triene 7 was first attempted at 165 °C in dichlorobenzene to produce a mixture of *endo/exo* products (*endo/exo*: 2.3 : 1) with 82% combined yield. To improve the *endo/exo* selectivity, we used BF₃–Et₂O to promote the IMDA reaction,²⁹ which resulted in decomposition. Fortunately, we found that Et₂AlCl (4.0 equiv.) was an effective promoter for the IMDA reaction to provide the desired *trans*-decalin as a mixture of 8a and 8b, which upon treatment with 37% HCl for removal of the TBS protecting group as well as lactonization generated



Scheme 1 Retrosynthetic analysis of anthracimycins.



and partial hydrogenation with the Lindlar catalyst. Aldehyde **11** was converted into alkyne **12a** with the Bestmann reagent, and then Sonogashira coupling with vinyl iodide **13a** followed by Lindlar hydrogenation provided the desired (*E,Z*)-diene **14a** in 64% yield over two steps. The installation of 1,3,5-tricarbonyls (**14a** → **16a**) was challenging: Claisen condensation of ester **14a** with 1,3-dicarbonyl **15a** was not successful, while a four-step sequence (DIBAL-H reduction, DMP oxidation, Aldol reaction, and DMP oxidation) delivered **16a** with 27% yield. Nevertheless, with tens of milligrams of **16a** in hand, we explored the BF₃·Et₂O-mediated macrolactonization, but we were not able to identify any desired product from many attempted reactions under various conditions (temperature and equivalent of BF₃·Et₂O). TLC and NMR analysis clearly suggested that **16a** decomposed in the presence of BF₃·Et₂O. Then, we investigated the second macrocyclization strategy: RCM. Because *Z*-selective Stork–Zhao olefination failed to produce *Z*-vinyl iodide for subsequent Stille coupling, we decided to take a risk: *Z*-alkene formation by RCM as the last step, which contrasts with the previous RCM for *E*-alkene formation.²⁷ To verify this macrocyclization method, we prepared two substrates **16b** and **16c**, corresponding to anthracimycin and anthracimycin B, respectively. Wittig olefination of **11** followed by ester reduction with DIBAL-H and DMP oxidation afforded aldehyde **13b** with 46% yield over 3 steps. The subsequent aldol reaction of **13b** with dienolate of 1,3-dicarbonyl **14b/14c** followed by DMP oxidation provided the key macrocyclization substrates **16b** and **16c** in 42% and 44% yield over two steps, respectively. The planned RCM of **16b** and **16c** proceeded smoothly with the Hoveyda–Grubbs II catalyst to furnish anthracimycin and anthracimycin B (exclusive *Z*-double bond), respectively, with >60% yield based on the crude NMR analysis (42–43% isolated yield). The NMR data of our synthetic samples were consistent with those reported for natural products (see ESI Tables S1–S4†). Additionally, single crystals of anthracimycin were obtained for X-ray diffraction analysis, which further confirmed the structure of our synthetic anthracimycin.

Since our total synthesis required 14 steps, we set out to further optimize our synthetic route (Scheme 3). First, we employed *N*-phenyl tetrazole sulfone **4b**,⁴⁷ instead of **4a** (Scheme 2), for Julia–Kocienski olefination with aldehyde **3** so that the late-stage conversion of lactone to vinyl ester (**9** → **12b**, Scheme 2) was unnecessary (and thus saving us 4 steps). Reductive removal of the Evans chiral auxiliary and DMP oxidation of the resulting alcohol afforded aldehyde **5b** (62% yield) with 6 : 1 *E/Z* selectivity for the newly-formed alkene. Next, we explored the possibility of one-pot sequential Mukaiyama vinylogous aldol (MVA)/intramolecular Diels–Alder (IMDA) reaction since both involved a Lewis acid as a promoter. After examining several acid promoters including TiCl₄, BF₃·Et₂O, CBS-HOTf, proline, Rh(I) and B(C₆F₅)₃, we found that the sequential MVA and IMDA reactions could be achieved in one pot by treating the reaction mixture first with B(C₆F₅)₃ and then with Et₂AlCl to provide a mixture of *trans*-decalins **8c** and **8d** (63% combined yield on a 1.1 g scale) with a 3 : 1 diastereomeric ratio in favor of the desired **8c**. Although the diastereoselectivity of the IMDA reaction was moderate as compared to substrate **7** probably due to the lack of a macrocyclic intermediate (or a transition state) for the highly diastereoselective transannular Diels–Alder reaction, the overall efficiency was much higher. Dehydration with the Burgess reagent allowed DIBAL-H reduction of the ester to be carried out in one pot. DMP oxidation of the resulting alcohol provided aldehyde **13b** (66% yield over two steps), which was elaborated to anthracimycin and anthracimycin B following the same protocol as in Scheme 2. The optimized route for **13b** allowed us to accomplish the total synthesis of anthracimycin and anthracimycin B in 10 steps with 3.3–3.6% isolated overall yield from commercially available Evans chiral auxiliary **1** without using protecting groups and with only 7 isolated intermediates. The concise and efficient route enabled us to prepare 25 mg of anthracimycin and 14 mg of anthracimycin B, which allowed us to evaluate their antibacterial activity.

It is well recognized that the bioactivity of newly isolated natural products might be different from that of the corresponding synthetic samples (possibly due to contamination of natural

Table 1 Antimicrobial and cytotoxic activities^a

Strains/cells	Ant	Ant B	Van
<i>A. baumannii</i> : B65371	>40	>40	>40
<i>E. cloacae</i> : NRRL-B-425	>40	>40	>40
<i>E. coli</i> : K12	>40	>40	>40
<i>K. pneumoniae</i> : NRRL-B-408	>40	>40	>40
MRSA: ATCC43300	0.03 ± 0.001	0.7 ± 0.13	0.8 ± 0.04
<i>S. aureus</i> : ATCC 25923	0.04 ± 0.004	0.3 ± 0.023	0.8 ± 0.002
MRSA ATCC29213	0.04 ± 0.004	1.0 ± 0.11	0.4 ± 0.045
MRSA ATCC29213 Biofilm (MBIC)	0.02 ± 0.006	0.6 ± 0.19	1.2 ± 0.094
MRSA Sa115	0.04 ± 0.005	1.0 ± 0.094	0.4 ± 0.038
<i>S. aureus</i> R2952	0.04 ± 0.047	1.0 ± 0.094	0.4 ± 0.023
<i>B. subtilis</i> : zk31	0.04 ± 0.005	0.5 ± 0.075	0.07 ± 0.007
<i>M. luteus</i> : ATCC 10040	0.02 ± 0.009	0.8 ± 0.024	1.6 ± 0.094
HaCaT (IC ₅₀)	14.74 ± 1.32	14.90 ± 1.93	

^a The experimental results are expressed as the MIC, MBIC and IC₅₀ (μg mL⁻¹). All the bioactivity assays of compounds were performed three times. In antimicrobial experiments, kanamycin and vancomycin were used as positive controls. Ant: anthracimycin; Ant B: anthracimycin B; Van: vancomycin; HaCaT cells are human keratinocytes.



samples). Therefore, it is imperative to verify their bioactivity using synthetic samples. The antibacterial activity of our synthetic anthracimycin and anthracimycin B was evaluated against several pathogenic strains, including Gram-negative strains: *A. baumannii* B-65371, *E. cloacae* NRRL-B-425, *E. coli* k12, and *K. pneumoniae* NRRL-B-408, and Gram-positive strains: MRSA ATCC 43300, *S. aureus* ATCC 25923, *B. subtilis* zk31 and *M. luteus* ATCC 10040. As shown in Table 1, synthetic anthracimycin and anthracimycin B were not effective against Gram-negative strains but they were potent against the Gram-positive bacteria (MIC: 0.03–0.8 $\mu\text{g mL}^{-1}$), which is consistent with MIC values reported for natural anthracimycins.^{15,17,18}

A cytotoxicity study using HaCaT (cell line from human skin) suggested that both synthetic anthracimycins possessed low cytotoxicity ($\text{IC}_{50} > 14 \mu\text{g mL}^{-1}$), which holds great promise for further drug development. More interestingly, we found that both anthracimycins significantly inhibited the MRSA biofilm formation (MRSA ATCC29213) with MBIC values (MBIC 0.02 $\mu\text{g mL}^{-1}$ and 0.6 $\mu\text{g mL}^{-1}$, respectively) even lower than the respective MIC values. As compared to vancomycin, a frontline antibiotic for treatment of MRSA infections, anthracimycin was 60 times more potent. This result is significant because MRSA biofilm formation^{48–51} is primarily responsible for the reduced efficacy or inefficacy of many clinically used antibiotics for MRSA infections, and anthracimycin demonstrated potential as a new effective anti-MRSA antibiotic.

Conclusion

In summary, we have accomplished a 10-step asymmetric total synthesis of anthracimycin and anthracimycin B (the first total synthesis) with 3.3–3.6% overall isolated yields. The key strategy is the use of alcohol as the latent alkene to enable convergent fragment coupling (vinylogous Mukaiyama aldol) and to enhance the *endo/exo* selectivity and yield of the intramolecular Diels–Alder reaction. Additionally, the ring-closing metathesis using the Hoveyda–Grubbs II catalyst allowed the efficient construction of a 14-membered macrolide with exclusive *Z*-alkene selectivity. Our short and efficient synthetic route enabled us to prepare 25 mg of anthracimycin and 14 mg of anthracimycin B and allowed us to evaluate their antibacterial activities using our synthetic sample, which were consistent with those reported for the natural anthracimycins. Finally, we found that our synthetic anthracimycin significantly inhibited the MRSA biofilm formation with great promise to be a new effective anti-MRSA antibiotic.

Data availability

Experimental procedures and characterization data are available within this article and its ESI.† Data are also available from the corresponding author on request.

Author contributions

P. T., X. Z. and Y. T. performed the synthetic experiments. W. Y. performed the biological experiments. P. Q. and R. T.

conceptualized and directed the project, and drafted the manuscript with the assistance from all co-authors.

Conflicts of interest

The authors declare no competing interests.

Acknowledgements

This research was financially supported by the Southern Marine Science and Engineering Guangdong Laboratory (Guangzhou) (SMSEGL20Sc01-B) and Research Grant Council of Hong Kong (C6026-19G, 16307219, 16304618, and 16303617). Prof. Wei Shen Aik and Prof. Jun (Joelle) Wang from Hong Kong Baptist University are gratefully acknowledged for X-ray crystallographic analysis.

Notes and references

- 1 C. L. Ventola, *Pharm. Ther.*, 2015, **40**, 277–283.
- 2 S. Baker, *Science*, 2015, **347**, 1064–1066.
- 3 E. J. Culp, N. Waglechner, W. Wang, A. A. Fiebig-Comyn, Y.-P. Hsu, K. Koteva, D. Sychantha, B. K. Coombes, M. S. Van Nieuwenhze, Y. V. Brun and G. D. Wright, *Nature*, 2020, **578**, 582–587.
- 4 G. L. Challis, *J. Med. Chem.*, 2008, **51**, 2618–2628.
- 5 Y.-X. Li, Z. Zhong, W.-P. Zhang and P.-Y. Qian, *Nat. Commun.*, 2018, **9**, 3273.
- 6 E. C. O'Neill, M. Schorn, C. B. Larson and N. Millán-Aguíñaga, *Crit. Rev. Microbiol.*, 2019, **45**, 255–277.
- 7 L. Foulston, *Curr. Opin. Microbiol.*, 2019, **51**, 1–8.
- 8 N. Ziemert, M. Alanjary and T. Weber, *Nat. Prod. Rep.*, 2016, **33**, 988–1005.
- 9 P. M. Wright, I. B. Seiple and A. G. Myers, *Angew. Chem., Int. Ed.*, 2014, **53**, 8840–8869.
- 10 S. E. Rossiter, M. H. Fletcher and W. M. Wuest, *Chem. Rev.*, 2017, **117**, 12415–12474.
- 11 A. Luther, M. Urfer, M. Zahn, M. Müller, S.-Y. Wang, M. Mondal, A. Vitale, J.-B. Hartmann, T. Sharpe, F. L. Monte, H. Kocherla, E. Cline, G. Pessi, P. Rath, S. M. Modaresi, P. Chiquet, S. Stiegeler, C. Verbree, T. Remus, M. Schmitt, C. Kolopp, M.-A. Westwood, N. Desjonquères, E. Brabet, S. Hell, K. LePoupon, A. Vermeulen, R. Jaisson, V. Rithié, G. Upert, A. Lederer, P. Zbinden, A. Wach, K. Moehle, K. Zerbe, H. H. Locher, F. Bernardini, G. E. Dale, L. Eberl, B. Wollscheid, S. Hiller, J. A. Robinson and D. Obrecht, *Nature*, 2019, **576**, 452–458.
- 12 T. F. Durand-Reville, A. A. Miller, J. P. O'Donnell, X. Wu, M. A. Sylvester, S. Guler, R. Iyer, A. B. Shapiro, N. M. Carter, C. Velez-Vega, S. H. Moussa, S. M. McLeod, A. Chen, A. M. Tanudra, J. Zhang, J. Comita-Prevoir, J. A. Romero, H. Huynh, A. D. Ferguson, P. S. Horanyi, S. J. Mayclin, H. S. Heine, G. L. Drusano, J. E. Cummings, R. A. Slayden and R. A. Tommasi, *Nature*, 2021, **597**, 698–702.
- 13 M. J. Mitcheltree, A. Pisipati, E. A. Syroegin, K. J. Silvestre, D. Klepacki, J. D. Mason, D. W. Terwilliger, G. Testolin, A. R. Pote, K. J. Y. Wu, R. P. Ladley, K. Chatman,



- A. S. Mankin, Y. S. Polikanov and A. G. Myers, *Nature*, 2021, **599**, 507–512.
- 14 Z. Wang, B. Koirala, Y. Hernandez, M. Zimmerman, S. Park, D. S. Perlin and S. F. Brady, *Nature*, 2022, **601**, 606–611.
- 15 K. H. Jang, S.-J. Nam, J. B. Locke, C. A. Kauffman, D. S. Beatty, L. A. Paul and W. Fenical, *Angew. Chem. Int. Ed.*, 2013, **52**, 7822–7824.
- 16 J. Larsen, C. L. Raisen, X. Ba, N. J. Sadgrove, G. F. Padilla-González, M. S. J. Simmonds, I. Loncaric, H. Kerschner, P. Apfalter, R. Hartl, A. Deplano, S. Vandendriessche, B. Černá Bolfiková, P. Hulva, M. C. Arendrup, R. K. Hare, C. Barnadas, M. Stegger, R. N. Sieber, R. L. Skov, A. Petersen, Ø. Angen, S. L. Rasmussen, C. Espinosa-Gongora, F. M. Aarestrup, L. J. Lindholm, S. M. Nykäsenoja, F. Laurent, K. Becker, B. Walther, C. Kehrenberg, C. Cuny, F. Layer, G. Werner, W. Witte, I. Stamm, P. Moroni, H. J. Jørgensen, H. de Lencastre, E. Cercenado, F. García-Garrote, S. Börjesson, S. Hæggman, V. Perreten, C. J. Teale, A. S. Waller, B. Pichon, M. D. Curran, M. J. Ellington, J. J. Welch, S. J. Peacock, D. J. Seilly, F. J. E. Morgan, J. Parkhill, N. F. Hadjirin, J. A. Lindsay, M. T. G. Holden, G. F. Edwards, G. Foster, G. K. Paterson, X. Didelot, M. A. Holmes, E. M. Harrison and A. R. Larsen, *Nature*, 2022, **602**, 135–141.
- 17 V. Rodríguez, J. Martín, A. Sarmiento-Vizcaíno, M. De La Cruz, L. García, G. Blanco and F. Reyes, *Mar. Drugs*, 2018, **16**, 406.
- 18 F. L. Sirota, F. Goh, K. N. Low, L. K. Yang, S. C. Crasta, B. Eisenhaber, F. Eisenhaber, Y. Kanagasundaram and S. B. Ng, *J. Genet. Genomics*, 2018, **6**, 63–73.
- 19 T. Fukuda, K. Nagai, A. Kanamoto and H. Tomoda, *J. Antibiot.*, 2020, **73**, 548–553.
- 20 J. W. Corcoran, *Mode of action and resistance mechanisms of macrolides*, Academic Press, Orlando, FL, 1984.
- 21 M. E. Hensler, K. H. Jang, W. Thienphrapa, L. Vuong, D. N. Tran, E. Soubih, L. Lin, N. M. Haste, M. L. Cunningham, B. P. Kwan, K. J. Shaw, W. Fenical and V. Nizet, *J. Antibiot.*, 2014, **67**, 549–553.
- 22 J. L. Freeman, M. A. Brimble and D. P. Furkert, *Org. Chem. Front.*, 2019, **6**, 2954–2963.
- 23 J. L. Freeman, PhD, University of Auckland, 2019.
- 24 G. Lodovici, PhD, University of York, 2019.
- 25 E. Harunari, H. Komaki and Y. Igarashi, *J. Antibiot.*, 2016, **69**, 403–405.
- 26 S. Alt and B. Wilkinson, *ACS Chem. Biol.*, 2015, **10**, 2468–2479.
- 27 E. K. Davison, J. L. Freeman, W. Zhang, W. M. Wuest, D. P. Furkert and M. A. Brimble, *Org. Lett.*, 2020, **22**, 5550–5554.
- 28 N. Rahn and M. Kalesse, *Angew. Chem., Int. Ed.*, 2008, **47**, 597–599.
- 29 K. Gerth, H. Steinmetz, G. Höfle and R. Jansen, *Angew. Chem., Int. Ed.*, 2008, **47**, 600–602.
- 30 B. Chatterjee, S. Bera and D. Mondal, *Tetrahedron: Asymmetry*, 2014, **25**, 1–55.
- 31 G. Casiraghi, F. Zanardi, G. Appendino and G. Rassu, *Chem. Rev.*, 2000, **100**, 1929–1972.
- 32 G. Casiraghi, L. Battistini, C. Curti, G. Rassu and F. Zanardi, *Chem. Rev.*, 2011, **111**, 3076–3154.
- 33 S. E. Denmark, J. R. Heemstra Jr and G. L. Beutner, *Angew. Chem., Int. Ed.*, 2005, **44**, 4682–4698.
- 34 M. Cordes and M. Kalesse, *Molecules*, 2019, **24**, 3040.
- 35 M. Kalesse, M. Cordes, G. Symkenberg and H.-H. Lu, *Nat. Prod. Rep.*, 2014, **31**, 563–594.
- 36 R. V. Hoffman and H. O. Kim, *J. Org. Chem.*, 1991, **56**, 1014–1019.
- 37 X. Xiong, Y. Wu and B. Liu, *Eur. J. Org. Chem.*, 2020, **2020**, 948–960.
- 38 C. W. Reed, M. G. Fulton, K. D. Nance and C. W. Lindsley, *Tetrahedron Lett.*, 2019, **60**, 743–745.
- 39 D. A. Evans, M. D. Ennis and D. J. Mathre, *J. Am. Chem. Soc.*, 1982, **104**, 1737–1739.
- 40 A. Fürstner, B. Fasching, G. W. O'Neil, M. D. B. Fenster, C. Godbout and J. Cecon, *Chem. Commun.*, 2007, 3045–3047, DOI: [10.1039/b707835h](https://doi.org/10.1039/b707835h).
- 41 S. Simsek, M. Horzella and M. Kalesse, *Org. Lett.*, 2007, **9**, 5637–5639.
- 42 M. Christmann and M. Kalesse, *Tetrahedron Lett.*, 2001, **42**, 1269–1271.
- 43 V. Koch, M. Nieger and S. Brase, *Org. Chem. Front.*, 2020, **7**, 2670–2681.
- 44 S. Poigny, S. Nouri, A. Chiaroni, M. Guyot and M. Samadi, *J. Org. Chem.*, 2001, **66**, 7263–7269.
- 45 G. Stork and K. Zhao, *Tetrahedron Lett.*, 1989, **30**, 2173–2174.
- 46 I. Janicki and P. Kielbasinski, *Adv. Synth. Catal.*, 2020, **362**, 2552–2596.
- 47 C. Jasper, R. Wittenberg, M. Quitschalle, J. Jakupovic and A. Kirschning, *Org. Lett.*, 2005, **7**, 479–482.
- 48 J.-O. Cha, J. I. Yoo, J. S. Yoo, H.-S. Chung, S.-H. Park, H. S. Kim, Y. S. Lee and G. T. Chung, *Osong Public Health Res. Perspect.*, 2013, **4**, 225–232.
- 49 A. Boudet, P. Sorlin, C. Pouget, R. Chiron, J.-P. Lavigne, C. Dunyach-Remy and H. Marchandin, *Front. Microbiol.*, 2021, **12**, 750489.
- 50 K. M. Craft, J. M. Nguyen, L. J. Berg and S. D. Townsend, *MedChemComm*, 2019, **10**, 1231–1241.
- 51 S. Cascioferro, D. Carbone, B. Parrino, C. Pecoraro, E. Giovannetti, G. Cirrincione and P. Diana, *ChemMedChem*, 2021, **16**, 65–80.

