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Total synthesis of landomycins Q and R and related core structures for exploration of the cytotoxicity and antibacterial properties†

Yao-Hsuan Lai,^{‡a} Soumik Mondal,^{‡ab} Hsin-Tzu Su,^{ab} Sheng-Cih Huang,^{ab} Mine-Hsine Wu,^c I.-Wen Huang,^c Tsai-Ling Yang Lauderdale,^c Jen-Shin Song,^c Kak-Shan Shia^c and Kwok-Kong Tony Mong^{id*ab}

Herein, we report the total synthesis of landomycins Q and R as well as the aglycone core, namely anhydrolandomycinone and a related core analogue. The synthesis features an acetate-assisted arylation method for construction of the hindered B-ring in the core component and a one-pot aromatization–deiodination–denbenzylation procedure to streamline the global functional and protecting group manipulation. Subsequent cytotoxicity and antibacterial studies revealed that the landomycin R is a potential antibacterial agent against methicillin-resistant *Staphylococcus aureus*.

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Introduction

Over the past few decades, the emergence of multidrug-resistant nosocomial pathogens has become a worldwide issue that is of concern for scientists and clinical practitioners.^{1,2} Consequently, new antibiotics and antibacterial strategies are urgently needed.^{3,4} However, the development of a new antibiotic is a slow process that is associated with a high risk of failure. Compounds that exhibit antimicrobial capacity *in vitro* may be unable to combat the infection *in vivo* or may be highly toxic to the host cells.⁵

Staphylococcus aureus (*S. aureus*) is a class of Gram-positive bacteria that is present on the skin and mucous membranes as a part of normal flora.⁶ If by chance *S. aureus* bacteria enter the bloodstream, they may cause severe infections including meningitides, endocarditis, and urinary tract infections.⁷ Staphylococcal infections are common in community and hospital-acquired settings and treatment of these infections has been complicated because of the rising incidence of methicillin resistant *S. aureus* (MRSA) infections.⁸ Although vancomycin is the antibiotic of choice, it may cause nephrotoxicity at high doses.^{9,10} Therefore, a less or non-toxic antibiotic is desirable.^{11–13}

The landomycins (LAs) are a class of angucyclines produced from *Streptomyces* species.¹⁴ The first report on LA described the isolation of LA A to D from *S. cyanogenus* S136.¹⁵ Through the use of the modern molecular biology techniques, the LA family has now grown to include more than 100 members.^{16,17} The general structure of LA compounds comprises a tetracyclic core, which is substituted with some hydroxyl groups and a 2-deoxysaccharide chain at the C8 position (Fig. 1a).^{14–17} Based on the B-ring structure and hydroxyl substitution pattern, four different core structures have been identified: tetrangulol, 5,6-anhydrolandomycinone, landomycinone, and 11-deoxylandomycinone. The combinations of different core structures and 2-deoxysaccharide chains provide a rich pool of natural products for structure and activity relationship studies.

Previous studies of LAs have mainly focused on the anti-cancer activity, and their potential antibacterial property has received far less attention.¹⁸ Given that the demand for non-toxic antibiotics is on the rise, we sought to establish a versatile synthetic route to procure the anhydrolandomycinone core (1), the B-ring unsaturated core analogue (2), the LA Q (3), and LA R (4) for exploration of the cytotoxic and antibacterial properties (Fig. 1b).

Results and discussion

Synthesis of anhydrolandomycinone (1), B-ring unsaturated core analogue (2), LA Q (3), and LA R (4)

The total synthesis of LA A and LA D has been reported by Yu and Yang, but the synthesis of LA Q (3) and LA R (4) with a different core structure has not yet been realized.¹⁹ From structural perspective, the synthesis of this class of compounds is non-trivial and yet challenging. Currently available protocols toward their aglycone core structure and glycone component

^aNational Chiao Tung University, Hsinchu 30010, Taiwan, Republic of China

^bNational Yang Ming Chiao Tung University, Hsinchu 30010, Taiwan, Republic of China. E-mail: tmong@nycu.edu.tw

^cNational Health Research Institutes, Miaoli County 35053, Taiwan, Republic of China

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‡ These authors contributed equally.



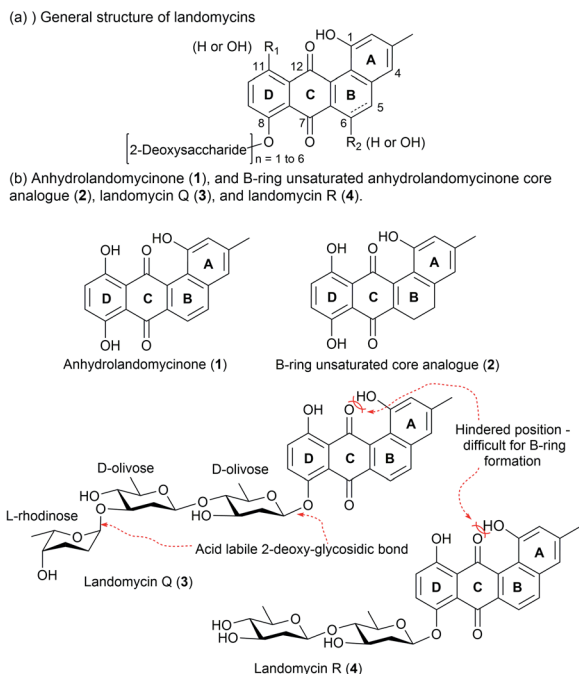
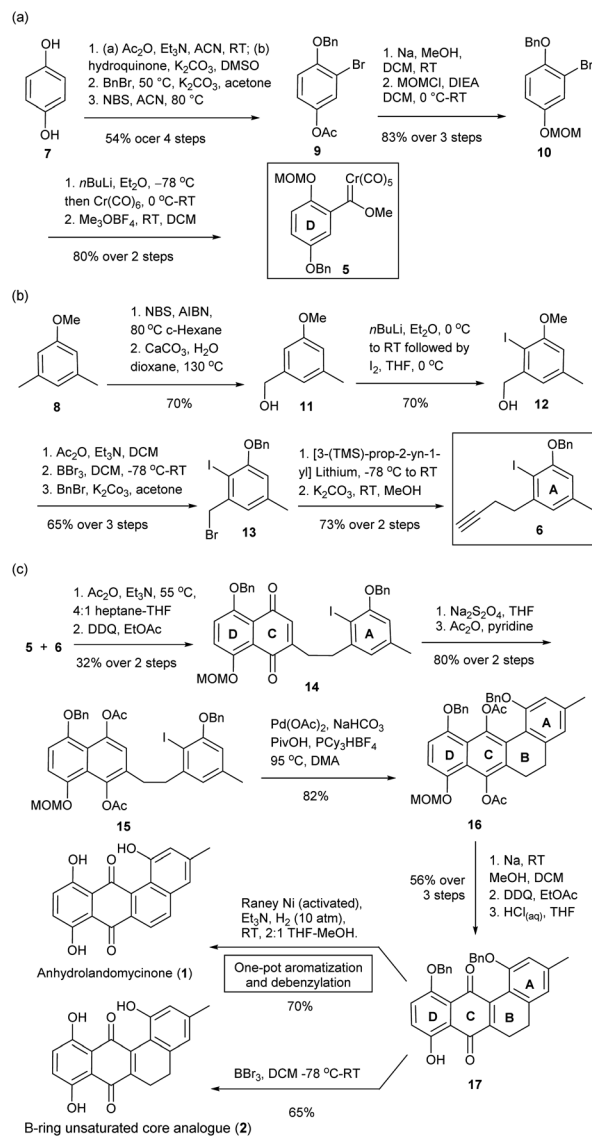


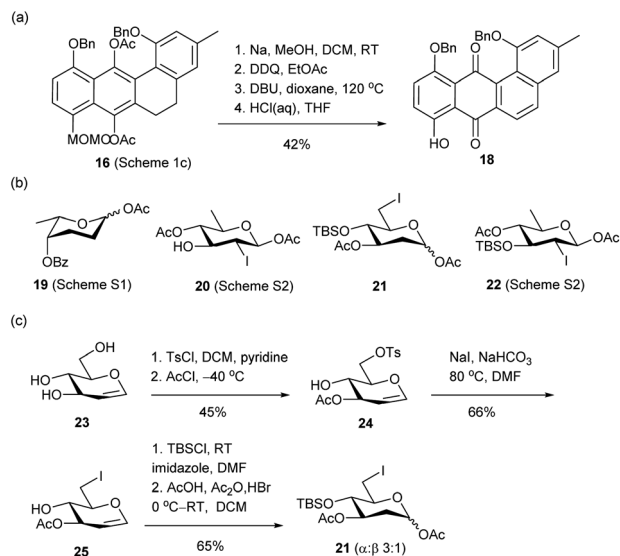
Fig. 1 (a) General Structure of LAs. (b) Structures of anhydrolandomycinone aglycone core (1), and B-ring unsaturated core analogue (2), LA R (3) and LA Q (4).

alone are inadequate for procurement of a complete LA scaffold.^{20,21} Some modifications are required for selective removal of protecting groups and subsequent coupling reactions; but such modifications may not be tolerable to the original protocols. Thereby, we envisaged some synthetic challenges to realize the structures of our targets. These include: (i) the construction of a highly hindered B-ring; (ii) stereocontrol of the formation of the 2-deoxy- β -glycosidic bonds; and (iii) the global deprotection in conditions tolerated by the rather fragile trideoxy- and dideoxy-glycosidic bonds (Fig. 1b).²²

To tackle these challenges, new synthetic routes were designed, which was based on the modifications of literature protocols.^{19–21} To access anhydrolandomycinone core (1) and its B-ring unsaturated analogue (2), the D-ring and A-ring building blocks 5 and 6 were prepared from available hydroquinone 7 and dimethylanisole 8, respectively. Based on the building blocks 5 and 6, the C and B-rings of the core scaffold were constructed.

In the preparation of the D-ring building block 5, regioselective acetylation of hydroquinone 7 with Zhang's procedure²³ followed by sequential benzylation and bromination gave the known 3-bromo-4-hydroxyphenyl acetate 9.²⁴ Then, 9 was converted to 1-(benzyloxy)-2-bromo-4-(methoxy-methoxy)-benzene 10 through deacetylation and methoxymethyl ether (MOM) protection (Scheme 1a). Subsequent lithiation of 10 in couple with Meerwein's salt methylation completed the preparation of building block 5 in 37% yield over seven steps.²⁵ Because of the poor stability of 5, it was freshly prepared prior to the Dötz benzannulation shown in Scheme 1c.



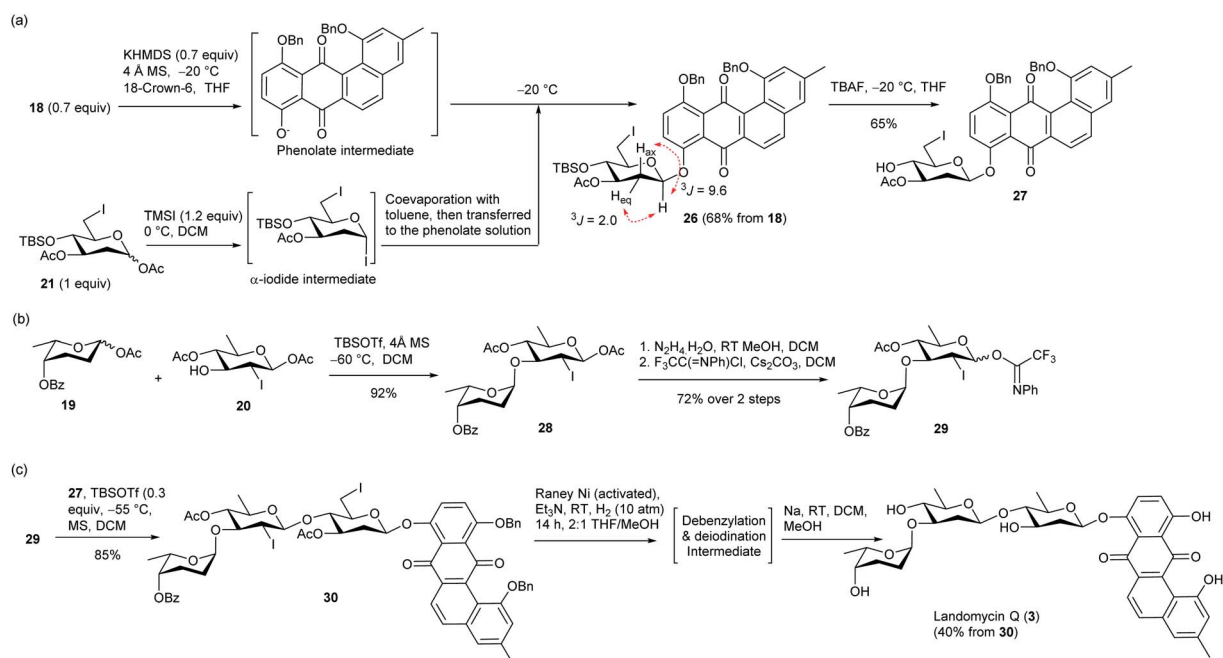


Scheme 2 (a) Preparation of anhydrolandomycinone acceptor **18**. (b) Structures of *L*-rhodosyl acetate **19** and olivosyl acetates **20–22**. (c) Preparation of *D*-olivosyl acetate **21**.

With the *D*- and *A*-ring building blocks **5** and **6** in hand, we embarked on the *C*-ring construction *via* Dötz benzannulation (Scheme 1c).²⁸ The annulation was followed by 2,3-dichloro-5,6-dicyano-benzoquinone (DDQ) oxidation to give the *A*-ring tethered naphthalene-1,4-dione **14**; which through $\text{Na}_2\text{S}_2\text{O}_4$ reduction and acetylation was converted to the *A*-ring tethered naphthalene-1,4-diacetate **15**. Subsequent acetate-assisted intramolecular directed arylation enabled construction of the *B*-ring in a highly congested environment.²⁹ The desired 5,6-

dihydrotetraphene-7,12-diyl-diacetate **16** was procured in a good 82% yield along with just a trace amount of the deiodination product of **15**. Removal of the acetyl groups of **16** followed by DDQ oxidation and acidic cleavage of the MOM protecting group gave 5,6-dihydrotetraphene-7,12-dione **17**. In subsequent hydrogenolytic debenzoylation, the modified RANEY®Ni catalyzed hydrogenation conditions were applied; whereas, the RANEY®Ni catalyst was freshly activated with $\text{NaOH}_{(\text{aq})}$ and triethylamine (Et_3N) was added as an acid scavenger.³⁰ Under the new conditions, no reduction of the carbonyl groups at *C*-ring occurred, while the *B*-ring was aromatized to give desired anhydrolandomycinone (**1**) in a one-pot fashion. In previous hydrogenation conditions, the carbonyl groups at the *C*-ring were also reduced, which had to be recovered by the DDQ oxidation.^{19,20d} In addition to the synthesis of (**1**), removal of the benzyl protecting groups of **17** with BBr_3 furnished the *B*-ring unsaturated core analogue (**2**).

For the synthesis of LA **Q** (**3**) and **R** (**4**), partially protected anhydrolandomycinone **18** was initially employed as the aglycone acceptor, which could be derived from advanced intermediate **16** *via* four functional- and protecting-group manipulation steps (Scheme 2a). The 2-deoxytrisaccharide component of LA **Q** (**3**) was assembled from *L*-rhodosyl acetates **19** (ref. 31) and, *D*-olivosyl acetates **20**,^{19a} and **21**, while the 2-deoxy-disaccharide component of LA **R** (**4**) was built from the *D*-olivosyl acetates **21** and **22** (ref. 19a) (Scheme 2b). *L*-Rhodosyl acetate **19** was derived from *L*-rhamnose (Scheme S1†), and *D*-olivosyl acetates **20** and **22** were derived from known glucal **23** according to literature procedures (Scheme S2†).³² 6-Iodo-olivosyl acetate **21** was also prepared from **23** *via* intermediates **24** and **25** in 20% overall yield (Scheme 2c).



Scheme 3 (a) Preparation of anhydrolandomycinonyl β -olivoside acceptor **27**. (b) Preparation of disaccharide imidate donor **29**. (c) Total synthesis of LA **Q** (**3**).

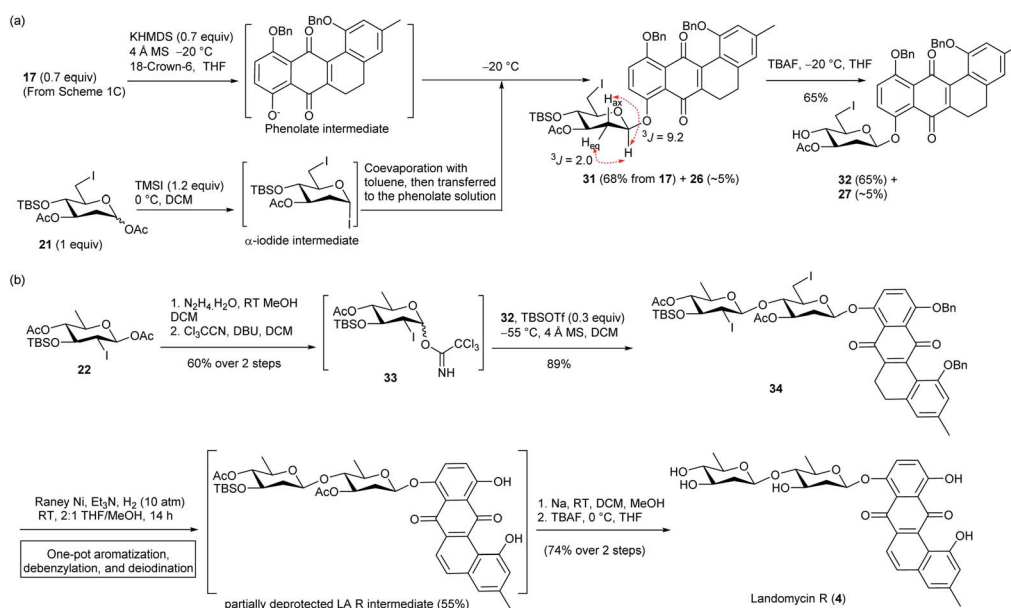


In the construction of LA Q (3), an anhydrolandomycinonyl β -olivoside acceptor was prepared, which was subsequently coupled with a non-reducing-end disaccharide donor. In the preparation of the β -olivoside acceptor, 6-iodo-olivosyl acetate **21** was converted to an α -olivosyl iodide by treatment with trimethylsilyl iodide (TMSI) (Scheme 3a).³³ After formation of the iodide intermediate, the DCM and residual TMSI were removed and the crude intermediate was reacted with the phenolate nucleophile, which was prepared *in situ* by deprotonation of **18** with potassium hexamethyldisilazane (KHMDs). Under the S_N2 glycosylation conditions, the desired anhydrolandomycinonyl β -olivoside **26** was obtained in a satisfactory 68% yield with perfect selectivity. Assignment of the configuration of the *O*-aryl 2-deoxy- β -glycosidic bond was supported by the $^3J_{H1-Hax}$ coupling constant of 9.6 Hz. Notably, the above substitution occurred exclusively at C1 position despite the presence of the iodo substituent at the C6 position. Subsequent removal of the silyl protecting group afforded the desired β -olivoside acceptor **27** in 65% yield.

For the preparation of the disaccharide donor, olivosyl acetate acceptor **20** was coupled with rhodinosyl acetate donor

19 to give α -rhodinosyl-(1,3)-olivosyl acetate **28** (Scheme 3b). In the glycosylation, a low -60°C temperature was applied to avoid the cleavage of the fragile 2,3,6-trideoxyglycosidic bond. Final replacement of the anomeric acetate group of **28** with *N*-phenyltrifluoroacetimidate completed the preparation of disaccharide imidate donor **29**.

The availability of donor **29** set the stage for the assembly of the target (Scheme 3c). Glycosylation of acceptor **27** with donor **29** gave protected LA Q **30** in a high 85% yield with excellent β -selectivity. The selectivity of the glycosylation is attributed to the participatory effect of the 2-iodo substituent group.³⁴ Subsequent one-pot deiodination and debenzoylation of **30** could be achieved under the same RANEY®Ni hydrogenation conditions as shown in Scheme 1c. Hydrolysis of the acyl protecting groups of the deiodinated and debenzoylated intermediate concluded the synthesis of LA Q (**3**) in 40% yield over two steps from **30**. The identity and structure of **3** were confirmed with HRMS and NMR spectroscopy. The proton chemical shifts of (**3**) were in full agreement with the data given from an authentic sample (Table S1†).^{16c}



Scheme 4 (a) Preparation of B-ring unsaturated anhydrolandomycinonyl core acceptor **32**. (b) Total synthesis of LA R (**4**).

Table 1 Inhibitory effects on normal/cancer cell proliferation by anhydrolandomycinone (**1**), B-ring unsaturated core analogue (**2**), LA Q (**3**), and LA R (**4**)^a

Entry	Compound	IC ₅₀ (μM)		
		NCI-H460	SF-268	Detroit 551
1	Core (1)	7.00 ± 0.70	8.57 ± 0.34	>10.0
2	B-ring unsaturated core (2)	>10.0	>10.0	9.58 ± 0.66
3	LA Q (3)	>10.0	>10.0	>10.0
4	LA R (4)	>10.0	9.82 ± 0.08	>10.0

^a Values represent the mean ± SD of three independent experiments.



Inspired by the one-pot debenzoylation and aromatization in Scheme 1c, and the one-pot debenzoylation and deiodination in Scheme 3c, we envisaged a simpler synthetic route for LA R (4), which would invoke a one-pot three-step, *i.e.* aromatization–debenzoylation–deiodination, procedure. With this aim, 5,6-dihydrotetraphene-7,12-dione **17** in Scheme 1c was coupled with 6-iodo oliviosyl acetate **21** according to the above S_N2 -glycosylation procedure to give 5,6-dihydrotetraphene-7,12-dione β -olivioside **31** in 68% yield along with *ca.* 5% of the inseparable aromatization product **26** (Scheme 4a). The crude β -olivioside **31** was subjected to desilylation to give the 5,6-dihydrotetraphene-7,12-dionyl β -olivioside acceptor **32** along with a small amount of olivioside **27**.

On the other hand, the non-reducing end oliviosyl acetate **22** was converted to the trichloroacetimidate donor **33** by standard procedures (Scheme 4b). Subsequent glycosylation of β -olivioside acceptor **32** with donor **33** furnished 8-disaccharidyl-5,6-dihydrotetraphene-7,12-dione **34** in 89% yield, though a tiny amount of glycosylation product arising from contaminant olivioside **27** was also produced. Without separation, the crude β -glycoside **34** was subjected to the one-pot aromatization–debenzoylation–deiodination procedure to afford a partially deprotected LA R intermediate in a satisfactory 55% yield. Final deprotection of the silyl and acetyl protecting groups concluded the total synthesis of LA R (4) in 74% yield over two steps.

Exploration of the cytotoxic and antibacterial properties

Having acquired anhydrolandomycinone (1), the B-ring unsaturated core analogue (2), LA Q (3) and LA R (4), we evaluated their cytotoxicity with the [3-(dimethylthiazol-2-yl)-5-(3-carboxymethoxyphenyl)-2-(4-sulfo-phenyl)-2H-tetrazolium] (MTS) assay (Table 1).^{35,36} Anhydrolandomycinone (1) was found to be cytotoxic against the cancer cell lines NCI-H460 and SF-268 with IC_{50} values at 7.00 ± 0.70 and 8.57 ± 0.34 μ M, respectively; but no appreciable cytotoxicity was noted for normal cell line Detroit 551 at 10 μ M (Entry 1). Intriguingly, such a toxicity pattern was reversed for the B-ring unsaturated core analogue (2) (Entry 2). At 10 μ M concentration of glycosylated LA Q (3) and R (4), no appreciable cytotoxic effect was detected for the normal cell line (Entries 3 and 4). Taken together, the unsaturated B ring and/or the 2-deoxysaccharide chain appear to play a role in the cytotoxicity of the LA compounds.

To elucidate the extent of the cytotoxicity, we evaluated the survival rates of Detroit 551 cell line at 10 μ M of the compounds, which were found to be $59.00\% \pm 3.26\%$ for the core structure (1), $104.2\% \pm 20.67\%$ for LA Q (3), and $93.56\% \pm 10.02\%$ for LA

R (4), implicating that the anhydrolandomycinone alone is more toxic than its glycosylated products.

After clarifying the cytotoxicity profiles of 1–4, we examined the antibacterial activity with MRSA 4N 216 and 7Y001 (Table 2).^{37,38} Among the compounds examined, only the disaccharide substituted LA R (4) exhibited inhibitory activity against MRSA 4N 216 and 7Y001 with the MIC values of 8 and 4 μ g mL^{−1}, respectively. Although, such a potency is inferior to that offered by vancomycin (VA), the non-toxic nature of LA R (4) renders it a potential candidate for further optimization.

Conclusion

In summary, a versatile strategy was established for the total synthesis of the anhydrolandomycinone core (1), the B-ring unsaturated anhydrolandomycinone analogue (2), landomycin Q (3), and landomycin R (4). The strategy employed the acetate-assisted intramolecular arylation for construction of the hindered B-ring of the anhydrolandomycin core and a one-pot aromatization, deiodination, and debenzoylation procedure was established to simplify the end-stage functional group manipulations. In subsequent MTS study, the structure of the B-ring and the degree of glycosylation of the landomycins appear to play a role in the cytotoxicity. Further exploration of the antibacterial properties revealed the potential of landomycin R (4) for inhibition of MRSA 4N 216 and 7Y001.

Conflicts of interest

There are no conflicts to declare.

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Table 2 *In vitro* inhibitory assay with anhydrolandomycinone (1), B-ring unsaturated core analogue (2), LA Q (3), and LA R (4)^a

MRSA strain	MIC (μ g mL ^{−1}) of compound				
	Core (1)	Analogue (2)	LA Q (3)	LA R (4)	VA
4N216	>8	>8	>8	8	1
7Y001	>8	>8	>8	4	1

^a Values are derived from two independent MIC experiments.



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