Abstract: Lemonomycin (I) was first isolated from the fermentation broth of Streptomyces candidus in 1964. The complete chemical structure was not elucidated until 2000 with extensive spectroscopic analysis. Lemonomycin is currently known as the only glycosylated tetrahydroisoquinoline antibiotic. Its potent antibacterial activity against Staphylococcus aureus and Bacillus subtilis and complex architecture make it an ideal target for total synthesis. In this short review, we summarize the research status of lemonomycin for biological activity, biosynthesis, and chemical synthesis. The unique deoxy aminosugar-lemonose was proposed to play a crucial role in biological activity, as shown in other antibiotics, such as arimetamycin A, nocathiacin I, glycothiohexide α, and thiazamycins. Given the self-resistance of the original bacterial host, the integration of biosynthesis and chemical synthesis to pursue efficient synthesis and further derivatization is in high demand for the development of novel antibiotics to combat antibiotic-resistant infections.

Keywords: aminosugar; antibiotic; biosynthesis; glycosylation; lemonomycin; total synthesis

1. Isolation and Structural Elucidation

The golden era of antibiotics propelled by the discovery of penicillin has been challenged by a growing crisis in antibiotic resistance worldwide and a decline in investments in the development of novel antibiotics [1]. Natural products are the mainstays of modern antibiotic drugs. Although, only approximately two hundred of 28,000 have been used clinically [2]. The chemical diversity of natural products, including those structures known for decades on the shelf, remains a huge space that has the potential to be a privileged starting point to explore. Lemonomycin (I) was first isolated from a fermentation broth of Streptomyces candidus (LL-AP191) in 1964 by Whaley and coworkers at the Lederle Laboratory of American Cyanamid Company (Figure 1) [3]. A few grams of lemonomycin hydrochloride were obtained after separation and purification. Its name was based on the color of lemon yellow about the corresponding globular substance. According to the primary 1H nuclear magnetic resonance (NMR) and infrared (IR) spectroscopy, the researchers concluded that the structure of lemonomycin should contain hydroquinone, as well as N-methyl and O-methyl functional groups. At the same time, lemonomycin released dimethylamine after the degradation experiment. The biological evaluation of lemonomycin revealed its broad antimicrobial activity against various microbial pathogens [3,4]. Unfortunately, lemonomycin was found to be fatal at a dose higher than the therapeutic dose, indicating why this potent antibiotic was shelved for decades without further characterization and derivatization.
The antitumor activity arises from the electrophilic iminium ion via a carbinolamine moiety of position of C18 through glycosylation [6,7]. Another feature is the existence of a hydrate feature of lemonomycin is that 2,6-dideoxy-4-amino sugar (lemonose) is connected at the position of C18 through glycosylation [6,7]. Another feature is the existence of a hydrate form of the aldehyde at the position of C16. One of the OH groups of the hydrate can form an intramolecular hydrogen bond, which stabilizes the acetal. The acetal was believed to be the warhead that participates in binding DNA in vivo, as shown in the mode of action of other tetrahydroisoquinolines (THIQs) [6,7]. Lemonomycin and its cyano derivative exhibited strong cytotoxic effects on a human colon cancer cell line (HTC116) in vitro, and their IC50 values were 0.36 μg/mL and 0.26 μg/mL, respectively. The minimal inhibitory concentration (MIC) was re-measured to exert 0.4 and 0.2 μg/mL for methicillin-resistant Staphylococcus aureus and vancomycin-resistant Enterococcus faecium. The absolute stereochemistry of lemonomycin remained elusive at this stage, although the configuration of the aglycon was tentatively assigned according to the absolute stereochemistry of the isoquinoline antibiotics [6,7].

Lemonomycin belongs to the family of tetrahydroisoquinoline (THIQ) antibiotics, a family of more than 60 congeners known to have various potent biological activities [6,7]. The antitumor activity arises from the electrophilic iminium ion via a carbinolamine moiety and its equivalents, which directly alkylate the N-2 residue of guanine in the GC-rich region of DNA [6–8] (Figure 2a, quinocarcin as an example). THIQ antitumor antibiotics are characterized by the diazabicyclic structure bridged in the tetrahydroisoquinoline ring system. According to the structural features of the bridged diazabicyle, these compounds can be divided into two categories, including two-carbon bridges (e.g., quinocarcin, cyanocycline A, tetrazomine, and lemonomycin) containing a 3,8-diazabicyclo[3.2.1]octane core (Figure 2b) and three-carbon bridges (e.g., saframycin A, renieramycin C, and ecterinasidin 743) containing a diazabicyle [3.3.1]nonane core (Figure 2c). Given their potent antitumor activity and great potential in the development of antineoplastic drugs, THIQ antibiotics have aroused the enormous interest of synthetic chemists [6,7]. At present, several research groups have completed the synthesis of the THIQ core of lemonomycin, named lemonomycinone amide (2, lemonomycin aglycon).

Figure 2. Representative THIQ antitumor antibiotics and mode of action (MoA). In the Cya17/19 complex, adenylated 5 with Cya19 was transferred to Cya17-PCP, followed by a Cya17-catalyzed Pictet–Spamann cyclization. (a) Chemical structure of lemonomycin (1), isolated sample, and lemon.
The biosynthesis of lemonomycin has not been fully illustrated and literally constructed through the glycosylation of lemonomycinone amide (2, aglycon) with lemonose (3) (Scheme 1a). Lemonomycin shares a common skeleton with other quinocarcin alkaloids despite the appended deoxy amino acid. Therefore, the biosynthetic pathway can be largely similar to the biosynthetic pathway dedicated to quinocarcins. As the first reported antibiotic in this category, naphthyridinomycin [9,10] (NDM, 4) has a tetracyclic ring in a nearly identical substitute pattern, except for the C4-amino group and the inverse configuration of the acetal at C15, which constitutes two additional heterocyclic rings in NDM (Scheme 1b). The labeled precursor feeding experiments have established the origin, and the molecular basis from L-tyrosine, L-methionine, serine (or glycine), D,L-ornithine, and ketose phosphates could be incorporated into NDM [11–16]. Despite extensive biosynthetic studies of the THIQ core as well as other rare deoxysugars, the biosynthesis of lemonose in vivo and the detailed enzymatic glycosylation remain unclear [17,18].

In 2013, Oikawa and coworkers performed gene cluster identification and biochemical analysis of substances in the tetrahydroisoquinoline family, from which the biosynthetic pathways quinocarcin and SF-1739 were proposed (Scheme 2) [19]. On the starting template of Cya13/12, fatty acid and L-Ala units were loaded, while the template Cya15 was likely responsible for carrying the glycol unit [14] from the ACP of D-xylulose-5-phosphate (10) to make its A domain active. Condensation with L-aminolevulinc acid and the subsequent reduction delivered aldehyde 11 by Cya17. In the separated pathway, L-tyrosine was converted to 5 through a series of C-methylation, hydroxylation, and O-methylation. By conducting in vitro assays, Cya18 was confirmed to be α-ketoglutarate-dependent dioxygenase catalyzing the dehydrogenation of L-Arg to form L-(E)-4,5-dehydroarginine (12). In the Cya17/19 complex, adenylated 5 with Cya19 was transferred to Cya17-PCP, followed by a Cya17-catalyzed Pictet–Spengler reaction with the leader peptide-tethered aldehyde 11 to generate aldehyde 13, with the assistance of reduction by the Cya17/19 complex. With the activation of 12 under the condition of adenylation provided by Cya16/17, the Cya17-PS domain was used to selectively catalyze the coupling of 12 and 13 via an intramolecular Mannich reaction, reductive cleavage, and spontaneous cyclization to release hemiaminal 14, bearing an endo-configuration at C15, providing an essential carbon framework of SF-1739 and congeners, including aclidinomycins and lemonomycin.
Ju and coworkers recently isolated eleven THIQ-like aclidinomycins from deep-sea-derived *Streptomyces niveus* SCSIO 3406 by genome orientation and the one-strain-many-compounds (OSMAC) strategy [20]. Aclidinomycin J and aclidinomycin K were isolated as amorphous yellow powders and found to have the characteristic core structure of lemonomycin, except for the inverse stereochemistry at C15 and saturation at C17. As indicated in Oikawa’s investigation [19], the specific route is possibly initiated from xylulose-5-phosphate (Xyl-5-P), aminolevulinic acid (Ala), myristoyl acid, a L-tyrosine derivative (5), and L-(E)-4,5-dehydroarginine (12) as the starting materials (Scheme 3). The discovery of aclidinomycins provides new insights into the molecular logic of the complex polycyclic architecture.

Very recently, Tang and coworkers reported an interesting mechanism of self-resistance when the bacterial host produced lemonomycin and naphthyridinomycin [21]. In the fermentation of *S. candidus* LL-AP191, 16-dehydroxy-LMM (1a), was identified instead of LMM (Scheme 4). Large-scale enzymatic reactions were performed to deliver the C17-reduced form 1b, indicating additional reduction and oxidation steps occurring at different times and locations. As a self-protection strategy, the pharmacophore of the
iminium ion at C17 was largely reduced in the cytoplasm to maintain a low concentration for the host. The additional oxidase NapU (flavoprotein) realized the extracellular oxidation step at C17 and C16 to generate the highly potent antibiotic LMM (1). The self-resistance strategy poses a challenge to develop a sophisticated biosynthetic pathway to reharvest such potent antibiotics in engineered biosynthetic machinery [22].

Scheme 3. Proposed biosynthetic pathway toward aclidinomycins J and K.

Scheme 4. SDR-catalyzed reduction of 16-dehydroxy-LMM (1a). (SDR: short-chain dehydrogenase/reductase).

3. Total Synthesis of Lemonomycin

The unique structure and potent antimicrobial activity have attracted considerable interest in several laboratories. Two groups completed the total synthesis, and five groups accessed the aglycon or the key intermediates to constitute the virtual synthesis. The Stoltz group was the first team to disclose the total synthesis of lemonomycin. As shown in Scheme 5 [4,23], the retrosynthetic plan features the Pictet–Spengler reaction of amide 18 with glycosylated aldehyde 19. Fragment 18 could be readily constructed by the aryl boronic ester 20 and vinyl iodide 21 via Suzuki cross-coupling. The characteristic diazo-bicyclic intermediate 21 was successfully established between dipolar precursor 23 and Oppolzer sultam derivative 24 by 1,3-dipolar cycloaddition, a novel tactic initially developed by Joule and coworkers [24–26] and widely adopted in the syntheses of other THIQ alkaloids [6,7]. Glycosylated aldehyde 19 bearing an amino sugar could be prepared from D-threonine in multiple steps.
Following the literature-known protocols, methyl ester 25 was prepared from D-threonine in three steps by the benzenesulfonyl group replacing the original Cbz protection. After the aminolysis of 25 with N,O-dimethylhydroxylamine, the resulting Weinreb amide was converted into methyl ketone 26 in 75% yield after two steps (Scheme 6). Aldol addition with ethyl acetate produced a virtually single diastereomer 27 via the Felkin–Ahn trajectory [27,28]. After treatment of β-hydroxy ester 27 with hydrochloric acid, the corresponding lactone 28 reacted with dimethoxymethane to form oxazolone 29 by using TMSOTf as a catalyst. The subsequent reduction by DIBAL-H and dehydration with allyl alcohol delivered α-allyl ether 30. The reductive cleavage of the benzenesulfonyl group by Red-Al and the subsequent reductive methylation formed allyl glycoside 32 in excellent yield. The oxidative cleavage of the alkene by Lemieux–Johnson reagent [29] proceeded smoothly to yield aldehyde 19 as a trifluoroacetate salt (55%) and in an acid-free form (30%).

Scheme 5. Stoltz’s retrosynthetic analysis of lemonomycin (1).

Scheme 6. Preparation of aldehyde 19 (the glycoside fragment).

Bromide salt 23 as a precursor for the requisite 1,3-dipole cycloaddition was readily prepared using benzyl bromide 34 and 6-methyl-2(1H)-pyrazinone in anhydrous ethanol and acetonitrile at the reflux in a 96% yield (Scheme 7). However, dipolarophile 24 was obtained by Kocierński’s protocol [30] with silylative activation and the subsequent acylation by acryloyl chloride in the presence of cupric chloride. Accordingly, N-Me-morpholine was used for the deprotonation of salt 23 to form an oxidopyrazinium salt for the 1,3-dipolar
cycloaddition with 24. Immediate treatment by NaBH$_4$ carried out the reductive removal of the sultam auxiliary to deliver enamide 36 in good yield and a high stereoselectivity (94% ee), indicating an excellent diastereoselectivity achieved in the cycloaddition event. The protection of the primary alcohol with a bulky silyl group and the next iodination produced vinyl iodide 21 as a single isomer in good yield.

![Scheme 7](image)

**Scheme 7.** Preparations of diazobicyclic amide 21 and arylboronate 20.

Arylboronate 20 was effectively synthesized from 2,6-dimethoxytoluene 22 in five straightforward steps, including Friedel–Crafts acylation, Baeyer–Villiger oxidation, tosylation, bromination, halogen exchange, and subsequent trapping by the boron ester (Scheme 6). With two fragments 20 and 21 in hand, Suzuki cross-coupling catalyzed by Pd(PPh$_3$)$_4$ proceeded to deliver the requisite aryl enamide 40 on the gram scale with a good isolated yield (69%) (Scheme 8). The hindered trisubstituted enamine was subjected to heterogeneous hydrogenation at 1000 psi to generate 41 with excellent diastereoselectivity. Reprotection of the amine group by the Cbz group and the subsequent removal of the $p$-toluenesulfonyl group by potassium trimethylsilolate (KOTMS)-promoted hydrolysis released the cyclic amide in 42. To release amine for the programmed Pictet–Spengler reaction, treatment with (Boc)$_2$O enabled the complete reduction of the amide group. The subsequent acid workup produced amino triol 43 as a form of trifluoroacetic acid (TFA) salt in 81% yield after three steps. The key Pictet–Spengler cyclization of 43 with lemonose-derived aldehyde 19 (TFA salt) was performed in a foil-wrapped vial for 63 h to give bis-trifluoronate 44 in a 95% yield. Again, the PS-cyclized product was isolated in the form of a single diastereomer at C1. Hydrogenolysis and Swern oxidation resulted in hemiaminal bis-trifluoronate 46 bearing a hydrate form of aldehyde. Finally, oxidation by cerium ammonium nitrate (CAN) to convert phenol into benzoquinone furnished bis-trifluoronate (−)-lemonomycin (1), which was completely identical to the reported data [5], and thus fully validated the stereochemistry of lemonomycin.

Given novel aminosugars in various potent antibiotics, the Fukuyama group proposed a late stage of glycosylation in the synthetic design to be versatile for future derivatization. In 2012, the total synthesis of lemonomycin was reported by this group [31]. As shown in Scheme 9, aglycon 48 was anticipated by reducing the enamine in 49 and its subsequent cyclization via the Pictet–Spengler reaction. The 3,8-diazabicyclo[3.2.1]octane skeleton was established by the Hosomi–Sakurai reaction, a strategy being executed in their previous syntheses toward quinocarcin and other alkaloids [32–34]. Critical intermediate 50 could be generated by a Perkin-type reaction of diketopiperazine 51 and aldehyde 52. The
latter aldehyde was derived from 2,6-dimethoxybenzene in five steps by a modification of Stoltz’s route [22].

Scheme 8. Complete synthesis of (−)-lemonomycin (1).

The synthesis commenced with the application of glycine-derived Oppolzer sultam 53 for asymmetric alkylation with propargylic iodide 54 [35] to deliver a single diastereomer of 55 after hydrolysis in an ether solution of HCl (Scheme 10). After condensation with N-tert-butyloxy carbonyl (Boc) glycine, the intermediate derived from the acidic removal of the Boc group was then cyclized to release diketopiperazine 56 in a combined 84% yield after three steps. Two secondary amides were masked by the Boc protection, and the resulting cyclic amide underwent Perkin-type condensation [36], with aryl aldehyde 52 to favor (Z)-enamine 57 (Z/E = 10:1) in an excellent yield. The triple bond in 57 was partially reduced by the Lindlar catalyst, while the carbonyl group at C11 was reduced to hemiami-
nal with NaBH₄ in an aqueous methanol solution. The acid-promoted Hosomi–Sakurai reaction successfully constructed the bridged ring of 59 with almost perfect stereoselectivity. Reprotection of the amine with the Cbz group gave lactam 59 in a 51% overall yield in three steps. In contrast to Stoltz’s synthesis, trisubstituted enamine was smoothly reduced by NaBH₃CN under acidic conditions, and the stereochemistry at C1 was satisfied. Ozonolysis and immediate reduction furnished a hydroxymethyl group at C15 in 60. Functional group manipulations, including protection by the tert-butyldiphenylsilyl (TBDPS) group and deprotection of the mesyl group by lithium bis(trimethylsilyl)amide (LiHMDS) [37], generated the phenol derivative 61. To improve the reactivity of the amine, the lactam group was first reduced by diisobutylaluminum hydride (DIBAL-H), and the resulting hemiaminal was converted to amino nitrile 62. To complete the synthesis of the aglycon, cinnamaldehyde was merged into the desired Pictet–Spengler cyclization in the presence of CSA and TMSCN, and then the tetracyclic skeleton 63 was formed at 92%. The high stereoselectivity achieved was found to be crucial under thermodynamic control. After acetylation, the requisite aglycon 65 was completed in an 84% yield after ozonolysis and subsequent reduction.

Scheme 10. Fukuyama’s synthesis of lemonomycin—preparation of aglycon 65.

The synthetic route of lemonose 71 was slightly modified from the previous procedure [22]. The initial amine protection group was chosen as the Cbz group instead of the benzenesulfonyl group in a previous synthesis (Scheme 11). After the formation of methyl ketone 67 through Weinreb amide, the aldol reaction and deprotection afforded lactone 68. The tertiary alcohol was protected by the tert-butyl(dimethylsilyl) (TBS) group, followed by the removal of the Cbz group and reductive amination in a one-pot method. The corresponding amine 69 was subjected to partial reduction by DIBAL-H to release acetol 70. Finally, fluorination by treatment with dimethylaminosulfur trifluoride (DAST) [38] resulted in glycosyl fluoride 71 in 84% yield.
With aglycon 65 and glycosyl fluoride 71 in hand, glycosylation was promoted by trimethylsilyl trifluoromethylsulfonate (TMSOTf) to give α-anomer 72 in 86% yield (Scheme 12). After the protection exchange, the fully protected compound 73 was subjected to debenzylation by hydrogenolysis and the subsequent removal of two silyl groups by tetra-n-butylammonium fluoride (TBAF) to generate amino alcohol 74 as a bistrifluoromethylsulfonate salt. Subsequent Swern oxidation smoothly converted the primary alcohol into hydronated aldehyde 75 after acidic workup. Silver nitrate was then adapted to remove the CN group to rebound the hemiaminal functionality, which was also tolerant to the final oxidation by CAN to complete the total synthesis of lemonomycin (1). This strategy diminished the separation steps of polar intermediates, which are clearly tedious and troublesome to handle because of the given salt forms in the late stage of total synthesis.

Scheme 12. Fukuyama’s synthesis of (−)-lemonomycin—preparation of glycosyl 71.

4. Other Studies toward Lemonomycin

The synthetic works contributed by the Magnus and Zhu groups were essentially inspired by the late stage of the biogenesis of quinocarcin alkaloids (Scheme 2). They designed the pathway, including the construction of lemonomycinone amide (2, aglycon) via a Mannich-type cyclization, to furnish the bridged ring system and final glycosylation with lemonose (Scheme 13).
was subjected to reduction \[50\] at the distal ester. The subsequent additional three steps of TBAF for desilylation to release the phenol group enabled the Pictet–Spengler reaction of 101 phase transfer catalyst potassium bis(trimethylsilyl)amide (KHMDS) as a base. To invert the stereochemistry of lemonomycinone amide (at C15 under acidic conditions. The final oxidation by CAN completed the synthesis of gies, they first established the C13 stereochemistry by the asymmetric alkylation of glycine-derived imine and silylated propargyl alcohol 77 under Castro conditions [42] and subsequent aromatization resulted in isoquinoline 78 in a 76% overall yield (Scheme 14). The subsequent addition of benzyloxymethyl lithium [43] and methyl chloroformate furnished the substitution on the B-ring. The removal of the trisopropylsilane (TIPS) group by TBAF also promoted cyclization to generate oxazolidinone, which was suitable for ionic hydrogenation by TFA-Et₃SiH and immediate hydrolysis to afford cis-amino alcohol 81 in a high yield. After structural confirmation by X-ray analysis, compound 82 was introduced into a glycine motif via an activated reagent, \(n\)-tert-butoxycarbonyl glycine. Swern oxidation initiated the formation of hemiaminal in a mixture of diastereomers (\(dr 3/2\)) [44], which was further converted into the thermodynamically stable thioaminal 84 (X-ray) in excellent yield. Single diastereomeric 86 was obtained from the alkylation of 84 and iodide 85 by potassium bis(trimethylsilyl)amide (KHMD) as a base. To invert the stereochemistry at C13, the treatment of 86 by \(^{1}\)BuLi and trapping by a hindered proton source, butylated hydroxytoluene (BHT), gave the \(\alpha\)-configured isomer which was desilylated to produce alcohol 87. After Swern oxidation, the resulting aldehyde was converted to silyl enolate 88 in 88% yield. The key Mannich-type cyclization was initiated by thioephile AgBF₄, and a single stereoisomer at C15 of the aldehyde was obtained. The high stereoselectivity is largely attributed to the steric bulk of the C1 substituent (BnOCH₂) in the axial orientation. After hydrogenation of two benzyloxyl groups by Pearlman’s catalyst Pd(OH)₂ [45], alcohol 89 was subjected to the removal of the Boc group and hydration of the aldehyde group at C15 under acidic conditions. The final oxidation by CAN completed the synthesis of lemonomycinone amide (2) in 35% yield as a racemic form.

In 2008, the Zhu group devised a similar Mannich-type strategy to synthesize lemonomycin amide (2) in an optical form [46]. In contrast to the abovementioned strategies, they first established the C13 stereochemistry by the asymmetric alkylation of glycine-derived imine 91 with functionalized benzyl bromide 90 in the presence of the Corey–Lygo phase transfer catalyst 93, [47–49] O-(9)-allyl-N-(9′-anthracenyl)methyl) cinchonidium bromide, producing amino ester 92 in an 85% yield (Scheme 15). The subsequent addition of TBAF for desilylation to release the phenol group enabled the Pictet–Spengler reaction of benzoyl acetaldehyde 94 to deliver a single cyclized product 95. After protection and ester exchange, compound 97 was obtained in excellent yield. To circumvent the linear manipulation in the Magnus synthesis [40], the Zhu group adopted L-glutamic acid (98) to introduce the four-carbon unit. However, the separate synthesis of 101 required seven steps to install the proper functional groups. After three steps of protection, compound 99 was subjected to reduction [50] at the distal ester. The subsequent additional three steps of protection and deprotection gave amino acid 101.

Scheme 13. A general bioinspired approach of Magnus and Zhu’s syntheses.
Scheme 14. Magnus’ synthesis of lemonomycinone amide (2).

Scheme 15. Zhu’s synthesis of lemonomycinone amide—preparations of 97 and 101.

The condensation of isoquinoline 97 and amino acid 101 was carried out to produce amide 102 without detectable epimerization in the presence of hexafluorophosphate az-
A particularly interesting phenomenon was discovered during the hydrogenolysis of 107a by the Perlman catalyst (Pd(OH)$_2$) [45] to remove two benzyl groups at room temperature (12 h at r.t. versus 6 h at 0 °C). In contrast to Magnus’s observation [39,40], the endo-isomer 108a (15-epi-108) was isolated in an 85% yield under identical conditions (Scheme 17). Although the endo-isomer is thermodynamically less favorable for steric reasons, the proximity of the C1-hydroxymethyl group may drive the equilibrium toward the final formation of hemiacetal 108b, providing mechanistic insights into the stereochemical outcome of the kinetically favored endo-adducts in the proposed biosynthesis [19].

Three years later, Zhu and coworkers continued to finish the asymmetric synthesis of lemonose (Scheme 18). [52] Threonine was again selected as the chiral pool, which was first converted to Weinreb amide 109 after three steps. Subsequently, two additions by different Grignard reactions readily furnished 111 with all requisite stereogenetic centers. The terminal double bond was then subjected to ozonolysis and a subsequent reduction to deliver alcohol 112 in an 81% yield for two steps. After global deprotection, double reductive amination with HCHO in the presence of NaBH$_3$CN furnished 114. To attenuate the impact of tertiary amines, the pretreatment of trifluoroacetic acid successfully chemoselectively converted the primary alcohol to aldehyde by O-iodoxybenzoic acid (IBX) [53] and then cyclized to form lactol 3 (lemonose) in a total of 10 steps and an 18% overall yield.
However, the complete assembly of lemonmycin (1) was not disclosed despite that both the given glycosylation of aglycon 2 with protection-free lemonose (3) were obtained. 

Scheme 17. A facile epimerization at C15 for biosynthetic implication.

Scheme 18. Zhu’s synthesis of lemonose (3).

Early in 2004, the Fukuyama group developed an intriguing Ugi-4CR to assemble four readily prepared fragments for the rapid construction of the polycyclic core of lemonomycin in a highly convergent manner, although no subsequent synthesis was reported [54]. Amino alcohol 115, amino acid 116, glyoxyaldehyde acetal 117, and isocyanide 118 were subjected to the Ugi reaction in CF₃CH2OH to generate dipeptide 119, followed by treatment with acid and cyclization under basic conditions to give N-acyloxazolidinone derivatives 120 in a good yield (Scheme 19). After the reduction and protection exchange, diacetate 121 was subjected to cross metathesis to introduce allyltrimethylsilane, which was elaborated to an intramolecular Hosomi–Sakurai reaction catalyzed by BF₃•Et₂O in an excellent yield, and complete stereochemical control at C15. To smoothly transform 122 into the desired alcohol 123, they had to operate a four-step sequence of exchanging two protecting groups. The oxidation of enamide 123 by dimethyl dioxirane (DMDO) resulted in 124, which upon treatment with TFA was reduced by NaBH₄ to furnish the stereogenic center at C3 from a less hindered exo-face. Enhancement of the nucleophilicity of the aromatic ring by KOSiMe₃ the promoted hydrolysis of the Ms group and benzylation in one pot. Alcohol 125 was oxidized by the Dess–Martin periodinane (DMP), and subsequent Friedel–Crafts cyclization yielded 126 with the requisite tetracyclic core of lemonomycin. Although they did not elaborate the current strategy to the final target, the key Hosomi–Sakurai cyclization was indeed successfully implemented into the later second-generation synthesis of lemonomycin (1) [31].
In 2008, Mulzer and coworkers completed a similar advanced intermediate by applying the intramolecular Hosomi–Sakurai reaction at different stages [55]. After a halogen exchange, the lithiation of arylbromide 127 reacted with aldehyde 128 to give a mixture of diastereomers of 129 (Scheme 20). An oxidation-reduction protocol secured a single diastereomer, which was then subjected to a protection–deprotection process to result in amino alcohol 130 in a good yield. After the removal of the benzyl group, the Pictet–Spengler reaction with aldehyde 131 proceeded under mild conditions to form the corresponding tetrahydroisoquinoline 132 in a highly selective manner. A Strecker-type condensation with cyanohydrin 133 was carried out to deliver hemiaminal 134 after oxidation of the primary alcohol at C16. Upon treatment with TFA, the reactive N-acyliminium ion-initiated Sakurai cyclization produced tetracyclic skeleton 135. The epimerization of the cyano group at C17 was largely attributed to the less demanding equatorial position via an iminium ion intermediate. A phenomenon was commonly encountered in the syntheses of THIQ alkaloids [6,7].

The latest effort devoted to the synthesis of LMM was reported in 2013 by Williams and coworkers [56]. They devised a 1,3-dipolar cycloaddition of functionalized oxidopyrazinium at a late stage of synthesis (Scheme 21). The synthesis was commenced from tetrahydroisoquinoline 137, which was derived from L-tyrosine methyl ester in 14 steps with a known protocol [57]. After acetylation and debromination, condensation with 139 and the subsequent transformation of the alcohol into aldehyde 140 resulted in a good overall yield. Treatment with TFA in CHCl₃ generated an iminium salt intermediate that was readily oxidized in the presence of (2,2,6,6-tetramethylpiperidin-1-yl) oxycarbonyl (TEMPO) and oxygen, followed by the addition of triethylamine to result in oxidopyrazinium 143. An immediate trap by tert-butyl acrylate smoothly afforded the requisite tetracyclic skeleton 144 with moderate facial selectivity (dr 2.4/1). The mixture was subjected to the following four steps: epimerization, partial reduction, benzylation, and isomerization at C1, and then the desired β-isomer at C1 was dominant. The reduction of aldehyde by NaBH₄, debenzylation by Pd/C, and hydrogenation from the bottom face by Raney nickel under a high pressure of hydrogen eventually delivered the advanced intermediate 146. The overall efficiency

**Scheme 19.** Fukuyama’s U-4CR approach toward the core of lemonomycin.
was largely detracted by the lengthy synthesis of aldehyde 140 and low diastereoselectivity of the intermolecular 1,3-dipolar cycloaddition. Therefore, a further transformation to aglycon 2 might not be facile due to the difficult accumulation of the key intermediate.

Scheme 20. Mulzer’s cyclization approach toward the core of lemonomycin.

Scheme 21. Williams’ intermolecular 1,3-dipolar cycloaddition approach [52].

5. Conclusions

The carbohydrate motifs in various natural products play a pivotal role for the development of bioactive glycoconjugates and carrier prodrugs by attenuating toxicity and improving bioavailability [58]. The rare amino sugar lemonose was essentially implemented in complex metabolites, such as nocathiacin I [59], thiazamycin [60], glycothiohexide α [61], arimetamycin A [62], daunorubicin and doxorubicin hybrids [63], and various oligosaccharides in antitumor saccharocarcins A–F [64] (Figure 3). These antibiotics have been found to have potent antimicrobial activity against Gram-positive and Gram-negative pathogens, implicating the crucial impact of deoxyaminosugars.
hexide α [61], arimetamycin A [62], daunorubicin and doxorubicin hybrids [63], and various oligosaccharides in antitumor saccharocarcins A–F [64] (Figure 3). These antibiotics have been found to have potent antimicrobial activity against Gram-positive and Gram-negative pathogens, implicating the crucial impact of deoxyaminosugars.

Figure 3. Antibiotics bearing 2,6-dideoxy-4-aminoglycoside (blue).

Although various synthetic methods have been developed for the construction of deoxy aminosugars [65–68], the syntheses of lemonose and its derivatives were based mainly on D-threonine as the starting material, and the lengthy synthetic routes may render further modification [69]. Furthermore, since the latest synthetic efforts devoted to the synthesis of the LMM core in 2013 by the Williams group, there have been no reported efforts in the past decade. Efficient synthesis remains in high demand. Given the notorious difficulties in the chemical synthesis of complex structures in this specific family, enzymes are, therefore, an appealing alternative with excellent chemoselectivity, regioselectivity, and stereoselectivity [70]. With the development of gene recombination technology in recent years, metabolic engineering has emerged as a powerful tool for the accumulation of bulky materials and even complex targets [17,18,71]. The integration of biological and chemical methods encourages access to lemonomycin and its derivatives by gaining insights into the critical role of aminosugars and defining potent analogs for future antibiotic development to combat antibiotic-resistant infections.

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