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New Broad-Spectrum Antibacterial Amphiphilic Aminoglycosides Active against Resistant Bacteria: From Neamine Derivatives to Smaller Neosamine Analogues

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Supporting Information

ABSTRACT: Aminoglycosides (AGs) constitute a major family of potent and broad-spectrum antibiotics disturbing protein synthesis through binding to the A site of 16S rRNA. Decades of widespread clinical use of AGs strongly reduced their clinical efficacy through the selection of resistant bacteria. Recently, conjugation of lipophilic groups to AGs generated a novel class of potent antibacterial amphiphilic aminoglycosides (AAGs) with significant improved activities against various sensitive and resistant bacterial strains. We have identified amphiphilic 3',6-dialkyl derivatives of the small aminoglycoside



neamine as broad spectrum antibacterial agents targeting bacterial membranes. Here, we report on the synthesis and the activity against sensitive and resistant Gram-negative and/or Gram-positive bacteria of new amphiphilic 3',4'-dialkyl neamine derivatives and of their smaller analogues in the 6-aminoglucosamine (neosamine) series prepared from N-acetylglucosamine.

INTRODUCTION

Neomycin B 1 (Figure 1) and its fragment neamine 2 are members of the major family of potent and broad-spectrum antibiotic drugs aminoglycosides (AGs) that are polycationic at physiological pH and act on bacteria through binding to the A site of 16S rRNA causing *in fine* protein synthesis alteration.^{1–10} Decades of widespread clinical use of AGs strongly reduced their clinical efficacy through the selection of resistant bacteria.^{8–19}

In the search for new antibiotics less toxic and susceptible to resistance than conventional AGs, Hanessian, Westhof, and coworkers reported the first example of ether-modified aminoglycoside paromomycin with in vivo antibacterial activities.^{20,21} Structural data with bacterial RNA have provided evidence of a novel binding mode, rendering these ether-substituted analogues less susceptible to inactivating enzymes.^{22,23} Subsequent studies from our laboratories^{24–29} and elsewhere^{11,30–45} have extended these results to other AGs named antibacterial amphiphilic AGs (antibacterial AAGs) resulting from the introduction of one to four lipophilic groups on AG cores. Such an increase in the AG lipophilicity results in a bacterial target shifting from rRNA to membranes and significantly improves activity against bacterial strains resistant to the parent AG drugs and to other classes of antibiotic drugs. Antifungal derivatives were also obtained in such an approach.^{44–46}

Targeting bacterial membrane functions remains an underexploited mechanism of action in the fight against resistant bacteria.⁴⁷ Persistent infections involving slow- and nongrowing bacteria are difficult to treat with antibiotics that target biosynthetic processes. In the fight against resistant bacteria, there is currently a growing interest for antibacterial cationic amphiphilic drugs acting on bacterial membranes through binding to their negatively charged constituents.^{24,48-56} For instance, the polycationic lipopeptide colistin, that is, one of the drug of last resort to combat multidrug-resistant (MDR) Pseudomonas aeruginosa infections, acts through binding to lipopolysaccharides (LPS) with displacement of divalent cations that stabilize LPS. 57-63 Insertion of the hydrophobic moiety of the lipopeptide in the membrane causes fusion of the inner leaflets of the outer membrane and the outer leaflet of the cytoplasmic membrane, causing osmotic imbalance and driving the antibiotic into the cytoplasmic membrane through polar and nonpolar channels.

Membrane-targeting antibacterials may use differences in the membrane composition to interact selectively with bacterial cells.^{51,64–69} Unlike most mammalian cell membranes, bacterial membranes are rich in negatively charged lipids (cardiolipin, phosphatidylglycerol, LPS, lipoteichoic acids) that could be

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Figure 1. Structure of the natural antibiotic aminoglycoside neomycin B, of neamine and its amphiphilic derivatives previously prepared.^{24–29}



Figure 2. Comparison of the structures of amphiphilic 3', 4'-dialkyl neamine derivatives and 1- α -allyl-3,4-dialkyl neosamine derivatives (allyl 3,4-dialkyl-6-amino- α -D-glucosaminides) prepared from *N*-acetyl-D-glucosamine.

selectively recognized by cationic amphiphiles (CAs) through ionic interactions and hydrophobic effects.

Membrane-active agents can interact with many targets in the bacterial membranes and inhibit the corresponding functions. Through binding to anionic lipids, they could modify the activity of membrane proteins such as efflux pumps and/or proteins involved in cell division for which a dependence upon negatively charged lipids has been demonstrated. For example, in the regulation of bacterial cell division, proteins MinD/MinE bind tightly to anionic lipids such as cardiolipin positioned at the cell pole, and thus anionic lipids can be useful targets for antimicrobial development.^{29,70,71}

Limited in vitro resistances to these amphiphiles have been observed due to their multiple modes of action.^{48,49,56} Membrane-active agents cannot be inactivated by intracellular bacterial enzymes and flushed out by efflux pumps. In addition, due to their interactions with many key membrane targets present in a great number of copies, biochemical modifications of such multiple targets should have a high cost for the bacteria and should result in a high sensitivity to other antibiotic drugs.

However, the clinical use of CA drugs such as antimicrobial or host defense peptides is limited due to protease susceptibility and toxicity.^{48–50} Over the years, the therapeutic potential of these amphiphiles has been improved by (i) reducing their ability to lyse red blood cells, (ii) increasing selectivity toward bacteria,^{51–55} (iii) reducing nonspecific binding to human serum proteins, and (iv) improving serum stability.⁷²

AAGs are expected to possess improved metabolic stability in regard to peptide-based CA antibiotics and are more difficult to modify by bacterial resistance-causing enzymes than AGs in regard to their expected mode of action. AAGs can also boost the innate immune response, specifically the recruitment of immune cells such as neutrophils required for the resolution of infections and can selectively control inflammatory responses induced in the presence of endotoxins to prevent septic shock.⁴¹

Several strategies for obtaining antibacterial AAGs have been developed including complete or partial conversion of the AG amine and hydroxyl functions into alkyl- or aryl-amide and -ether groups, respectively.²⁴

In our approach in the field of antibacterial AAGs, we assumed that the presence of a large number of amine functions

in AG derivatives like in neomycin 1, which carries six amine functions, can be a source of toxicity through nonspecific binding to the target.^{24–29} Neamine **2** carrying four amine functions is less toxic than neomycin,^{73–76} and the neamine core corresponds to the minimum scaffold necessary for binding to 16S rRNA.^{5,6} This core was used in the synthesis of new AGs in the search for antibacterial and antiviral agents targeting RNA, gene therapy vectors, and for the treatment of the Méniere's disease of the inner ear characterized by recurring attacks of disabling vertigo, hearing loss, and tinnitus.^{10,7} Therefore, for obtaining amphiphilic AGs targeting rRNA, we have modified selectively the small AG neamine 2 on one, two, three, and the four hydroxyl functions in order to keep unchanged the four amine functions potentially protonated at physiological pH, at least partially, in regard to their major role in the binding to anionic targets in bacterial membranes and rRNA. We have identified a first antibacterial amphiphilic neamine derivative, namely 3',4',6-tri2-naphthylmethylene (3',4',6-tri2NM) neamine 3 (Figure 1), having a broad spectrum of activity²⁵ and targeting LPS in the outer membrane of *P. aeruginosa*.²⁶ Structure–activity and structure–cytotoxicity relationships were delineated from various amphiphilic 3',6dialkyl neamine derivatives for obtaining compounds 4-7 (Figure 1) more active than 3 against sensitive and resistant Gram-positive and/or Gram-negative bacteria and strongly less toxic against eukaryotic cells.²⁷ A critical window of lipophilicity appeared to be necessary for optimal antibacterial effects. The study of the mode of action of compounds 5-7 confirmed a strong binding to LPS of P. aeruginosa as well as membrane depolarization.²⁸ Compound 7 has been found the most efficient neamine derivative against Gram-negative bacteria. It appeared to also be able to inhibit growth of P. aeruginosa biofilms and be active against P. aeruginosa strains resistant to colistin, suggesting a different mode of action from the one of colistin.²

In our first report on the identification of the tri-2NM neamine derivative 3, we have described the 3',6- and 3',4'-di2NM neamine derivatives 4 and 8 showing similar antibacterial effects against sensitive and resistant strains of *Staphylococcus aureus* and weak activity against Gram-negative bacteria.²⁵ Herein, we report on the synthesis and the antibacterial activities of new 3',4'-dialkyl neamine derivatives





^{*a*}Reagents and conditions: (a) R = 2NP: NaH/DMF, 3-(2'-naphthyl)propyl bromide (2NPBr),²⁷ rt, 5 h, 44%; R = Nn: TBAF (2 equiv), 50% aq NaOH/toluene, 1-nonyl bromide (1NnBr), rt, 43%. (b) TFA/CH₂Cl₂, anisole, 0 °C; 10, 45%; 11, 55%.

Scheme 2. Preparation of the α -Allyl-3',4'-di2NM Neosamine Derivative 17 and the Corresponding Reference Compound α -Allylneosamine 19^a



^aReagents and conditions: (a) 2NMBr, BEMP, DMF, rt, 48 h, 28%. (b) KOH, EtOH, reflux or $Ba(OH)_2 \cdot 8H_2O$, H_2O , reflux, 8 h, 78%. (c) TrCl, DMF, Et₃N, rt, 8 h, 82%. (d) 2NMBr, NaH, DMF, rt, 10 h, 79%. (e) Ph₃P, THF/H₂O (19/1), 80 °C, 6 h, 91%. (f) TFA/anisole (1/1), 0 °C, 3 h. (g) Dowex resin (Cl⁻ ion exchange); 17, 62%; 19, 84% (3 steps).

active against sensitive and resistant Gram-positive and Gramnegative bacteria in comparison to their active 3',6-dialkyl neamine isomers 5 and 7 previously described.

In the previously identified antibacterial 3',4'-di2NM neamine derivative 8 (Figure 1), both lipophilic groups are attached on the glucosamine ring II. Therefore, small AAGs in which ring II carries at the 3- and 4-positions two lipophilic groups and at the 1-position acyclic side chains generated from an allyl group (Figure 2) could have attractive antibacterial properties. Such an approach has been previously developed from ring I or ring II of neamine in the search for antibiotic aminosugars targeting 16S rRNA.^{97–100}

We have also previously conjugated ring II of neamine to a peptide nucleic acid targeting transactivation response element of HIV-1 RNA genome that shows a high bioavailability in human cells and strongly inhibits Tat-mediated transactivation of HIV-1 transcription.¹⁰¹

Amphiphilic 3,4-dialkyl derivatives of 6-amino-6-deoxyglucosamine named neosamine were synthesized from Nacetylglucosamine. First, derivatives carrying an allyl group introduced in α -configuration at the anomeric position corresponding to the location of ring I on ring II in neamine were prepared (Figure 2). Second, the reactive allyl group of the 3,4-dinonyl neosamine derivative was chemically modified in order to adjust the lipophilicity/hydrophily balance of the resulting AAGs, balance previously identified in the neamine series as a key parameter for obtaining a broad spectrum antibacterial activity. This group was converted to an epoxide ring (oxirane) in order to introduce by ring-opening hydroxyl and/or amine functions like those found in ring I of the corresponding 3',4'-dialkyl neamine derivatives. The route using epoxides as intermediates was selected for the resulting obtention of diasteroisomers that extends the molecular diversity in the antibacterial evaluation. Herein, we compared the antibacterial effects of the prepared 3,4-dialkyl neosamine derivatives to those of the corresponding novel 3',4'-dialkyl neamines described here and to those of the previously described 3',6-dialkyl neamines. *P. aeruginosa* inner membrane permeabilization assays and MIC changes against *P. aeruginosa* induced upon exposure to some of the most active AAGs identified are also reported.

SYNTHESIS

Synthesis of New 3',4'-Dialkyl Neamine Derivatives. In regard to the structure of the 3',6-dialkyl neamine derivatives previously identified as interesting antibacterials, the 3',4'-di-2naphthylpropyl (2NP) and 3',4'-dinonyl (Nn) neamine derivatives were prepared for evaluation of their antibacterial effects (Scheme 1). The *N*-tetratrityl neamine derivative 9^{90} selectively protected at the 6-position by the *p*-methoxybenzyl group was first prepared in good yield under phase transfer conditions with TBAF as a phase transfer agent from *N*tetratritylneamine.⁹⁶ Then, the 3',4'-di2NP (10) and 3',4'-diNn Scheme 3. Preparation of α -Allyl-3,4-di2NP (28) and -DiNn (29) Neosamine Derivatives^a



^aReagents and conditions: (a) TBAF (2 equiv), 50% aq NaOH/toluene or NaH/DMF, C₉H₁₉Br (3 equiv), rt, 5 h, 57%; NaH, DMF, 2NPBr,³² 50 °C, 20 h, 28%. (b) Boc₂O, DMAP, THF, 45 °C, 6 h; **22**, 86%; **23**, 77%. (c) (CH₃O⁻, Na⁺)/CH₃OH, rt, 6 h. (d) PPh₃, THF/H₂O, rt, 4 h; **26**, 83%; **27**, 81% (2 steps). (e) TFA, DCM, rt, 4 h; **28**, 98%; **29**, 97%.

Scheme 4. Synthesis of Azido and Amino Derivatives of the α -Allyl-3,4-diNn Neosamine 27 through Epoxidation and Ring-Opening^a



"Reagents and conditions, PMBnoc = *p*-methoxybenzyloxycarbonyl. (a) *p*-methoxybenzyl-*S*-(4,6-dimethylpyrimidin-2-yl) thiocarbonate (1.2 equiv), DCM, rt, 14 h, 90%. (b) mCPBA (2.5 equiv), DCM, rt, 14 h, 98%. (c) NaN₃ (excess), DMF, 70 °C, 14 h; **32**, 57%; **33**, 29%. (d) PPh₃ (3 equiv), THF/H₂O, rt, 14 h; **35**, 87%; **36**, 83%. (e) TFA, DCM, rt, 4 h; **34**, 98%; **37**, 94%; **38**, 92%.

(11) neamine derivatives were obtained with moderate yields in two steps: (i) alkylation of 9 with the corresponding bromoalkane RBr under phase transfer conditions (R = Nn) or in the presence of NaH in DMF (R = 2NP) and (ii) deprotection with TFA/anisole.

Synthesis in the Neosamine Series. The neosamine core corresponding to ring II in neamine has been previously used in the search for antibacterial agents targeting 16S ribosomal Asite RNA.^{82,97–99} For instance, a library of compounds based upon this core has been synthesized through a combinatorial approach and screened for binding specifically to 16S rRNA by the Wong group to lead to effective binders to a model of A-site 16S RNA in the micromolar range.⁹⁷ One of the key

intermediates used in the different reported approaches has been compound **12**, which was one of our key intermediates in the preparation of amphiphilic neosamine derivatives.

Synthesis of α -Allyl-3,4-di(2'-naphthylmethylene) Neosamine Derivative **17** and of the Corresponding Reference Compound α -Allylneosamine **19**. N-Acetylglucosamine was converted to compound **12** (Scheme 2), possessing an allyl group at the anomeric position in order to allow further modifications, using a three steps sequence adapted from the method described by Wong et al.:⁹⁷ (i) conversion of Nacetylglucosamine to the corresponding α -allyl glycoside, (ii) selective tosylation of the primary alcohol function, and (iii) displacement of the tosyl group with sodium azide. Then the alkylation of 12 with 2-bromomethylnaphthalene (2NMBr) was somewhat troublesome (Scheme 2), and under various basic conditions (NaH or K₂CO₃ or lutidine) was ineffective; using an hindered organic base (BEMP: 2-tert-butylimino-2diethylamino-1,3-dimethylperhydro-1,3,2-diazaphosphorine), compound 13 was obtained in a low 28% yield. The deprotection of the 2-amine function in 13 could not be achieved using either potassium hydroxide in ethanol or barium hydroxide in water at reflux. Thus, compound 12 was first deprotected by refluxing with aqueous barium hydroxide to provide 14 in 78% yield,⁹⁷ and then the free amino group was protected by reaction with trityl chloride to give compound 15 (82%). Then, 15 was alkylated with 2NMBr in DMF and sodium hydride to obtain 16 in 79% yield. The 6-azido group was reduced with triphenylphosphine in a THF/water mixture, and then the trityl group was removed under acidic conditions at 0 °C. A final treatment with an ion-exchange resin afforded the 3,4-di2NM neosamine derivative 17 as a dichlorohydrate (62%, 3 steps). To evaluate the role of the 3,4-dialkyl groups in the antibacterial activity of the prepared amphiphilic neosamine derivatives, the parent compound 1999 was prepared from 15 using the same three-step sequence (76% from 15) through preparation of the intermediate amine 18.

Synthesis of α -Allyl-3,4-di(2'-naphthylpropyl) and -Dinonyl Neosamine Derivatives **28** and **29** (Allyl 3,4-dialkyl-6amino- α -D-glucosaminides). α -Allyl-3,4-dialkyl neosamine derivatives carrying the same alkyl groups as the prepared 3',4'-dialkyl neamines were synthesized for antibacterial evaluation. Two 2-naphthylpropyl and two nonyl groups were introduced at the 3 and 4 positions of α -allyl neosamine starting from compound **12** (Scheme 3). Previously, the removal of the *N*-acetyl group in the 3,4-di2NM intermediate **13** has failed under different drastic basic and nucleophilic conditions and also with other prepared *N*-acetyl-O-3,4-dialkylneosamines. Thus, a Boc group was introduced on the *N*-acetyl group in order to facilitate the final deprotection of the 2-amine function according to the method developed by Chapleur et al. (Scheme 3).¹⁰²

Compound 12 was dialkylated with 2-(3'-bromopropyl)naphthalene²⁷ and 1-bromononane under phase transfer conditions⁹⁶ or in the presence of NaH in DMF, and then the Boc group was introduced to lead to compound 22 and 23, respectively. These compounds were deacetylated in the presence of sodium methoxide to lead to compounds 24 and 25 carrying a 2-NHBoc group. Reduction of the azido group afforded compounds 26 and 27, which were deprotected in the presence of TFA to lead to the AAGs 28 and 29 incorporating two 2NP and two nonyl groups, respectively.

Synthesis of Hydrophilic 3,4-Dialkyl Neosamine Derivatives through Epoxidation of the Allyl Group and Opening of the Obtained Oxiranes with Nucleophiles. According to the higher antibacterial activity against *P. aeruginosa* strains of the 3',6-dinonyl neamine derivative 7 in comparison to the 3',6di2-naphthylpropyl (5) and 3',6-di2-naphthylbutyl (6) derivatives (Figure 1),²⁷ the allyl group found in 3,4-diNn neosamine **29** was modified from compound **27** (Scheme 4) in which the 2-*N*-acetyl group was replaced by a Boc group in order to avoid the *N*-deacetylation difficulty encountered previously with compound **13**, for example (Scheme 2, step b).

In a first step (Scheme 4), the 6-amine function present in 27 was protected in good yield by reaction with a stoichiometric amount of p-methoxybenzyl-S-(4,6-dimethylpyrimidin-2-yl) thiocarbonate.

In a second step, the double bond of the allylic group was oxidized with mCPBA to lead in high yield to the 2:1 diasteroisomeric mixture of (R)- and (S)- α -epoxypropyl derivatives 31a,b characterized by NMR. The configuration of these isomers could not be determined by NOESY experiments at 500 and 600 MHz, and the next steps were performed with the isolated mixture. Such an asymmetric induction has been reported in the epoxidation by mCPBA of β -alkenyl glycosides of O-acetyl- and/or O-benzyl-N-acetyl- β -D-glucosamines.¹⁰³ In this study, the R configuration has been assigned to the major isomers through correlation by ¹H NMR with (R)-glycerinaldehvde. In the mixture of (R)- and (S)- α -epoxides 31a.b. we observed, by ¹H NMR, for one proton of the methylene group in the epoxide ring (C3'-H), a signal more shielded in the minor isomer than in the major isomer (CDCl₃, $\Delta\delta = 0.07$ ppm). Previously, a similar difference in the corresponding chemical shifts (CDCl₃) has been observed after the selective synthesis of (R)- and (S)-(2',3'-epoxy)propyl 2,3,4,6-tetra-Obenzyl- α -D-glucopyranosides by glycosylation with the corresponding chiral 2,3-epoxide alcohols (CDCl₃, $\Delta \delta \approx 0.1$ ppm).¹⁰⁴ Therefore, we assigned the (R) configuration to the major epoxide 31a.

Both (R)- and (S)-epoxides in mixture were opened with sodium azide, 1,2-aminoethanol, 1,3-aminopropanol, 1,2diaminoethane, and 1,3-diaminopropane, respectively, in order to introduce hydrophilic side chains positively charged at physiological pH attached to the anomeric position (Schemes 4 and 5). After opening of the epoxide mixture

Scheme 5. Hydrolysis and Opening of the Isolated 2:1 Epoxide Mixture 31a,b to Lead to Hydrophilic Neosamine Derivatives of AAG $44-47^a$



^aReagents and conditions: (a) TFA/H₂O, rt, 14 h; **39a,b**, 93%. (b) Amine in excess, DCM, 35 °C, 6 h; **40a,b**, 91%; **41a,b**, 99%; **42a,b**, 86%; **43a,b**, 96%. (c) TFA, DCM, rt, 4 h; **44a,b**, 91%; **45a,b**, 93%; **46a,b**, 89%; **47a,b**. 72%.

with sodium azide, it was possible to isolate by chromatography on silica gel the two main products in good yields (57 and 29%). Those compounds were characterized by NMR and mass spectrometry as being two diasteroisomers **32** (major) and **33** (minor) (Scheme 4). However, it was not possible to determine by NMR their configuration. The 2:1 ratio in isolated diasteroisomers was similar to the ratio of both starting epoxides determined by NMR, and thus the (*R*) configuration was assigned to the major isomer **32** according to the epoxide Table 1. clogP Values Calculated Using MarvinSketch Software 5.11.4¹⁰⁵ for the Tetraprotonated Amphiphilic Neamine and Neosamine Derivatives Used in This Study in Comparison to Reference Compounds

antibacterials					
reference antibacterials selected as comparators and 3′,6- and 3′,4′-dialkyl neamines				neosamine derivatives	
compd	clogP	neosamines	clogP	$1-(CH_2-CHOH-CH_2R)-3,4-dinonyl neosamines R =$	clogP
neamine 2	-19.4	α -allyl-3,4-di2NP 28	-0.92	N ₃ 34	-2.83
colistin	-29.7	α-allyl-3,4-diNn 29	-1.39	ОН 39а, b	-3.55
neomycin B 1	-29.9	α -allyl-3,4-di2NM 17	-2.21	NH ₂ 37, 38	-7.04
		lpha-allyl neosamine 19	-8.93	NH(CH ₂) ₃ OH 45a,b	-7.49
Di2NP 5 ²⁷ and 10	-11.4			NH(CH ₂) ₂ OH 44a,b	-7.54
DiNn 7 ²⁷ and 11	-11.9			$NH(CH_2)_3NH_2$ 47a,b	-10.98
Di2NM 4 and 8 ²⁵	-12.7			NH(CH ₂) ₂ NH ₂ 46a,b	-11.03

Table 2. Minimum Inhibitory Concentrations (MIC) of the Neamine and Neosamine Derivatives Synthesized and Some Representative AG against Sensitive and Resistant *Staphylococcus aureus* and *Pseudomonas aeruginosa* Strains^d

	MIC µg/mL							
Aminoglycosides	S. aureus				P. aeruginosa			
	ATCC 25923	SA-1 Pump NorA	ATCC 33592 HA-MRSA	ATCC 27853	Psa. FO3ª	PA22 ^b	PA406 ^c	
Amikacin	4	2	8-16	2-4	8	8	1	
Gentamicin	0.25-0.5	0.5	1	1	>128	4	< 0.25	
Tobramycin	0.25	0.5	64	< 0.25	128	0.5	< 0.25	
Neomycin B 1	1-2	0.5	>128	64	128	32-64	2	
Neamine 2	16-32	8	>128	>128	>128	>128	64	
			Dialkyl neamine	s	10			
3',6-di2NM 4 ²⁵	8	4-8	8-16	64-128	64-128	>128	2-4	
3',4'-di2NM 8 ²⁵	2-4	2-4	4-8	16	128	128	2-4	
3',6-di2NP 5 ²⁷	2	2	2	8-16	16-32	16-32	2	
3',4'-di2NP 10	2-4	0.5-1	1-2	4	8	16	4	
3',6-diNn 7 ²⁷	1-2	2	2-4	2-4	4-8	4	2-4	
3',4'-diNn 11	4	2	4	8-16	8-16	8-16	8-16	
Allyl neosamines								
allyl neosamine 19	>128	>128	>128	>128	>128	>128	>128	
3,4-di2NM 17	8	8	8	>128	>128	>128	32	
3,4-di2NP 28	4-8	4	4-8	128	>128	>128	32	
3,4-diNn 29	4	4	4	>128	>128	>128	>128	
	1- (0	CH2-CHOH-C	H ₂ R)-3,4-Dinon	yl neosamines,	R=			
N ₃ (major) 34	2	2	2	>128	>128	128	>128	
NH2 (major) 37	2	2	2	8	8	8	8	
NH ₂ (minor) 38	2	1	2	4-8	8	8	4-8	
OH 39a,b	2	2	2	64	64	64	64	
NH(CH ₂) ₂ OH 44a,b	4	1-2	1-2	8	8	8	8	
NH(CH ₂) ₃ OH 45a,b	2	1-2	1-2	8	8	8	8	
NH(CH ₂) ₂ NH ₂ 46a,b	2	2	2	8	4-8	4-8	8	
NH(CH2)3NH2 47a,b	2	2	2	8	8	8	4	

^{*a*}Psa.F03 AAC6'-IIA. ^{*b*}Surexp MexXY. ^{*c*}PAO509.5 Δ triABC. ^{*d*}MIC values of AAGs strains lower than or equal to 2–4 μ g/mL against *S. aureus* are highlighted in blue and MIC values equal or lower than 4-8 μ g/mL against *P. aeruginosa* strains are highlighted in yellow.

opening from a nucleophilic attack on the methylene group that was confirmed by NMR (for example DEPTQ experiments). For biological evaluation, the azide **32** was deprotected by treatment with TFA to give the azido AAGs **34** (major *R*). The reduction of azides **32** and **33** with PPh₃/H₂O led to the

corresponding amine 35 and 36, which were deprotected in the presence of TFA to afford AAGs 37 and 38, respectively.

NOE experiments were performed at 500 and 600 MHz with 37 and 38 in CD_3OD . The NOESY spectra revealed only for the minor isomer 38 a correlation between the anomeric C1-proton and the proton carried by the asymmetric 2'-carbon

Table 3. Minimum Inhibitory Concentrations (MIC) of the Neamine and Neosamine Derivatives Synthesized and Some Representative AG against Selected Bacterial Sensitive and Resistant Acinetobacter lwoffi, Escherichia coli, and Klebsiella pneumonia Strains^d

	MIC µg/mL						
Aminoglycosides	A. lwoffi		E. coli			K. pneumonia	
	ATCC 17925	AI.88-483 ^a	ATCC 25922	PAZ505H8101 ^b	L8058.1 ^e	ATCC 700603	
Amikacin	< 0.25	128	4	64	2	1	
Gentamicin	< 0.25	2	0.5	1	64	4-8	
Tobramycin	< 0.25	0.5-1	1	32	64	4-8	
Neomycin B 1	< 0.25	64-128	2	4	1	4-8	
Neamine 2	0.5	>128	32	>128	32	32	
			Dia	lkyl neamines			
3',6-di2NM 4 ²⁵	4-8	128	32-64	32	64-128	128	
3',4'-di2NM 8 ²⁵	4	>128	16-32	8-16	16-32	>128	
3',6-di2NP 5 ²⁷	2	32-64	16	8	16	32	
3',4'-di2NP 10	2-4	32-64	16-32	2	4-8	64	
3',6-diNn 7 ²⁷	1-2	32	2-4	2-4	2-4	2-4	
3',4'-diNn 11	2-4	32	4-8	2-4	4-8	16-32	
	Allyl neosamines						
allyl neosamine 19	>128	>128	>128	>128	>128	>128	
3,4-di2NM 17	8	64	64	64	128	>128	
3,4-di2NP 28	4-8	64	32-64	64	64	128	
3,4-diNn 29	4-8	32	32-64	>128	>128	128	
	1- (CH ₂ -CHOH-CH ₂ R)-3,4-Dinonyl neosamines, R=						
N ₃ (major) 34	4	>128	32	>128	>128	>128	
NH ₂ (major) 37	2	8	4	4	4	8-16	
NH ₂ (minor) 38	2	4	4	4	4-8	8	
ОН 39а, b	1	8	8	16	8	64	
NH(CH ₂) ₂ OH 44a,b	2	4-8	4-8	8	4-8	8	
NH(CH ₂) ₃ OH 45a,b	1	8	4-8	4	4	8-16	
NH(CH ₂) ₂ NH ₂ 46a,b	2	16-32	8	4-8	4-8	8-16	
NH(CH ₂) ₃ NH ₂ 47a,b	2	8	8	4	8	32	

^{*a*}APH3'-VIA. ^{*b*}AAC6'-IB. ^{*c*}ANT2"-IA. ^{*d*}MIC values of AAGs lower than or equal to $4 \mu g/mL$ are highlighted in green and MIC values equal to 4-8 or $8 \mu g/mL$ highlighted in purple.

atom in the side chain, whereas correlations between the H-1 and H-1' were detected for both isomers. Such an additional correlation should be optimal in conformations in which the C1, H1, O1, C1', C2', and H2' atoms are nearly coplanar. In such conformations, the sugar intracyclic oxygen atom and the 2'-oxygen atom of the 2'-hydroxyl group are in a relative trans position in the (S)-isomer minimizing the electronic repulsive effects induced by the oxygen atoms, whereas their relative cis position in the (R)-isomer should disadvantage a NOE correlation. Thus, the observed NOE effect confirmed the (S)-configuration assigned to the minor isomer from the proposed asymmetric induction observed in the epoxidation.

Because **37** and **38** could be used to evaluate the role of the stereochemistry of the side chain in the biological properties, the next synthesis were performed without separation of the diasteroisomers formed in the opening of the mixture of epoxides (Scheme 5). Hydrolysis of the epoxide mixture **31a**,**b** and deprotection were performed in the presence of TFA to lead to the mixture of AAGs **39a**,**b**. The opening of **31a**,**b** with different amines was performed by heating at 35 °C in DCM in the presence of an excess of amine. The reactions with 1,2-

aminoethanol, 1,3-aminopropanol, and with 1,2-diaminoethane led in excellent yields under mild conditions to the mixtures of isomers **40a,b**, **41a,b**, and **42a,b**, respectively, which cannot be separated on TLC. They were deprotected in TFA to afford AAGs **44a,b**, **45a,b**, and **46a,b**, respectively. The opening of the epoxides **31a,b** with 1,3-diaminopropane in DCM led surprisingly to the isomers **43a,b** in which an additional methylene group was observed by ¹H and ¹³C NMR and mass spectrometry. The presence of this methylene group results from cyclization of 1,3-diamino group with the solvent DCM (Scheme 5). It was removed concomitantly with the other carbamate protecting groups by treatment with TFA to give AAG **47a,b**. In the mixtures **40a,b–47a,b**, it was not possible to detect the presence of the diasteroisomers by NMR.

LIPOPHILICITY OF THE SYNTHESIZED DERIVATIVES

In our previous report on the antibacterial activity of amphiphilic neamine derivatives, their lipophilicity expressed by clogP values has been identified to be a key parameter in the

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antibacterial activity as well as in their cytotoxicity. The clogP values of the completely protonated derivatives used in this study were calculated using the MarvinSketch software 5.11.4 and are compared in Table 1.¹⁰⁵

ANTIBACTERIAL ACTIVITY, RESULTS AND DISCUSSION

Three modes of bacterial resistance to AGs have been identified: (i) reduction in the intracellular concentration of the antibiotics by efflux pump proteins or through reduced membrane permeability, (ii) deactivation by AG-modifying enzymes, and (iii) structural modifications of the 16S rRNA binding site that lead to reduced target affinity.⁸⁻¹⁹ Many AGinactiving enzymes that modify the hydroxyl and/or amine functions have been identified and are classified in three families: AG nucleotidyltransferases (ANTs), AG phosphotransferases (APHs), and AG acetyltransferases (AACs).8-Regarding the modifications of 16S rRNA, the methylation of specific nucleotides within the A-site hampers binding of AGs and appeared more and more to be a serious threat to the aminoglycoside antibiotics through the action of plasmidmediated methyltransferases (r-methylases).¹⁶⁻¹⁹ These enzymes that are spreading to different species confer high levels of resistance to clinically useful AG such as amikacin, tobramycin, and gentamicin. In many cases, AG-resistant bacteria have selected combinations of resistance mechanisms that render them very difficult to eradicate.

The synthesized amphiphilic aminoglycosides were evaluated against a large panel of Gram-positive and Gram-negative bacteria. In the former class of bacteria, the mimimum inhibitory concentrations (MIC) were measured against sensitive *Staphylococcus aureus* ATCC 25923 and two resistant strains, the SA-1 strain surexpressing resistance pump (NorA) and the methicillin-resistant strains ATCC 33592 HA-MRSA (Table 2). In the Gram-negative class of bacteria, the effects were evaluated against sensitive and resistant strains of *P. aeruginosa* (Table 2), *Acinetobacter lwoffi*, and *Escherichia coli* surexpressing aminoglycoside-modifying enzymes or efflux pumps and against the sensitive ATCC 700603 *Klebsiella pneumoniae* strain (Table 3).

Antibacterial Activities of the 3',4-Dialkyl Neamine Derivatives 8, 10, and 11. Against the selected *S. aureus* strains (Table 2), the 3',6-2NM neamine 4 showed the highest MIC values (4–16 μ g/mL) and the 3',4'-2NM neamine 8 exhibited lower MIC values (2–8 μ g/mL). The 3,4'-di2NP (10) and -diNn (11) neamine derivatives showed MIC values slightly lower than those measured with the 3',4'-di2NM derivative 8 (MIC = 0.5–4 and 2–8 μ g/mL, respectively). No significant difference in the MIC values were observed between 10 and 11 and their 3',6-isomers 5 and 7, respectively. Among the six evaluated dialkyl neamine derivatives, the best derivatives appeared to be the 3',4'-di2NP derivative 10 reported here.

Against *P. aeruginosa* strains (Table 2), in contrast to the 3',4'-di2NM (8) and 3',6-di2NM (4) derivatives which are inactive, the 3,4'-di2NP 10 showed low MIC values slightly lower than those observed with the 3',4'-diNn derivative 11 (MIC= 4–16 and 8–16 μ g/mL, respectively). As previously observed in the 3',6-dialkyl series, increases of the clogP values from the 2NM to the Nn and 2NP derivatives (Table 1) strongly enhance the antibacterial activity against Gramnegative bacteria. In the 3',4'-series, the most active derivative is the 2NP derivative (10), whereas, in the 3',6-series, the diNn

derivative 7 appeared to be the best anti-*P. aeruginosa* agent. These Nn and 2NP derivatives have close lipophilicities (Table 1). Interestingly, all derivatives showed an activity against *P. aeruginosa* PAO509.5 Δ triABC (MIC = 2–16 µg/mL).

As shown in Table 3, low MIC values against the sensitive strain of A. lwoffi were obtained with the 3',4'-dialkyl derivatives evaluated (MIC = $2-4 \mu g/mL$) as well as with the 3',6-dialkyl derivatives. Higher MIC values against the resistant strain AI.88–483 (MIC = 32 to >128 μ g/mL) were measured and the di2NM derivatives 4 and 8 were found to be inactive (MIC = 128 and >128 μ g/mL, respectively). For the same lipophilic group introduced, the measured MIC values in the 3',4'- and 3',6-series were close. The difference of activity against sensitive and resistant A. lwoffi observed for the 3',4'- and 3',6-dialkyl derivatives is surprising. This difference cannot be related to the overexpression of the APH3'-VIA enzyme in the AI.88-483 strain and the corresponding AAG 3'-modification because an alkyl group is present at the 3'-position. It could be related to difference in the membrane structure of sensitive and resistant A. lwoffi.

Against the sensitive and resistant *E. coli* strains (Table 3), among the three 3',4'-dialkyl neamines evaluated, **8**, **10**, and **11**, the 3',4'-diNn derivative **11** appeared to be the most active (MIC = $2-8 \ \mu g/mL$). Similarly, in the 3',6-series, the diNn derivative 7 showed the lowest MIC values (MIC = $2-4 \ \mu g/mL$) as well as against the selected *P. aeruginosa* strains.

Against sensitive *K. pneumonia*, the 3',4'-di2NM compound 8 and its 3',6-isomer 4 were inactive as well as against the resistant strains of *A. lwoffi* (MIC > 128 μ g/mL). The MIC values measured with the 3',4'-di2NP (10), 3',4'-diNn (11), and 3',6-di2NP (5) neamines appeared to be high (MIC = 16–64 μ g/mL for) in comparison to the MIC obtained with the 3',6-diNn derivative 7 showing low 2–4 μ g/mL MIC.

Antibacterial Activities of α -Allyl-3,4-dialkyl Neosamines 17, 28, and 29 and Hydrophilic 3,4-Dialkyl Neosamines 34, 37–39, and 44–47. The allyl derivatives 17, 28, and 29 showed only a good activity against *S. aureus* and sensitive *A. lwoffi* strains (MIC = 4–8 μ g/mL). The weak or lack of antibacterial activity against the other Gram-negative strains of these compounds should be related to their high lipophilicity (Table 1; clogP = -0.9 to -2.2) as previously observed in the 3',6-dialkyl neamine series (in this series, clogP value have to be lower than –9 for obtaining a good and broadspectrum activity).²⁷ The lack of antibacterial activity of the synthesized hydrophilic parent compound 19 (MIC > 128 μ g/mL) confirmed the essential role of the lipophilic groups.

As shown in Table 2, the neosamine derivatives 34, 37, 38, 39a,b, and 44a,b-47a,b, more hydrophilic than the allyl derivatives 17, 28, and 29 (Table 1), showed close and low MIC values against the S. aureus strains (MIC = $1-4 \mu g/mL$, mainly 2 μ g/mL). The most lipophilic derivatives in this hydrophilic series of neosamine derivatives, the azide 34 (Table 1), appeared to be inactive against the selected Gram-negative bacteria (Tables 2 and 3) except against the sensitive A. lwoffi and *E. coli* strains (MIC = 4 and 32 μ g/mL, respectively). This azide carries one or two amine function(s) protonated at physiological pH less than the other derivatives in the series and its lipophilicity is probably too high (clogP = -2.8) as previously observed in the neamine series.²⁷ Among the less lipophilic neosamine derivatives than 34 (Table 3), the diols 39a,b that are, after 34, the most lipophilic compounds in the series (clogP = -3.6), showed high MIC values against P. aeruginosa and K. pneumonia strains (Tables 2 and 3; MIC = 64 μ g/mL) and lower good MIC values against all *A. lwoffi* and *E. coli* sensitive and resistant strains (Table 3; MIC = 1–8 and 8–16 μ g/mL, respectively).

All other hydrophilic neosamine derivatives **37**, **38**, and **44a,b–47a,b** showed low to medium MIC values against *P. aeruginosa, A. lwoffi, E. coli,* and *K. pneumonia* strains (MIC = 4-8, 1-16, 4-16, and $8-32 \mu g/mL$, respectively). Clearly, the decrease of the lipophilicity (Table 1; clogP = -7 to -11) and the addition of at least one amine function protonated at physiological pH in comparison to the allyl derivatives and the azide **34** strongly increase the antibacterial activity against Gram-negative bacteria. Indeed, at physiological pH, compounds **46** and **47** can bear only three positive charges due to the proximity of the amine functions in their flexible 1-alkyl chain.

Concerning the role in the antibacterial effects observed of the stereochemistry of the hydrophilic side chain attached to the 1-anomeric position (R and S isomers), the evaluation of the isomeric α -aminoalcohols **37** and **38** led to close MIC values showing weak effect on the antibacterial activity of the configuration of the asymmetric 2'-carbon atom found in the 1hydrophilic side chain. Moreover, weak variations in the antibacterial activity were observed by replacement in the hydrophilic side chain of a terminal hydroxyl group by an amine function (from compounds **44a**,**b** to **46a**,**b** and from **45a**,**b** to **47a**,**b**) and by addition a methylene group between the terminal hydroxyl or amine function and the central amino function (from compounds **44a**,**b** to **45a**,**b** and from **46a**,**b** to **47a**,**b**).

Overall, except for **39a,b**, all derivatives retain antimicrobial activity against all sensitive and resistant strains, even those against which conventional AGs are inactive (for examples, gentamicin against *P. aeruginosa* expressing AAC6'-IIA or *E. coli* expressing ANT2"-IA).

CYTOTOXICITY

Using the MTT assay, the viability of murine J774 macrophages were evaluated in the presence of 10 and 30 μ M of the 3',4'-dialkyl neamines (8, 10, and 11) and neosamines (34, 37–39, 44–47) derivatives described here. The viability values were compared in Table 4 to those measured under the same conditions in the presence of the antibacterial 3',6-dialkyl neamines 4, 5, and 7 previously described, of conventional AG drugs and the prepared reference AG in the neosamine series, compound 19, which does not carry lipophilic chains.

At 10 μ M, the lipophilic neosamine derivative **34** carrying an azido group showed the lowest viability (62%) whereas all other evaluated derivatives exhibited viability higher than 78%.

Among them, the less lipophilic neosamine derivatives having good and broad-spectrum antibacterial activity **37–39** and **44–47** showed 85–100% viability.

At 30 μ M, the viability decreased for the majority of the evaluated compounds including conventional AGs drugs with the exception of amikacin (92% viability), the 3',6-di2NP neamine derivative **5** (90% viability), and the neosamine reference compound **19** (82% viability) that has no antibacterial activity.

In the neamine series, the viability strongly decreased from 10 to 30 μ M in the presence of the 3',4'- and 3',6-di2Nn neamine derivatives 11 and 7 (from 83 to 49% and from 78 to 39%, respectively). It decreased to a lower extent in the presence of the other neamine derivatives, from 88 to 67% in the presence of the active 3',4'-di2NP neamine 10 whereas

Table 4. Viability (%) of Murine J774 Macrophages Determined Using the MTT Assay in the Presence of 10 and 30 μ M of the Prepared Neamine and Neosamine Derivatives^a

Aminoglycosides	Viability				
	10 µM	30 µM			
Amikacin	81.2 (3)	92.0 (2)			
Gentamicin	88.2 (9)	74.2 (2)			
Tobramycin	99.7 (4)	83.2 (2)			
Neomycin B 1	87.3 (10)	69.8 (2)			
Neamine 2	94.8 (9)	84.4 (2)			
Dialkyl neamine					
3',6-di2NM 4 25	86.0 (7)	75.9 (1)			
3',4'-di2NM 8 25	79.1 (2)	/			
3',6-di2NP 5 27	91.1 (13)	89.5 (2)			
3',4'-di2NP 10	87.7 (6)	66.9 (2)			
3',6-diNn 7 ²⁷	78.4 (10)	38.8 (2)			
3',4'-diNn 11	82.7 (4)	48.6 (2)			
Allyl neosamines					
allyl neosamine 19	82.2 (5)	82.1 (2)			
3,4-di2NM 17	86.9 (4)	28.3 (2)			
3,4-di2NP 28	105.3 (7)	7.0(1)			
3,4-diNn 29	91.4 (6)	6.6(1)			
1-(CH2-CHOH-CH2R)-3,4-dinonyl neosamines, R=					
N3 (major) 34	62.2 (4)	5.9 (2)			
NH ₂ (major) 37	88.0 (5)	14.4 (1)			
NH ₂ (minor) 38	85.1 (5)	7.2 (1)			
ОН 39а, b	89.7 (4)	8.8 (2)			
NH(CH ₂) ₂ OH 44a,b	99.1 (7)	10.1 (2)			
NH(CH2)3OH 45a,b	89.9 (5)	13.7 (2)			
NH(CH2)2NH2 46a,b	87.5 (4)	53.0 (2)			
NH(CH2)3NH247a,b	100.9 (8)	71.6 (2)			

"The numbers of independent experiences are mentioned after the viability values in brackets.

unchanged in the presence of the 3',6-di2NP isomer 5 (91 and 90% from 10 to 30 μ M).

In the neosamine series, the viability strongly decreases for the majority of the derivatives (17, 28, 29, 34, 37, 38, 39a,b, and 44a,b,45a,b; viability values from 85 to 100% at 10 μ M to 6–28% at 30 μ M). For the tetra-amino derivatives 46a,b and 47a,b, the viability decreased less strongly, from 88 to 53 in the presence of 46a,b and from to 100 to 72% for 47a,b.

Thus, the less cytotoxic AAGs at 30 μ M are the 3',6-di2NM (4), 3',6-di2NP (5), and 3',4'-di2NP (10) neamines (76, 90, and 67% viability, respectively) and the diamino neosamine derivatives 47a,b (72%).

P. AERUGINOSA INNER MEMBRANE PERMEABILIZATION

To investigate if antibacterial activity could be related with a membrane permeabilizing effect, propidium iodide (PI)^{29,106,107} was used to investigate *P. aeruginosa* inner membrane permeabilization induced by 3',6-diNn (7) and 3',4'-diNn (11) neamines and 3,4-diNn neosamine 47a,b (Figure 3). When PI passes through the cell membrane and binds to nucleic acids, fluorescence intensity increases.

The 3',6-diNn neamine derivative (7) induced a dosedependent permeabilization of sensitive *P. aeruginosa* membranes, reaching a plateau value at 5 μ M. This derivative (7)



Figure 3. Inner membrane permeabilization of *P. aeruginosa* induced by 3',6-diNn neamine 7 (∇), 3',4'-diNn neamine 11 (\triangle), and 3,4-diNn neosamine 47a,b (\odot) as assessed by enhancing propidium iodide fluorescence.

induced a higher effect compared to that obtained with 3',4'diNn neamine (11) or 3,4-diNn neosamine 47a,b (75% and 25% permeabilization both, respectively).

Interestingly, these results demonstrated the critical role of the position of the hydrophobic substituents (3', 6 versus 3', 4'), with a higher effect on inner membrane permeabilization induced when the substituents are more distant. When the hydrophobic substituent were adjacent (3',4'-diNn neamine 11 and neosamine derivatives 47a,b), no major difference was observed upon modification of ring I. The small effect on inner membrane permeabilization induced by neamine 11 and neosamines 47a,b is probably sufficient to provoke membrane depolarization and bacterial cell death. A similar behavior has been described for some ceragenins,¹⁰⁸ which showed membrane depolarization and bactericidal effect without effect on inner membrane permeabilization to probes such as Onitrophenyl- β -D-galactoside. Elucidation of the respective roles of membrane permeabilization and depolarization is therefore critical for understanding the mechanisms of action of antimicrobial agents and for providing evidence of the role of membrane integrity in bacterial viability. The absence of correlation between bacterial inner membrane permeabilisation and low values of MIC could be explained by differences in binding to bacterial outer membranes or to other mechanisms like changes in lipid environment required for proper activities of proteins inserted within lipid bilayers.

MIC CHANGES AGAINST P. AERUGINOSA UPON LONG EXPOSURE TO AAGS

To study MIC changes upon exposure to 3',6-di2NP (5) and 3',4'-di2NP (10) neamines in comparison to the fluoroquinolone ciprofloxacin, *P. aeruginosa* ATCC 27853 were grown in the continuous presence of a drug concentration corresponding to half of the MIC.¹⁰⁹

The changes in MIC were observed during the 12 days of exposure to half-MIC concentrations of 3',6-di2NP (5) and 3',4'-di2NP (10) neamines in comparison to ciprofloxacin (Figure 4). Ciprofloxacin induced a marked increase of MIC (~15-fold) at day 4. In comparison, the effect afforded by both di2NP neamines 5 and 10 appeared slower with a 15-fold increase observed after days 7 and 9, respectively.

These results show that exposure of *P. aeruginosa* to subinhibitory concentrations of ciprofloxacin, 3',6-di2NP (5), and 3',4'-di2NP (10) neamines caused a decrease in susceptibility which appears later for new amphiphilic neamine



Figure 4. Evolution of the MIC against *P. aeruginosa* of 3',6-di2NP neamine **5** (dotted line) and 3',4'-di2NP neamine **10** (dots) in comparison with MIC of ciprofloxacin (solid line) after exposure to half-MIC concentrations for the indicated times. The concentration of the antibiotic was readjusted each day to remain equivalent to half the MIC. Results are expressed in changes in MICs over initial value.

derivatives as compared to ciprofloxacin. The resistance mechanisms will be investigated in another study.

DISCUSSION AND CONCLUSION

Comparison of the Antibacterial Activity along the Bacteria Strains and Structure-Activity Relationships. Among the AAGs compared here for their antibacterial activity, the 3',6-diNn neamine 7 previously described is the most active against all selected bacteria strains excepted the resistant A. *lwoffi* strain (MIC = $1-8 \mu g/mL$ against sensitive and resistant S. aureus, P. aeruginosa, E. coli and sensitive A. lwoffi and K. pneumonia and MIC= 32 μ g/mL against the resistant A. lwoffi strain). Its 3',4'-diNn isomer 11 showed similar good activities against sensitive and resistant S. aureus and E. coli strains and sensitive A. lwoffi and lower activities against sensitive and resistant P. aeruginosa strains and sensitive K. pneumonia. The 3',4'-di2NP derivative 10 having close lipophilicity to 7 and 11 has good activities similar to the ones of 7 against sensitive and resistant S. aureus and P. aeruginosa, resistant E. coli and sensitive A. lwoffi, activities stronger than those of its 3,'6di2NP isomer 5.

Concerning the neosamine derivatives prepared, the more hydrophilic 37, 38, 39a,b, 44a,b-47a,b having clogP values between -7 to -11 (Table 1) showed similar good antibacterial activity against all selected sensitive and resistant strains (MIC= $1-16 \ \mu g/mL$). Their antibacterial activities are weaker than the ones of 7 mainly against sensitive *K. pneumonia* (MIC= 8-16 and $2-4 \ \mu g/mL$, respectively) but better against resistant *A. lwoffi* (MIC= 8-16 and $32 \ \mu g/mL$, respectively). They are the most active derivative against this resistant strain.

These results point out the existence of different structure– activity relationships against the selected Gram-negative bacteria strains in regard to the role of ring I and its acyclic scaffold mimics. On the contrary, good and close activities were obtained with neamine and neosamine derivatives against *S. aureus* strains in the three series of compounds studied (3',4'and 3',6-dialkyl neamines and 3,4-dialkyl neosamine derivatives).

Regarding Gram-negative bacteria and compounds having similar lipophilicity near to -11 (Table 1), the results obtained are not very different in the neamine and neosamine series against *P. aeruginosa* and *E. coli* strains. Against *A. lwoffi* and *K. pneumonia* strains, flexibility appears to increase the antibacterial activity.

In the neamine series, the di2NP and diNn derivatives have close clogP values (Table 1). Among them, the 3',4'-di2NP (10) and 3',6-di2NP (5) derivatives show the best activity against *S. aureus* strains. Against *P. aeruginosa* strains, in 3',4'-series, the 2NP and Nn substituents led to similar results whereas, in the 3',6-series, the diNn substituent produces better antibacterial effects than the di2NP group. Better activities of the 3',4'-di2NP (10) and 3',6-diNn (7) neamine derivatives are also observed against *E. coli* strains. A comparative study of the modes of action of these derivatives on bacterial membranes could explain this result that merits attention for future lead selection.

In the neamine and neosamine series, the viability of murine J774 macrophages appeared to be mainly higher or close to 80% at 10 μ M. It decreases significantly at 30 μ M in the neamine series in the presence of the strongly active 3',4'- and 3',6-di2Nn neamine derivatives **11** and 7 and decreases even more drastically in the presence of the active neosamine derivatives **37**, **38**, **39a,b** and **44a,b**-**47a,b**. The viability remains close to 90% at 30 μ M in the presence of the active 3',6-di2NP neamine **5** and decreased to be near to 70% in the presence of the 3',4'-di2NP **10** and the tetra-amino derivatives **47a,b**. These three compounds have strong antibacterial activity against the sensitive and resistant *S. aureus* strains and showed viability near or higher than 90% at 5 to 15-fold the corresponding MIC, respectively.

The amphiphilic hydrophilic neosamine derivatives synthesized extend the spectrum of action of the antibacterial AAGs in the class of Gram-negative bacteria to the resistant *A. lwoffi*. Focusing on *A. lwoffi*, all dialkyl neamine derivatives are weakly active or inactive against the selected resistant strain. The molecular basis of this result probably related to the composition and structure of the membranes of the resistant strain merits to be studied because multidrug-resistant *A. lwoffi* clinical isolates are increasingly reported worldwide.

Concerning the chemical modifications performed in the acyclic scaffolds of the neosamine derivatives, the results point out that (i) the lipophilicity of the derivatives and/or the presence of at least three amine functions are parameters especially critical for a good activity against P. aeruginosa strains, (ii) the clogP values of the antibacterial neosamine derivatives have to be lower than -3.5 for exhibiting a large spectrum of action, and (iii) the modifications made in the acyclic scaffold weakly affect the antibacterial activity in the -7.5 to -11.0 clogP range for compounds having at least at physiological pH one protonated amine function in the side chain. Thus, we can conclude that ring I is the main pharmacophoric element in the antibacterial activity of amphiphilic neamine and neosamine derivatives and that the presence at physiological pH of at least three protonated amine functions and not more are necessary for a good antibacterial activity. This conclusion completes the structure-activity relationships delineated in our previous report, leading to the conclusion that the lipophilicy of dialkyl neamine derivatives have to be included in the window -12.5to -9 for obtaining good antibacterial activities.²⁷ We are not far from this range in the neosamine series.

In conclusion, we show here that the broad spectrum of antibacterial activity observed previously in the 3',6'-dialkyl and 3',4',6-trialkyl neamine series can be extended to the 3',4'-dialkyl neamine and 3,4-dialkyl neosamine series. Membrane permeabilization assays performed with the most active dinonyl derivative in each series (7, 11, and 47a,b) showed permeabilization of the *P. aeruginosa* inner membrane with

emphasis for the 3',6-dinonyl neamine derivatives. Regarding the development of resistance, first measurements of MIC changes against *P. aeruginosa* upon long exposure to the 3',6and 3',4'-di2NP neamine derivatives showed a slower increase of MICs in comparison to ciprofloxacin. The mechanisms of action and resistance will be studied further.

EXPERIMENTAL SECTION

Calculation of clogP Values. The lipophilicity character of the neamine derivatives prepared was estimated through the calculation of clogP values (octanol/water partition coefficients) using the MarvinSketch software [Marvin 5.11.4, 2012, ChemAxon (http://www.chemaxon.com)]. The clogP plug-in in this software calculates the octanol/water partition coefficient, which is used in QSAR analysis and rational drug design as a measure of molecular hydrophobicity. The calculation method used here is based on a modification of the method published by Viswanadhan and Ghose et al. (VG method).¹⁰⁵ The clogP of the substituents were determined through calculation with the same method from the structure of the corresponding alkanes.

Synthesis. General Procedures. Procedure 1. General procedure for the deprotection of the alkylated tetra-N-tritylated neamine and Boc-N-neosamine derivatives. The protected compound was dissolved at 0 °C or at room temperature in CH₂Cl₂/TFA (4/1, v/v). For the tetra-N-tritylated neamines, anisole (0.1 mL/mL) was added. After 2 h stirring at rt, the solvents were evaporated under reduced pressure. H₂O and Et₂O were added and the aqueous phase was washed twice with Et₂O before being evaporated to dryness and then the residue was chromatographed on C18 reversed phase eluting with a H₂O/MeOH gradient and obtained pure as a TFA salt.

Procedure II. General procedure for opening of epoxides **31a**,**b** by amine substrates. To a solution of epoxides **31a**,**b** dissolved in CH₂Cl₂ were added the amines in excess. After 16 h at 35 °C, the solvent was evaporated under reduced pressure. The crude product was diluted with ethyl acetate and washed with water (×3). The dried organic layer was evaporated to dryness. The residue was chromatographed on silica gel with ethyl acetate/methanol (5–10%) to give the β-aminoalcohol derivatives.

Purification. The aminosugar purity of the evaluated compounds was \geq 95%. Before the final deprotection step under acidic conditions, careful purifications of tritylated or Boc derivatives allowing minor isomers or impurities removal were performed by chromatography on silica gel.

The purities were measured by HPLC for the derivatives carrying chromophores and was controlled by ¹H NMR spectrometry and TLC on silica gel (eluent, EtOH/H₂O/(NH₃, H₂O) (20%) 80:10:10; TLC visualizations, sulfuric acid spray (5 mL in 100 mL of EtOH) and ninhydrin spray (0.3 g, 3 mL AcOH, 100 mL of EtOH)). For example, under these TLC conditions, the retardation factors of the 3',6-dinonyl derivative 7 and its 3',4'-isomer 11 were 0.5 and 0.3, respectively. It was not possible to detect by NMR at 400 MHz the presence of two diastereoisomers in compounds 39a,b and 44a,b-47a,b.

3',**4**'-**Di-O-alkylneamines. 3**',**4**'-*Di-O-*(**3**"-(2^m-*naphthyl*)*propyl*)-*neamine* (**10**). To a solution of compound 9^{96} (0.75 g, 0.53 mmol) in dry DMF under argon were added NaH (60%, 213 mg, 5.31 mmol), and then, after 30 min at rt, 2-(3'-bromopropyl)naphthalene (0.53 g, 2.13 mmol). After 1 h stirring at rt, the solvent was evaporated under reduced pressure. The crude product was diluted with ethyl acetate and washed with water (\times 3). The organic layer was dried over MgSO₄ and evaporated to dryness. The residue was chromatographed on silica gel with toluene/ethyl acetate (95:5) to give the N-tetratrityl 3',4'-Odi(2-naphthylpropyl)-6-O-para-methoxybenzyl neamine and 3',4',6-Oalkyl derivatives with 46% yield (0.41 g, 0.24 mmol). HRMS (ESI⁺) *m/z*: [M + K]⁺ calcd, 1785.8319; found, 1785.8327. HRMS (ESI⁺) *m/* z: [M + Na]⁺ calcd, 1769.85797; found, 1769.8570. The deprotection of 0.30 g of this product was achieved following procedure II. 10: 97% yield (0.17g, 0.17 mmol white solid). ¹H NMR (400 MHz, CD₃OD) δ 7.74-7.16 (m, 14H, H ar), 5.89 (d, 1H, J = 3.7 Hz, H-1'), 4.07 (td, 1H, J = 2.5,6.2 Hz, H-5'), 3.95 (t, 1H, J = 6.5 Hz, H-4), 3.88 (dd, 1H, J

= 8.6, 10.4 Hz, H-3'), 3.81–3.75 (m, 1H, CH₂O), 3.72–3.61 (m, 2H, CH₂O), 3.56 (t, 1H, *J* = 9,1 Hz, H-5), 3.52–3.47 (m, 1H, CH₂O), 3.43–3.36 (m, 2H, *H* = 6.3), 3.31–3.26 (m, 2H, H-6', H-2), 3.19–3.11 (m, 2H, H-4, H-1), 3.06 (dd, 1H, *J* = 9.5, 13.1 Hz, H-6'), 2.68 (t, 2H, *J* = 7.4 Hz, CH₂ np), 2.61 (t, 2H, *J* = 7.6 Hz, H-CH₂ np), 2.40 (td, 1H, *J* = 4.1, 12.5 Hz, H-2), 2.02–1.87 (m, 3H, H-2, CH₂), 1.77–1.67 (m, 2H, CH₂). ¹³C NMR (100 MHz, CD₃OD) δ 133.6–135.1 (6C np), 126.3–129.0 (14 CH np), 96.8 (C1'), 81.4 (C4'), 79.2 (C4), 77.8 (C3'), 77.3 (C5), 74.4 (C6), 73.7, 74.1 (2CH₂O), 70.9 (C5'), 54.7 (C2'), 51.5 (C1), 50.1 (C3), 41.6 (C6'), 33.1–33.2 (2CH₂ np), 32.3, 32.6 (2CH₂), 30.0 (C2). HRMS (ESI⁺) *m/z*: [M + H]⁺ calcd 659.3803, found 659.3803. HRMS (ESI⁺) *m/z*: [M + Na]⁺ calcd 681.3622, found 681.3631.

3',4'-Di-O-(1"-nonyl)neamine (11). To a solution of compound 9^{96} (1.73 g, 1.23 mmol) in toluene (85 mL) were added TBAF 3H_2O (1.55 g, 4.92 mmol), 1-bromononane (1.04 mL, 5.55 mmol), and an aqueous solution of NaOH (50% w/w, 42.5 mL). The resulting mixture was stirred vigorously for 24 h at rt. The organic solution was diluted with ethyl acetate and then washed twice with an aqueous saturated ammonium chloride solution before being dried over MgSO4 and evaporated to dryness. The residue was chromatographed on silica gel with toluene/ethyl acetate (95:5) to give the tetratrityl 3',4'-Odinonyl-6-O-para-methoxybenzyl neamine with 43% yield (0.88 g, 0.53 mmol). HRMS (ESI⁺) m/z: [M + K]⁺calcd, 1701.9259; found, 1701.9297. The deprotection of 0.50 g of this product was achieved following procedure II. 11: 98% yield (0.30 g, 0.29 mmol, white solid). ¹H NMR (400 MHz, CD₃OD) δ 5.88 (d, 1H, J = 3.7 Hz, H-1'), 4.06 1H, J = 8.6, 10.4 Hz, H-3'), 3.80–3.73 (m, 1H, CH₂O), 3.72–3.60 (m, 2H, CH₂O), 3.57 (t, 1H, J = 9.1 Hz, H-5), 3.53–3.47 (m, 1H, CH₂O), 3.44-3.35 (m, 2H, H-6, H-3), 3.32-3.25 (m, 2H, H-6a', H-2'), 3.19-3.10 (m, 2H, H-4, H-1), 3.06 (dd, 1H, J = 9.5, 13.1 Hz, H-6'b) 2.41 (td, 1H, J = 4.1, 12.5 Hz, H-2a), 2.04–1.86 (m, 1H, H-2b) 1.79–1.60 (m, 4H, 2CH₂), 1.47–1.28 (m, 24H, 12CH₂), 0.94 (m, 6H, 2CH₂). ¹³C NMR (100 MHz, CD₃OD) δ 96.7 (C1'), 82.3 (C4'), 79.0 (C4), 77.7 (C5), 77.5 (C3'), 75.1 (CH₂O), 74.7 (CH₂O), 73.5 (C6), 71.7 (C5'), 54.7 (C2'), 50.6 (C1), 50.3 (C3), 41.9 (C6'), 33.1 (2CH₂), 31.2 (CH₂), 31.1 (CH₂), 30.7 (4CH₂), 30.4 (2CH₂), 30.1 (C2), 27.1 (CH₂), 26.9 (CH₂), 23.7 (2CH₂), 14.5 (2CH₃). HRMS (ESI⁺) m/z: [M + K]⁺ calcd 597.4562, found 597.4573. HRMS (ESI⁺) m/z: [M + H]⁺ calcd 575.4742, found, 575.4740.

3,4-Di-O-alkylneosamines. Allyl 6-Azido-2,6-dideoxy-2-tritylamino- α -D-glucopyranoside (15). To a stirred solution of 14⁹⁷ (0.3 g, 1.23 mmol) in DMF (8 mL) and Et₃N (0.5 mL) were added Et₃N (0.5 mL) and trityl chloride (1.03 g, 3.68 mmol, 3 equiv) in DMF (3 mL), and the mixture was stirred at room temperature for 8 h under argon atmosphere. A saturated NH₄Cl solution (10 mL) was added, and the mixture was extracted with EtOAc $(3 \times 10 \text{ mL})$. The combined organic layers were dried over MgSO4 and filtered, and the filtrate was concentrated under reduced pressure. The residue obtained was purified by chromatography on silica gel in ethyl acetate/cyclohexane (1:3) with a few drops of Et_3N to give compound 15 (0.49 g, 82%) as a white crystalline solid; mp 128-130 °C. ¹H NMR (400 MHz, CDCl₃) δ 7.54–7.56 (m, 5H ar), 7.23–7.32 (m, 10H ar), 5.81 (m, 1H, CH all), 5.21 (dd, 1H, J = 4.0, 16.0 Hz, CH₂ all), 5.13 (dd, 1H, J = 4.0, 12.0 Hz, CH2 all), 3.75-3.80 (m, 2H, H-3, OCH2), 3.60 (m, 1H, H-5), 3.43-3.48 (m, 2H, H-4, H-6), 3.27-3.38 (m, 2H, H-6', OCH₂), 3.07 (d, 1H, J = 4.0 Hz, H-1), 2.96 (dd, 1H, J = 4.0, 12.0 Hz, H-2). ¹³C NMR (100 MHz, CDCl₃) δ 146.7, 134.0 (CH all), 129.0, 128.2, 126.9, 117.1 (CH₂ all), 97.5 (C-1), 73.9 (C-3), 71.6 (C-4), 70.4 (C-5, CPh₃), 68.7 (OCH₂), 57.6 (C-2), 51.7 (C-6). HRMS (ESI⁺) m/z: [M+K] calcd 525.1941, found 525.1931; [M + Na]⁺ calcd 509.2165, found 509.2164.

Allyl 6-Azido-2,6-dideoxy-3,4-di-O-[(2-naphthyl)methyl]-2-tritylamino- α -D-glucopyranoside (16). To a stirred solution of 15 (1.22 g, 2.51 mmol) in DMF (10 mL) were added NaH (0.4 g, 10.04 mmol, 4 equiv) and 2-bromomethylnaphthalene (2.22 g, 10.04 mmol, 4 equiv). The resulting mixture was stirred at ambient temperature for 10 h under argon atmosphere. A saturated NH₄Cl solution was added, and the mixture was extracted with EtOAc (3 × 10 mL). The combined organic layers were dried over MgSO_4 and filtered, and the filtrate was concentrated under reduced pressure. The residue obtained was purified by chromatography on silica gel in ethyl acetate/cyclohexane (1:45) with a few drops of Et_3N to afford compound 16 (1.52 g, 79%) as a white crystalline solid; mp 118-120 °C. ¹H NMR (400 MHz, CDCl₃) δ 7.17–8.01 (m, 29H ar), 5.80 (m, 1H, CH all), 5.70 (m, 1H, CH₂ np), 5.18–5.23 (m, 2H, CH₂ all, CH₂ np), 5.11–5.14 (m, 2H, CH_2 all, CH_2), 4.76 (m, 1H, CH_2 np), 3.94 (t, 1H, J = 8.0 Hz, H-3), 3.65–3.72 (m, 2H, H-5, OCH₂ all), 3.50 (dd, 1H, J = 8.0, 12.0 Hz, H-4), 3.42 (m, 1H, H-6), 3.14-3.32 (m, 3H, H-2, H-6', OCH₂ all), 2.71 (d, 1H, J = 4.0 Hz, H-1), 2.46 (d, J = 10.7 Hz, 1H). ¹³C NMR (100 MHz, CDCl₃) δ 147.4, 136.6, 135.7, 134.2 (CH all), 133.6, 133.4, 133.1, 133.0, 129.2, 128.4, 128.3, 128.1, 127.9, 127.8, 126.7, 126.6, 126.3, 126.2, 126.1, 126.0, 125.9, 117.0 (CH₂ all), 97.9 (C-1), 83.4 (C-3), 79.7 (C-4), 77.2 (CH₂ np), 75.5 (CH₂ np), 70.8 (CPh₃), 70.0 (C-5), 68.8 (OCH₂ all), 58.3 (C-2), 51.6 (C-6). HRMS (ESI⁺) m/z: [M + Na]⁺ calcd 789.3417, found 789.3434; [M - N2 + Na]⁺ calculated 761.3353, found 761.3339.

Allyl 2,6-Diamino-2,6-dideoxy-3,4-di-O-[(2-naphthyl)methyl]- α -*D-glucopyranoside* (17). To a stirred solution of 16 (0.41 g, 0.53 mmol) in a 19:1 THF/H₂O mixture was added Ph₂P (0.21 g, 0.80 mmol, 1.5 equiv), and the mixture was heated at 80 °C for 8 h. After evaporation to dryness, the resulting residue was purified by chromatography on silica gel in ethyl acetate/cyclohexane (1:4) with a few drops of Et₃N to afford the detritylated compound. This compound was treated with a 1:1 TFA/anisole mixture for 3 h at 0 $^\circ C$ under argon atmosphere. The mixture was co-evaporated twice with toluene. The residue obtained was washed with dry Et_2O (3 × 5 mL) to obtain a yellow compound. This compound was chromatographed on an ion-exchange resin (Dowex resin Cl⁻ form) in methanol to afford the hydrochloride salt of compound 17 (0.19 g, 62% for the 3 steps) as a yellowish white crystalline solid; mp 183–185 °C. ¹H NMR (400 MHz, CD₂OD) δ 7.34–7.81 (m, 14H ar), 6.03 (m, 1H, CH all), 5.43 (m, 1H, CH₂ all), 5.32 (m, 1H, CH₂ all), 5.21 (d, 1H, J = 4.0 Hz, H-1), 5.07 (dd, 2H, CH₂ np), 4.91 (m, 2H, CH₂ np), 4.37 (dd, 1H, J = 8.0, 12.0 Hz, OCH₂ all), 4.20 (dd, 1H, J = 8.0, 12.0 Hz, OCH₂ all), 3.99-4.15 (m, 2H, H-3, H-5), 3.68 (t, 1H, J = 8.0 Hz, H-4), 3.53 (dd, 1H, J = 4.0, 12.0 Hz, H-2), 3.35-3.40 (m, 1H, H-6), 3.04 (m, 1H, H-6′). ¹³C NMR (100 MHz, D₂O) δ 136.7, 136.3, 134.8, 134.7, 134.6 (CH all), 134.5, 129.4, 129.2, 129.1, 128.8, 127.7, 127.4, 127.3, 127.2, 126.8, 126.7, 119.5 (CH2 all), 96.4 (C-1), 81.5 (C-4), 79.5 (C-3), 76.5 (OCH₂ np), 76.4 (OCH₂ np), 70.8 (OCH₂ all), 69.8 (C-5), 54.9 (C-2), 41.8 (C-6). HRMS (ESI⁺) m/z: [M + Na]⁺ calcd 521.2416, found 521.2404; [M + H]⁺ calcd 499.2597, found 499.2594. Elemental Analysis for C₃₁H₃₆N₂O₄ + 1.5H₂O Calcd: C, 62.20; H, 6.57; N, 4.68. Found: C, 62.44; H, 6.30; N, 4.88.

Allyl 6-Amino-2,6-dideoxy-2-N-tritylamino- α -D-glucopyranoside (18). To a stirred solution of 15 (0.22 g, 0.45 mmol) in a 19:1 THF/H₂O mixture was added Ph₃P (0.15 g, 0.59 mmol, 1.3 equiv), and the mixture was heated at 80 °C for 6 h. The solvent was evaporated to dryness, and the resulting residue was purified by chromatography on silica gel in methanol/ethyl acetate (1:9 to 1:1) with a few drops of Et_3N to afford compound 18 (0.19 g, 91%) as a white crystalline solid; mp 139-141 °C. ¹H NMR (400 MHz, CD₃OD) δ 7.66-7.69 (m, 5H ar), 7.15-7.27 (m, 10H ar), 5.85 (m, 1H, CH all), 5.25 (dd, 1H, J = 4.0, 16.0 Hz, CH₂ all), 5.12 (m, 1H, CH₂ all), 3.84 (dd, 1H, J = 4.0, 12.0 Hz, OCH₂), 3.74 (t, 1H, J = 8.0, 12.0 Hz, H-3), 3.58 (m, 1H, H-5), 3.24 (m, 2H, H-6, OCH₂), 3.15 (m, 1H, H-4), 2.90 (dd, 1H, J = 4.0, 12.0 Hz, H-6'), 2.87 (d, 1H, J = 4.0 Hz, H-1), 2.75 (dd, 1H, J = 4.0, 12.0 Hz, H-2). ¹³C NMR (100 MHz, CD₃OD) δ 148.7, 135.5 (CH all), 130.3, 129.0, 127.6, 117.0 (CH₂ all), 99.4 (C-1), 74.5 (C-3), 73.8 (C-4), 71.7 (CPh₃), 69.8 (OCH₂), 69.3 (C-5), 59.2 (C-2), 42.4 (C-6). HRMS (ESI⁺) m/z: $[M + Na]^+$ calcd 483.2260, found 483.2260; [M + H]⁺ calcd 461.2440, found 461.2442.

Allyl 2,6-Diamino-2,6-dideoxy- α -D-glucopyranoside (19).⁹⁹ A solution of 18 (0.11 g, 0.24 mmol) in a 1:1 TFA/anisole mixture was stirred for 6 h at 0 °C under argon atmosphere. The mixture was coevaporated twice with toluene. The residue was washed with dry Et₂O (3 × 5 mL) to obtain a gummy liquid, which was chromatographed on an ion-exchange resin (Dowex resin Cl⁻ form)

in H₂O to afford the hydrochloride salt of compound **19** (0.06 g, 84% for both steps) as a yellow gummy liquid. ¹H NMR (400 MHz, D₂O) δ 5.96 (m, 1H, CH all), 5.36 (dd, 1H, *J* = 4.0, 16.0 Hz, CH₂ all), 5.28 (dd, 1H, *J* = 4.0, 12.0 Hz, CH₂ all), 5.21 (d, 1H, *J* = 4.0 Hz, H-1), 4.26 (m, 1H, OCH₂), 4.05 (m, 1H, OCH₂), 3.85–3.93 (m, 2H, H-4, H-5), 3.36–3.46 (m, 3H, H-2, H-3, H-6), 3.12 (dd, 1H, *J* = 4.0, 12.0 Hz, H-6'). ¹³C NMR (100 MHz, D₂O) δ 133.0 (CH all), 118.7 (CH₂ all), 94.3 (C-1), 71.1 (C-3), 69.4 (C-4), 69.0 (OCH₂ all), 68.0 (C-5), 53.6 (C-2), 40.2 (C-6). HRMS (ESI⁺) *m*/*z*: [M + H]⁺ calcd for C₉H₁₈N₂O₄ 219.1345, found 219.1353.

Allyl 2-Acetamido-6-azido-2,6-dideoxy-3,4-di-O-[(2-naphthyl)propyl]- α -D-glucopyranoside (20). To a solution of compound 12 (1.0 g, 3.49 mmol) in dry DMF (20 mL) under argon were added NaH (60%, 0.42 g, 10.5 mol) and, then, after 30 min at rt the 2naphthylpropyl bromide (1.74 g, 6.99 mmol). After 6 h stirring at rt, the solvent was evaporated under reduced pressure. The crude product was diluted with ethyl acetate and washed with water (x3). The organic layer was dried over MgSO4 and evaporated to dryness. The residue was chromatographed on silica gel with toluene/ethyl acetate (80:20) to give compound 20 with 29% yield (0.64 g, 1.02 mmol, white solid). 20: ¹H NMR (400 MHz, CDCl₃) δ 7.75-7.18 (m, 14H, H-ar), 5.93-5.78 (m, 1H, H-2'), 5.46 (d, 1H, J = 4.2 Hz, NH), 5.23 (dd, 1H, J = 1.2, 17.5 Hz, H-3a'), 5.20 (dd, 1H, J = 1.1, 10.9 Hz, H-3b'), 4,70 (d, 1H, J = 6.5 Hz, H-1), 4.23 (dd, 1H, J = 5.3, 12.8 Hz, H-1a'), 4.03 (dd, 1H, J = 6.0, 12.8 Hz, H-1b'), 3.88 (dd, 1H, J = 3.5, 10.1 Hz, H-2), 3.75–3.69 (m, 1H, H-5), 3.63–3.56 (m, 3H, CH₂O, H-6), 3.50-3.48 (m, 2H, CH₂O), 3.44-3.23 (m, 3H, H-3, H-4, H-6), 2.70-2.55 (m, 4H, CH₂ ar), 1.93–1.76 (m, 7H, 2CH₂, CH₃).¹³C NMR (100 MHz, CDCl₃) δ 168.7 (C=O), 139.5–135.0 (6C ar), 132.1 (C2'), 126.1-128.7 (14CH ar), 118.3 (C3'), 96.8 (C1), 81.2 (C3), 79.02 (C4), 72.5 (C5), 71.1, 68.4 (2CH₂O), 62.3 (C1'), 52.4 (C2), 51.4 (C6), 34.1, 32.5, 32.2, 31.8 (4CH₂), 23.5 (CH₃). HRMS (ESI⁺) m/z: $[M + H]^+$ calcd. 623.3233; found. 623.3229.

Allyl 2-Acetamido-6-azido-2,6-dideoxy-3,4-di-O-nonyl- α -D-glucopyranoside (21). To a solution of compound 12 (1.0 g, 3.49 mmol) in toluene (50 mL) were added TBAF.3H₂O (2.75 g, 8.72 mmol), the 1bromononane (2.66 mL, 13.9 mmol) and an aqueous solution of NaOH (50% w/w, 25 mL). The resulting mixture was stirred vigorously for 16 h at rt. The organic solution was diluted with ethyl acetate and then washed twice with an aqueous saturated ammonium chloride solution before being dried over MgSO4 and evaporated to dryness. The residue was chromatographed on silica gel with toluene/ ethyl acetate (80:20) to give compound 21 with 57% yield (1.069 g, 1.98 mmol white solid). 21: ¹H NMR (400 MHz, CDCl₃) δ5.92-5.82 (m, 1H, H-2'), 5.61 (d, 1H, NH), 5.28 (dd, 1H, J = 1.5, 17.2 Hz, H-3a'), 5.21 (dd, 1H, J = 1.4, 10.4 Hz, H3-b'), 4.78 (d, 1H, J = 3.7 Hz, H-1), 4.22-4.14 (m, 2H, H-2, H-1a'), 3.97 (ddt, 1H, J = 1.2, 6.3, 12,8 Hz, H1b'), 3.81-3,67 (m, 3H, CH₂), 3.54-3,33 (m, 5H, H-3, H-4, H-5, H-6, CH₂), 3.27-3.23 (m, 1H, H-6), 2.00 (s, 3H, CH₃), 1.57-1.46 (m, 4H, CH₂), 1.35-1.19 (m, 24H, CH₂), 0.86 (dt, 6H, J = 3.3, 7.0Hz, CH₃). ¹³C NMR (100 MHz, CDCl₃) δ 169.8 (C=O), 133.6 (C2'), 118.3 (C3'), 97.0 (C1), 81.6 (C3), 79.1 (C4), 73.5, 73.4 (2CH₂O), 71.4 (C5), 68.6 (C1'), 52.7 (C2), 51.6 (C6), 32.1-23.7 $(CH_3, 14CH_2), 22.9 (CH_3), 14.3 (2CH_3). HRMS (ESI⁺) m/z: [M +$ Na]⁺ calcd 561.3992, found 561.3990. HRMS (ESI⁺) m/z: [M + H]⁺ calcd 539.4167, found 539.4167.

Compounds 22 and 23. To a solution of compound 20 or 21 (0.10 g, 0.159 or 0.186 mmol) in THF (6 mL) were added Boc_2O (0.30 g, 1.33 mmol) and DMAP (63 mg, 0.52 mmol). After 6 h at 70 °Cthe solvent was evaporated under reduced pressure. The crude product was diluted with ethyl acetate and washed with water (×3). The organic layer was dried over MgSO₄ and evaporated to dryness. The residue was chromatographed on silica gel with cyclohexane/ethyl acetate (90:10) to give compound 22 with 86% yield (0.10 g, 0.14 mmol colorless oil) or 23 with 88% yield (0.10 g, 0.16 mmol colorless oil).

Allyl 2-[N-(tert-Butoxycarbonyl)acetamido]-6-azido-2,6-dideoxy-3,4-di-O-[(2-naphthyl)propyl]- α -D-glucopyranoside (**22**). ¹H NMR (400 MHz, CDCl₃) δ 7.84–7.30 (m, 14H, H ar), 5.94–5.84 (m, 1H, H-2'), 5.34 (dd, 1H, *J* = 1.3, 17.2 Hz, H-3a'), 5.26 (dd, 1H, *J* = 1.0, 10.4 Hz, H-3b'), 4,90 (d, 1H, *J* = 3.6 Hz, H-1), 5.70 (s, 1H, H-2), 4.52 (t, 1H, J = 9.3 Hz, H-3), 4.20 (dd, 1H, J = 4.9, 13.1 Hz, H-1a'), 4.07 (dd, 1H, J = 6.8, 13.1 Hz, H-1b'), 3.92–3.78 (m, 4H, H-5, CH₂), 3.67–3.61 (m, 1H, CH₂), 3.53 (dd, 1H, J = 2.2, 12.9 Hz, H-6), 3.45 (dd, 1H, J = 6.0, 13.0 Hz, H-6), 3.31–3.27 (m, 1H, H-4), 2.86–2.71 (m, 4H, CH₂), 2.31 (s, 3H, CH₃), 2.01–1.85 (m, 4H, CH₂), 1.54 (s, 9H, CH₃). ¹³C NMR (100 MHz, CDCl₃) δ 172.5 (C=O), 154.9 (COO), 139.3–131.9 (6C ar), 133.7 (C2'), 128.0–125.1 (14CH ar), 118.3 (C3'), 97.3 (C1), 81.6 (C3), 79.6 (C–O), 79.3 (C4), 72.6 (C5), 71.1, 68.5 (2CH₂O), 54.2 (C1'), 54.1 (C2), 51.4 (C6), 32.7, 32.5, 32.0, 31.7 (4CH₂), 23.2 (CH₃). HRMS (ESI⁺) m/z: [M + H]⁺ calcd 723.3758, found 723.3760.

Allyl 2-[N-(tert-Butoxycarbonyl)acetamido]-6-azido-2,6-dideoxy-3,4-di-O-nonyl- α -D-glucopyranoside (23). ¹H NMR (400 MHz, $CDCl_3\delta$ 5.92-5.82 (m, 1H, H-2'), 5.31 (dd, 1H, J = 1.5, 17.2 Hz, H-3a'), 5.23 (dd, 1H, I = 1.4, 10.4 Hz, H3-b'), 4.86 (d, 1H, I = 3.7 Hz, H-1), 4.63 (d, 1H, J = 9.2 Hz, H-2), 4.48-4.39 (m, 1H, H-3), 4.17 (ddd, 1H, J = 3.2, 5.0, 7.3 Hz, H-1a'), 4.05 (dd, 1H, J = 6.8, 13.1 Hz,H-1b'), 3.87-3,69 (m, 4H, H-5, CH₂), 3.58-3,49 (m, 2H, H-6, CH₂), 3.43 (dd, 1H, J = 6.1, 12.9 Hz, H-6), 3.22 (dd, 1H, J = 8.6, 9.9 Hz, H-4), 2.35 (s, 3H, CH₃), 1.64-1.51 (m, 13H, CH₂, CH₃), 1.37-1.25 (m, 24H, CH₂), 0.93 (dt, 6H, J = 3.2, 6.8 Hz, CH₃). ¹³C NMR (100 MHz, $CDCl_3$) δ 172.9 (C=O), 153.9 (COO), 133.5 (C2'), 118.3 (C3'), 96.9 (C1), 83.6 (C-O), 81.0 (C3), 78.4 (C4), 73.2, 71.9 (2CH₂O), 71.0 (C5), 68.3 (C1'), 57.4 (C2), 51.5 (C6), 31,9-22.7 (4CH₃, 14CH₂), 14.1 (2CH₃). HRMS (ESI⁺) m/z: [M + H]⁺ calcd 639.4691, found 639.4687. HRMS (ESI⁺) *m*/*z*: [M + Na]⁺ calcd 661.4512, found 661.4507.

Compounds **24** *and* **25**. To a solution of **22** (138 mg, 0.19 mmol) or **23** (89 mg, 0.14 mmol) in anhydrous MeOH (14 or 9 mL) was added MeONa (15.6 mg, 0.29 mmol or 11.4 mg, 0.21 mmol). After 6 h at rt, the solvent was evaporated under reduced pressure. The crude product was diluted with ethyl acetate and washed with water (×3). The organic layer was dried over MgSO₄ and evaporated to dryness. The residue was chromatographed on silica gel with cyclohexane/ethyl acetate (90:10) to give compound **24** with 96% yield (125 mg, 0.18 mmol, white solid) or **25** with 99% yield (82.3 mg, 0.14 mmol, white solid).

Allyl 6-Azido-2-(tert-butoxycarbonylamino)-2,6-dideoxy-3,4-di-O-[(2-naphthyl)propyl]-α-*D*-glucopyranoside (**24**). ¹H NMR (400 MHz, CDCl₃) δ 7.85–7.31 (m, 14H, H-ar), 6.03–5.93 (m, 1H, H-2'), 5.38 (dd, 1H, *J* = 1.3, 17.2 Hz, H-3a'), 5.30 (dd, 1H, *J* = 1.2, 10.4 Hz, H-3b'), 4,89 (d, 1H, *J* = 3.3 Hz, H-1), 4.83 (d, 1H, *J* = 10.1 Hz, NH), 4.26 (dd, 1H, *J* = 5.3, 12.8 Hz, H-1a'), 4.07 (dd, 1H, *J* = 6.1, 12.8 Hz, H-1b'), 3.98–3.91 (m, 2H, H-6, CH₂), 3.85–3.62 (m, 4H, CH₂O, H-5), 3.58–3.40 (m, 3H, H-3, H-6), 3.35 (t, 1H, *J* = 9.3 Hz, H-4), 2.89–2.77 (m, 4H, CH₂ ar), 2.04–1.94 (m, 4H, 2CH₂), 1.48(s, 9H, CH₃). ¹³C NMR (100 MHz, CDCl₃) δ 155.4 (COO), 139.6–132.0 (6C ar), 133.7 (C2'), 128.0–125.2 (14CH ar), 118.3 (C3'), 97.3 (C1), 81.6 (C3), 79.7 (C–O), 79.3 (C4), 72.6 (C5), 71.1, 68.4 (2CH₂O), 54.2 (C1'), 53.5 (C2), 51.4 (C6), 32.6, 32.5, 32.0, 31.8 (4CH₂), 28.4 (3CH₃). HRMS (ESI⁺) *m*/*z*: [M + H]⁺ calcd 681.3652, found 681.3649.

Allyl 6-Azido-2-(tert-butoxycarbonylamino)-2,6-dideoxy-3,4-di-O-nonyl- α -D-glucopyranoside (25). ¹H NMR (400 MHz, \dot{CDCl}_3) δ 5.99-5.89 (m, 1H, H-2'), 5.34 (dd, 1H, J = 1.5, 17.2 Hz, H-3a'), 5.26 (dd, 1H, J = 1.2, 11.6 Hz, H3-b'), 4.84 (d, 1H, J = 3.4 Hz, H-1), 4.73 (d, 1H, J = 10.8 Hz, NH), 4.21 (dd, 1H, J = 5.3, 12,8 Hz, H-1a'), 4.03 $(dd, 1H, J = 6.1, 12,8 Hz, H-1b'), 3.91-3,82 (m, 3H, H-2, CH_2),$ 3.80–3,70 (m, 2H, H-5, CH₂), 3.63 (dd, 1H, J = 7.3, 15.1 Hz, CH₂), 3.57-3.51 (m, 2H, H-6, CH₂), 3.45-3.38 (m, 2H, H-3, H-6), 3.26 (t, 1H, J = 9.4 Hz, H-4), 1.62–1.54 (m, 4H, CH₂), 1.51 (s, 9H, CH₃), 1.38–1.25 (m, 24H, CH₂), 0.93 (dd, 6H, J = 6.4, 7.1 Hz, CH₃). ¹³C NMR (100 MHz, CDCl₃) δ 155.3 (COO), 133.6 (C2'), 117.7 (C3'), 97.3 (C1), 81.5 (C3), 79.5 (C-O), 79.1 (C4), 73.6, 73.4 (2CH₂O), 71.1 (C5), 68.4 (C1'), 54.2 (C2), 51.4 (C6), 31.9-26.1 (14CH₂), 22.7 (3CH₃), 14.1 (2CH₃). HRMS (ESI⁺) m/z: [M + H]⁺ calcd 597.4586, found 597.4582. HRMS (ESI⁺) m/z: [M + Na]⁺ calcd 619.4405, found 619.4400.

Compounds 26 and 27. To a solution of 24 (100 mg, 0.15 mmol) or 23 (93 mg, 0.16 mmol) in THF (10 mL) were added water (3.3

mL) and PPh₃ (116 mg, 0.44 mmol or 123 mg, 0.47 mmol). After 4 h at rt, the solvent was evaporated under reduced pressure. The crude product was diluted with ethyl acetate and washed with water (×3). The organic layer was dried over $MgSO_4$ and evaporated to dryness. The residue was chromatographed on silica gel with ethyl acetate/MeOH (90:10) to give compound **26** with 86% yield (82.7 mg, 0.13 mmol, white solid) or **27** with 96% yield (85.4 mg, 0.15 mmol, white solid).

Allyl 6-Amino-2-(tert-butoxycarbonylamino)-2,6-dideoxy-3,4-di-O-[(2-naphthyl)propyl]- α -D-glucopyranoside (26). ¹H NMR (400 MHz, CD₃OD) δ 7.72–7.17 (m, 14H, H-ar), 5.97–5.87 (m, 1H, H-2'), 5.32 (dd, 1H, *J* = 1.5, 17.3 Hz, H-3a'), 5.17 (dd, 1H, *J* = 1.2, 10.4 Hz, H-3b'), 4,72 (d, 1H, *J* = 3.4 Hz, H-1), 4.17 (dd, 1H, *J* = 5.1, 13.1 Hz, H-1a'), 3.97 (dd, 1H, *J* = 6.1, 13.1 Hz, H-1b'), 3.75 (dt, 1H, *J* = 6.7, 15.5 Hz, CH₂O), 3.66–3.43 (m, 6H, H-2, H-3, H-5, CH₂O), 3.06–3.00 (m, 2H, H-4, H-6), 2.78–2.60 (m, 5H, H-6, CH₂ ar), 1.86–1.71 (m, 4H, CH₂), 1.34 (s, 9H, CH₃). ¹³C NMR (100 MHz, CD₃OD) δ 158.2 (COO), 141.0–133.5 (6C ar), 135.4 (C2'), 129.0–126.1 (14CH ar), 117.9 (C3'), 98.4 (C1), 82.1 (C3), 82.0 (C4), 80.45 (C–O), 73.7, 73.4 (2CH₂O), 72.2 (C5), 54.2 (C1'), 69.4 (C2), 43.4 (C6), 33.6, 33.4, 33.3, 32.9 (4CH₂), 28.9 (3CH₃). HRMS (ESI⁺) *m*/*z*: [M + H]⁺ calcd 655.3747, found 655.3749.

Allyl⁶ 6-Amino-2-(tert-butoxycarbonylamino)-2,6-dideoxy-3,4-di-O-nonyl-α-D-glucopyranoside (**27**). ¹H NMR (400 MHz, CDCl₃) δ 5.92–5.82 (m, 1H, H-2'), 5.25 (dd, 1H, *J* = 1.5, 17.2 Hz, H-3a'), 5.17 (dd, 1H, *J* = 1.3, 10.4 Hz, H-3b'), 4.74 (d, 1H, *J* = 3.4 Hz, H-1), 4.70 (d, 1H, *J* = 10.0 Hz, NH), 4.13 (dd, 1H, *J* = 5.3, 12,8 Hz, H-1a'), 3.92 (dd, 1H, *J* = 6.0, 12,6 Hz, H-1b'), 3.80–3.45 (m, 6H, H-2, H-5, CH₂O), 3.36 (t, 1H, *J* = 9.6 Hz, H-3), 3.09 (t, 1H, *J* = 9.3 Hz, H-4), 3.00 (d, 1H, *J* = 12.4 Hz, H-6), 2.77 (dd, 6H, *J* = 6.6, 12.6 Hz, H-6), 2.05–1.94 (m, 2H, NH), 1.59–1.48 (m, 4H, CH₂), 1.43 (s, 9H, CH₃), 1.34–1.15 (m, 24H, CH₂), 0.85 (t, 6H, *J* = 6.7 Hz, CH₃). ¹³C NMR (100 MHz, CDCl₃) δ 155.4 (COO), 133.9 (C2'), 117.4 (C3'), 97.3 (C1), 81.6 (C3), 79.9 (C4), 79.4 (C–O), 73.5, 73.3 (2CH₂O), 72.6 (C5), 68.1 (C1'), 54.4 (C2), 42.9 (C6), 31.9–26.1 (14CH₂), 22.7 (3CH₃), 14.1 (2CH₃). HRMS (ESI⁺) *m*/*z*: [M + H]⁺ calcd 571.4686, found 571.4684.

Compounds 28 and 29. The deprotection of 26 (73 mg, 0.11 mmol) or 27 (80 mg, 0.14 mmol) was achieved following procedure I. Allyl 2,6-Diamino-2,6-dideoxy-3,4-di-O-[(2-naphthyl)propyl]- α -Dglucopyranoside (28). Yield 98% (85.6 mg, 0.11 mmol, white solid). ¹H NMR (400 MHz, CD₃OD) δ 7.77–7.20 (m, 14H, H-ar), 6.06– 5.96 (m, 1H, H-2'), 5.40 (dd, 1H, J = 1.4, 17.2 Hz, H-3a'), 5.30 (dd, 1H, J = 1.3, 10.4 Hz, H-3b'), 5.15 (d, 1H, J = 3.2 Hz, H-1), 4.33 (dd, 1H, J = 5.5, 12.5 Hz, H-1a'), 3.97 (dd, 1H, J = 6.0, 13.0 Hz, H-1b'), 3.91-3.79 (m, 2H, H-5, CH₂O), 3.75-3.64 (m, 3H, H-3, CH₂O), 3.61-3.52 (m, 1H, CH₂O), 3.39-3.28 (m, 2H, H-2, H-6), 3.24 (t, 1H, *J* = 9.3 Hz, H-4), 3.10 (dd, 1H, *J* = 9.5, 13.0 Hz, H-6), 2.73 (t, 2H, *J* = 7.4 Hz, CH₂ ar), 2.65 (t, 2H, J = 7.6 Hz, CH₂ ar), 2.02–1.93 (m, 2H, CH₂), 1.85–1.71 (m, 2H, CH₂). ¹³C NMR (100 MHz, CD₂OD) δ 140.4-133.6 (6C ar), 135.1 (C2'), 129.1-126.3 (14CH ar), 119.4 (C3'), 96.0 (C1), 81.7 (C4), 79.4 (C3), 74.2, 73.8 (2CH₂O), 70.4 (C1'), 69.7 (C5), 54.6 (C2), 41.6 (C6), 33.1, 33.1, 32.6, 32.3 (4CH₂). HRMS (ESI⁺) m/z: $[M + H]^+$ calcd 555.3223, found 555.3217.

Allyl 2,6-Diamino-2,6-dideoxy-3,4-di-O-nonyl-α-D-glucopyranoside (**29**). Yield 97% (0.95 g, 0.14 mmol, white solid). ¹H NMR (400 MHz, CD₃OD) δ 5.93–5.83 (m, 1H, H-2'), 5.27 (dd, 1H, *J* = 1.4, 17.2 Hz, H-3a'), 5.15 (dd, 1H, *J* = 1.1, 10.4 Hz, H-3b'), 5.03 (d, 1H, *J* = 3.5 Hz, H-1), 4.20 (dd, 1H, *J* = 5.5, 12.5 Hz, H-1a'), 3.98 (dd, 1H, *J* = 6.5, 12.5 Hz, H-1b'), 3.79–3.63 (m, 3H, H-5, CH₂O), 3.53– 3.44 (m, 3H, H-3, CH₂O), 3.26–3.15 (m, 2H, H-2, H-6), 3.11 (t, 1H, *J* = 9.3 Hz, H-4), 3.00 (dd, 1H, *J* = 9.5, 12.9 Hz, H-6), 1.60–1.43 (m, 4H, CH₂), 1.33–1.10 (m, 24H, CH₂), 0.79 (t, 6H, *J* = 6.8 Hz, CH₃). ¹³C NMR (100 MHz, CD₃OD) δ 135.2 (C2'), 119.3 (C3'), 96.0 (C1), 81.7 (C4), 79.6 (C3), 74.1, 73.8 (2CH₂O), 70.3 (C1'), 69.5 (C5), 54.7 (C2), 40.9 (C6), 32.1–26.0 (14CH₂), 14.1 (2CH₃). HRMS (ESI⁺) *m*/*z*: [M + Na]⁺ calcd 577.3941, found 577.3939.

Allyl 2-(tert-Butoxycarbonylamino)-2,6-dideoxy-6-(p-methoxybenzyloxycarbonylamino)-3,4-di-O-nonyl- α -D-glucopyranoside (**30**). Compound **27** (2.5 g, 4.4 mmol) was dissolved in CH₂Cl₂ (15 mL). p-Methoxybenzyl-S-(4,6-dimethylpyrimidin-2-yl) thiocarbonate (1.47 g, 4.8 mmol) was added, and the reaction was stirred for 16 h at rt. The solvent was evaporated under reduced pressure, and the crude product was chromatographed on silica gel with toluene/ethyl acetate (95:5) to give compound **30** with 90% yield (2.89 g, 3.95 mmol, white paste). ¹H NMR (400 MHz, CDCl₃) δ 7.29 (d, 2H, J = 8.5 Hz, H ar), 6.87 (d, 2H, J = 8.6 Hz, H ar), 5.89–5.79 (m, 1H, H-2'), 5.23 (dd, 1H, *J* = 1.5, 17.2 Hz, H-3a'), 5.17 (dd, 1H, *J* = 1.3, 10.4 Hz, H-3b'), 5.09– 4.95 (m, 3H, CH₂ ar, NH), 4.71 (d, 1H, J = 3.0 Hz, H-1), 4.67 (d, 1H, *J* = 10.0 Hz, NH), 4.05 (dd, 1H, *J* = 5.4, 12,8 Hz, H-1a'), 3.90 (dd, 1H, J = 6.0, 12,7 Hz, H-1b'), 3.80 (s, 3H, CH₃), 3.78-3,63 (m, 3H, H-2, CH₂O), 3.61-3.45 (m, 4H, H-5, H-6, CH₂O), 3.43-3.34 (m, 2 H, H-3, H-6), 3.07 (t, 1H, J = 9.2 Hz, H-4), 1.61–1.48 (m, 4H, CH₂), 1.44 (s, 9H, CH₃), 1.34-1.20 (m, 24H, CH₂), 0.87 (t, 6H, J = 6.8 Hz, CH₃). ¹³C NMR (100 MHz, CDCl₃) δ 159.6 (C ar), 156.4 (COO), 155.4 (COO), 133.7(C2'),129.9 (CH ar), 128.8 (C ar), 117.7 (C3'), 113.9 (CH ar), 97.4 (C1), 81.3 (C3), 79.5 (C4, C-O), 73.7, 73.5 (2CH₂O), 70.3 (C5), 68.4 (C1'), 66.7 (CH₂ ar), 55.3 (CH₃), 54.2 (C2), 41.6 (C6), 31.9-22.7 (14CH₂, 4CH₃), 14.1 (2CH₃). HRMS $(ESI^+) m/z: [M + H]^+$ calcd 735.5154, found 735.5153.

(2'R/S)-2',3'-Epoxypropyl 2-(tert-Butoxycarbonylamino)-2,6-dideoxy-6-(p-methoxybenzyloxycarbonylamino)-3,4-di-O-nonyl- α -Dglucopyranoside (**31***a*,**b**). To a solution of **30** (2.88 g, 3.92 mmol) in CH₂Cl₂ (28 mL) was added mCPBA (2.33 g, 9.8 mmol). After one night of stirring at 75 °C, CH₂Cl₂ (42 mL) was added and the solution was washed with $NaOH_{aq}$ (5%, 50 mL) and twice with water (50 mL). The organic layer was dried over MgSO₄ and evaporated to dryness to give mixture 31a,b with 98% yield (2.89 g, 3.85 mmol). ¹H NMR (400 MHz, CDCl₃) δ 7.29 (d, 2H, J = 8.5 Hz, H ar), 6.87 (d, 2H, J = 8.6 Hz, H ar), 5.13–4.99 (m, 3H, CH₂ ar, NH), 4.73–4.67 (m, 2H, H-1, NH), 3.80 (s, 3H, CH3), 3.78-3,48 (m, 8H, H-2, H-5, H-6, H-1', CH₂O) 3.43-3.29 (m, 3H, H-3, H-6, H-1'), 3.12-3.05 (m, 2H, H-4, H-2'), 2.89-2.81 (m, 1H, H-3'), 2.66 (dd, 0.66H, J = 2.7, 4.9 Hz, H-3a'), 2.59 (dd, 0.33H, J = 2.6, 4.8 Hz, H-3b'), 1.59–1.49 (m, 4H, CH₂), 1.45 (s, 9H, CH₃), 1.32–1.23 (m, 24H, CH₂), 0.88 (dd, 6H, J = 6.3, 7.0 Hz, CH₃). ¹³C NMR (100 MHz, CDCl₃) δ 159.6 (C ar), 156.5 (COO), 155.4 (COO), 130.0 (CH ar), 128.8 (C ar), 113.9 (CH ar), 98.6 (C1a), 98.3 (C1b), 81.2 (C3), 79.7, 79.5 (C4, C-O), 73.8, 73.5 (2CH₂O), 70.5 (C5), 69.0 (C2, C1'), 66.5 (CH₂ ar), 55.3 (CH₃), 50.5 (C2'b), 50.3 (C2'a), 44.5 (C3'a), 44.4 (C3'b), 41.7 (C6), 31.9-22.7 $(14CH_2, 4CH_3)$, 14.1 $(2CH_3)$. HRMS $(ESI^+) m/z$: $[M + H]^+$ calcd 751.5103, found 751.5101.

Compounds 32 and 33. To a solution of 31a,b (99 mg, 0.13 mmol) in DMF (5 mL) was added NaN₃ (260 mg, 3.98 mmol). After one night of stirring at 75 °C, the solvent was evaporated under reduced pressure. The crude product was diluted with ethyl acetate, filtered, and washed with water (×3). The organic layer was dried over MgSO₄ and evaporated to dryness. The residue was chromatographed on silica gel with toluene/ethyl acetate (80:20) to give compound 32 with 57% yield (60 mg, 75 μ mol, white solid) and 33 with 29% yield (30 mg, 38 μ mol, white solid).

3'-Azido-(2'*R*)-2'-hydroxypropyl 2-(tert-Butoxycarbonylamino)-2,6-dideoxy-6-(p-methoxybenzyloxycarbonylamino)-3,4-di-Ononyl-α-D-glucopyranoside (**32**). ¹H NMR (400 MHz, CDCl₃) δ 7.29 (d, 2H, *J* = 8.5 Hz, H ar), 6.87 (d, 2H, *J* = 8.6 Hz, H ar), 5.10– 5.02 (m, 3H, CH₂ ar, NH), 4.77–4.70 (m, 2H, H-1, NH), 3.93–3.85 (m, 1H, H-2'), 3.80 (s, 3H, CH₃), 3.77–3,65 (m, 3H, H-2, CH₂O), 3.62–3.26 (m, 10H, H-3, H-1', H-3', H-5, H-6, CH₂O), 3.07 (t, 1H, *J* = 9.2 Hz, H-4), 1.60–1.49 (m, 4H, CH₂), 1.44 (s, 9H, CH₃), 1.35– 1.20 (m, 24H, CH₂), 0.87 (t, 6H, *J* = 6.8 Hz, CH₃). ¹³C NMR (100 MHz, CDCl₃) δ 159.6 (C ar), 156.5 (COO), 155.5 (COO), 130.0 (CH ar), 128.7 (C ar), 114.0 (CH ar), 98.5 (C1), 81.0 (C3), 79.8 (C– O), 79.5 (C4), 73.7, 73.5 (2CH₂O), 70.5 (C5), 69.7 (C1'), 69.5 (C2'), 66.6 (CH₂ ar), 55.3 (CH₃), 54.3 (C2), 53.6 (C3'), 41.6 (C6), 31.9–22.7 (14CH₂, 4CH₃), 14.1 (2CH₃). HRMS (ESI⁺) *m*/*z*: [M + H]⁺ calcd 794.5274, found 794.5269.

3'-Azido-(2'S)-2'-hydroxypropyl 2-(tert-Butoxycarbonylamino)-2,6-dideoxy-6-(p-methoxybenzyloxycarbonylamino)-3,4-di-Ononyl-α-D-glucopyranoside (**33**). ¹H NMR (400 MHz, CDCl₃) δ 7.30 (d, 2H, J = 8.5 Hz, H ar), 6.88 (d, 2H, J = 8.6 Hz, H ar), 5.065.00 (m, 3H, CH₂ ar, NH), 4.79–4.64 (m, 2H, H-1, NH), 3.96–3.87 (m, 1H, H-2'), 3.80 (s, 3H, CH₃), 3.77–3.67 (m, 3H, H-2, CH₂O), 3.63–3.27 (m, 10H, H-3, H-1', H-3', H-5, H-6, CH₂O), 3.07 (t, 1H, J = 9.2 Hz, H-4), 1.60–1.50 (m, 4H, CH₂), 1.45 (s, 9H, CH₃), 1.35–1.25 (m, 24H, CH₂), 0.87 (t, 6H, J = 6.8 Hz, CH₃). ¹³C NMR (100 MHz, CDCl₃) δ 159.6 (C ar), 156.5 (COO), 155.5 (COO), 130.0 (CH ar), 128.7 (C ar), 114.0 (CH ar), 98.8 (C1), 81.0 (C3), 79.8 (C–O), 79.5 (C4), 73.7, 73.5 (2CH₂O), 70.5 (C5), 69.7 (C1'), 69.6 (C2'), 66.6 (CH₂ ar), 55.3 (CH₃), 54.3 (C2), 53.5 (C3'), 41.7 (C6), 31.9–22.7 (14CH₂, 4CH₃), 14.1 (2CH₃). HRMS (ESI⁺) m/z: [M + H]⁺ calcd 794.5274, found 794.5272.

3'-Azido-(2'*R*)-2'-hydroxypropyl 2,6-diamino-2,6-dideoxy-3,4-di-O-nonyl-α-D-glucopyranoside (**34**). The deprotection of **32** (70 mg, 88.2 μmol) was achieved following procedure II. **34**: 71% yield (52 mg, 68.6 μmol, white solid). ¹H NMR (400 MHz, CD₃OD) δ 5.12 (s, 1H, H-1), 4.07–3.99 (m, 1H, H-2'), 3.96–3.76 (m, 4H, H-5, H-1', CH₂O), 3.75–3.67 (m, 2H, H-3, CH₂O), 3.66–3.56 (m, H-1', CH₂O), 3.43 (d, 2H, *J* = 5.5 Hz, H-3'), 3.38–3.31 (m, 2H, H-2, H-6), 3.25 (t, 1H, *J* = 9.3 Hz, H-4), 3.20–3.09 (m, 1H, H-6), 1.75–1.57 (m, 4H, CH₂), 1.44–1.24 (m, 24H, CH₂), 0.92 (t, 6H, *J* = 6.9 Hz, CH₃). ¹³C NMR (100 MHz, CD₃OD) δ 95.5 (C1), 80.0 (C4), 77.6 (C3), 73.6, 73.2 (2CH₂O), 69.6 (C1'), 68.9 (C2'), 68.4 (C5), 53.4 (C2), 53.2 (C3'), 40.0 (C6), 31.6–22.3 (14CH₂), 12.9 (2CH₃). HRMS (ESI⁺) *m*/*z*: [M + H]⁺ calcd 530.4276, found 530.4277.

Compounds **35** *and* **36**. To a solution of **32** (33 mg, 41.6 μ mol) or **33** (69 mg, 86.9 μ mol) in THF (4 or 8 mL) were added water (1 or 2 mL) and PPh₃ (34 mg, 0.13 mmol or 70 mg, 0.27 mmol). After 4 h at rt, the solvent was evaporated under reduced pressure. The crude product was diluted with ethyl acetate and washed with water (×3). The dried organic layer was evaporated to dryness. The residue was chromatographed on silica gel with ethyl acetate/MeOH (80:20) to give compound **35** with 75% yield (23.9 mg, 31.1 μ mol, white paste) or **36** with 96% yield (35 mg, 45.6 μ mol, white paste).

3'-Amino-(2' Ř)-2'-hydroxypropyl 2-(tert-Butoxycarbonylamino)-2,6-dideoxy-6-(p-methoxybenzyloxycarbonylamino)-3,4-di-Ononyl-α-D-glucopyranoside (**35**). ¹H NMR (400 MHz, CDCl₃) δ 7.29 (d, 2H, J = 8.5 Hz, H ar), 6.87 (d, 2H, J = 8.6 Hz, H ar), 5.27– 4.56 (m, 7H, H-1, CH₂ ar, NH, NH₂), 4.06–3,20 (m, 15H, H-2, H-3, H-5, H-6, H-1', H-2', CH₂O, CH3), 3.14–2.69 (m, 3H, H-4, H-3'), 1.89 (s, NH₂), 1.66–1.37 (m, 13H, CH₂, CH₃), 1.36–1.15 (m, 24H, CH₂), 0.87 (t, 6H, J = 6.8 Hz, CH₃). ¹³C NMR (100 MHz, CDCl₃) δ 159.6 (C ar), 156.5 (COO), 155.6 (COO), 130.0 (CH ar), 128.7 (C ar), 114.0 (CH ar), 98.7 (C1), 81.1 (C3), 79.5 (C–O), 79.4 (C4), 73.7, 73.5 (2CH₂O), 70.3 (C5, C1'), 68.9 (C2'), 66.6 (CH₂ ar), 55.3 (CH₃), 54.4 (C2), 43.3 (C3'), 41.6 (C6), 31.9–22.7 (14CH₂, 4CH₃), 14.1 (2CH₃). HRMS (ESI⁺) *m*/*z*: [M + H]⁺ calcd 768.5369, found 768.5369.

3'-Amino-(2'S)-2'-hydroxypropyl 2-(tert-Butoxycarbonylamino)-2,6-dideoxy-6-(p-methoxybenzyloxycarbonylamino)-3,4-di-Ononyl-α-D-glucopyranoside (**36**). ¹H NMR (400 MHz, CDCl₃) δ 7.29 (d, 2H, *J* = 8.5 Hz, H ar), 6.87 (d, 2H, *J* = 8.6 Hz, H ar), 5.24– 4.96 (m, 4H, CH₂ ar, NH), 4.69 (s, 1H, H-1), 4.15–3,24 (m, 17H, H-2, H-3, H-5, H-6, H-1', H-2', CH₂O, CH₃, NH₂), 3.10–2.64 (m, 3H, H-4, H-3'), 1.96 (s, NH₂), 1.59–1.38 (m, 13H, CH₂, CH₃), 1.35–1.09 (m, 24H, CH₂), 0.87 (t, 6H, *J* = 6.8 Hz, CH₃). ¹³C NMR (100 MHz, CDCl₃) δ 159.6 (C ar), 156.5 (COO), 155.5 (COO), 130.0 (CH ar), 128.7 (C ar), 114.0 (CH ar), 98.7 (C1), 81.1 (C3), 79.5 (C4, C–O), 73.7, 73.5 (2CH₂O), 70.3 (C5, C1'), 69.6 (C2'), 66.6 (CH₂ ar), 55.3 (CH₃), 54.4 (C2), 43.5 (C3'), 41.7 (C6), 31.9–22.7 (14CH₂, 4CH₃), 14.1 (2CH₃). HRMS (ESI⁺) *m*/*z*: [M + H]⁺ calcd 768.5369, found 768.5363.

Compounds **37** and **38**. The deprotection of **35** (21 mg, 27.3 μ mol) or **36** (28 mg, 36.5 μ mol) was achieved following procedure I. 3'-Amino-(2'R)-2'-hydroxypropyl 2,6-Diamino-2,6-dideoxy-3,4-di-O-nonyl- α -D-glucopyranoside (**37**). Yield 95% (21.9 mg, 25.9 μ mol, colorless paste). ¹H NMR (400 MHz, CD₃OD) δ 5.14 (d, 1H, J = 3.2 Hz, H-1), 4.14–4.05 (m, 1H, H-2'), 3.98–3.68 (m, 6H H-3, H-5, H-1', CH₂O), 3.66–3.51 (m, 2H, H-1', CH₂O), 3.36–3.29 (m, 2H, H-2, H-6), 3.27–3.09 (m, 3H, H-4, H-6, H-3'), 3.02 (dd, 1H, J = 9.5, 12.8 Hz, H-3'), 1.70–1.55 (m, 4H, CH₂), 1.42–1.24 (m, 24H, CH₂),

0.91 (t, 6H, J = 6.8 Hz, CH₃). ¹³C NMR (100 MHz, CD₃OD) δ 97.1 (C1), 81.4 (C4), 78.9 (C3), 75.1, 74.7 (2CH₂O), 71.1 (C1'), 69.9 (C5), 67.7 (C2'), 54.9(C2), 43.0 (C3'), 41.5 (C6), 33.1–23.8 (14CH₂), 14.4 (2CH₃). HRMS (ESI⁺) m/z: [M + H]⁺ calcd 504.4371, found 504.4367.

3'-Amino-(2'S)-2'-hydroxypropyl 2,6-Diamino-2,6-dideoxy-3,4di-O-nonyl-α-D-glucopyranoside (**38**). Yield 93% (28.7 mg, 33.9 μmol, colorless paste). ¹H NMR (400 MHz, CD₃OD) δ 5.10 (d, 1H, J = 3.5 Hz, H-1), 4.10–4.01 (m, 1H, H-2'), 3.89 (td, 1H, J = 2.6, 9.6 Hz, H-5), 3.85–3.72 (m, 3H, H-1', CH₂O), 3.71–3.61 (m, 2H, H-3, CH₂O), 3.60–3.47 (m, 2H, H-1', CH₂O), 3.34–3.25 (m, 2H, H-2, H-6), 3.20 (t, 1H, J = 9.3 Hz, H-4), 3.15–3.03 (m, 2H, H-6, H-3'), 2.96 (dd, 1H, J = 9.5, 12.8 Hz, H3'), 1.67–1.51 (m, 4H, CH₂), 1.36–1.21 (m, 24H, CH₂), 0.87 (t, 6H, J = 6.9 Hz, CH₃). ¹³C NMR (100 MHz, CD₃OD) δ 97.2 (C1), 81.4 (C4), 79.0 (C3), 75.0, 74.6 (2CH₂O), 71.3 (C1'), 69.8 (C5), 67.8 (C2'), 54.8 (C2), 42.8 (C3'), 41.5 (C6), 33.1–23.8 (14CH₂), 14.4 (2CH₃). HRMS (ESI⁺) *m*/*z*: [M + H]⁺ calcd 504.4371, found 504.4365.

(2'R/S)-2',3'-Dihydroxypropyl 2,6-Diamino-2,6-dideoxy-3,4-di-Ononyl- α -D-glucopyranoside (**39a**,**b**). The mixture **31a**,**b** (198 mg, 0.26 mmol) was dissolved in CH₂Cl₂/TFA (4/1 mL) in the presence of water (2 mL). After 4 h stirring, the solvents were evaporated under reduced pressure. MeOH (2 \times 5 mL) and H₂O (2 \times 5 mL) were added and evaporated. Then the residue was chromatographed on C18 reversed phase eluting with a H₂O/MeOH (55:45). 39a,b were obtained with 93% yield as a TFA salt (180 mg, 0.25 mmol, colorless solid). **39a,b**: δ 5.05 (dd, 1H, I = 3.2, 7.4 Hz, H-1), 3.93–3.76 (m, 5H, H-5, H-1', CH₂O), 3.73-3.11 (m, 6H, H-3, H-2', H-3', CH₂O), 3.36-3.28 (m, 1H, H-6), 3.23-3.02 (m, 3H, H-2, H-4, H-6), 1.72-1.75 (m, 4H, CH₂), 1.42-1.24 (m, 24H, CH₂), 0.90 (t, 6H, J = 7.0 Hz, CH₃). ¹³C NMR (100 MHz, CD₃OD) δ 97.4 (C1), 81.6 (C4), 80.0 (C3), 75.0, 74.6 (2CH₂O), 71.7 (C1'), 70.3 (C2'), 69.8 (C5), 63.5 (C3'), 55.3 (C2), 41.5 (C6), 33.1-23.8 (14CH₂), 14.4 (2CH₃). HRMS (ESI⁺) m/z: $[M + H]^+$ calcd 505.4211, found 505.4211.

3'-(2"-Hydroxyethylamino)-(2'R/S)-2'-hydroxypropyl 2-(tert-Butoxycarbonylamino)-2,6-dideoxy-6-(p-methoxybenzyloxycarbonylamino)-3,4-di-O-nonyl- α -D-glucopyranoside (40a,b). The opening of 31a,b (200 mg, 0.27 mmol) by ethanolamine (2 mL) was achieved following procedure II. 40a,b: 91% (197 mg, 0.24 mmol, colorless paste). ¹H NMR (400 MHz, CDCl₃) δ 7.25 (d, 2H, J = 8.5 Hz, H ar), 6.83 (d, 2H, J = 8.6 Hz, H ar), 5.74-5.17 (m, NH), 5.07-4.92 (m, 2H, CH₂ ar, NH), 4.67–4.60 (m, 1H, H-1, NH), 3.87–3.20 (m, 17H, H-2, H-3, H-5, H-6, H-1', H-2', H-5', CH₂O, CH₃, NH), 3.06-2.98 (m, 1H, H-4), 1.94 (N-H), 2.77-2.52 (m, 4H, H-3', H-4'), 1.57-1.35 $(m, 13H, CH_2, CH_3), 1.33-1.17 (m, 24H, CH_2), 0.84 (t, 6H, J = 6.8)$ Hz, CH₃). ¹³C NMR (100 MHz, CDCl₃) δ 159.5 (C ar), 156.6 (COO), 155.7 (COO), 129.9 (CH ar), 128.7 (C ar), 113.9 (CH ar), 98.6, 98.5 (C1), 81.0 (C3), 79.5 (C-O), 79.4 (C4), 73.5 (2CH₂O), 70.6(C1'), 70.2 (C5), 68,6 (C2'), 66.5 (CH₂ ar), 60.8 (C5'), 55.3 (CH₃), 54.4 (C2), 52.0 (C3'), 51.4 (C4'), 41.7 (C6), 31.9-22.7 $(14CH_2, 4CH_3), 14.1 (2CH_3)$. HRMS $(ESI^+) m/z$: $[M + H]^+$ calcd 812.5631, found 812.5638

3'-(3"-Hydroxypropylamino)-(2'R/S)-2'-hydroxypropyl 2-(tert-Butoxycarbonylamino)-2,6-dideoxy-6-(p-methoxybenzyloxycarbonylamino)-3,4-di-O-nonyl- α -*p*-glucopyranoside (41a,b). The opening of 31a,b (200 mg, 0.27 mmol) by propanolamine (2 mL) was achieved following procedure II. 41a,b: 99% yield (217 mg, 0.26 mmol, colorless paste). ¹H NMR (400 MHz, CDCl₃) δ 7.22 (d, 2H, J = 8.5 Hz, H ar), 6.80 (d, 2H, J = 8.6 Hz, H ar), 5.99-5.24 (m, NH), 4.99-4.91 (m, 2H, CH₂ ar, NH), 4.61-4.57 (m, 1H, H-1, NH), 4.06-3.95 (m, 1H, H2'), 3.80-3.25 (m, 15H, H-2, H-3, H-5, H-6, H-1', CH₂O, CH₃, NH), 3.10–2.72 (m, 5H, H-4, H-3', H-4'), 1.89–1.73(m, 2H, H-5'),1.53-1.31 (m, 13H, CH₂, CH₃), 1.28-1.08 (m, 24H, CH₂), 0.80 (dd, 6H, J = 5.4, 6.9 Hz, CH₃). ¹³C NMR (100 MHz, CDCl₃) δ 159.5 (C ar), 156.6 (COO), 156.0 (COO), 129.9 (CH ar), 128.7 (C ar), 113.9 (CH ar), 98.9, 98.8 (C1), 81.0 (C3), 79.6 (C-O), 79.2 (C4), 73.6, 73.5 (2CH₂O), 70.2 (C5, C1'), 66.7 (C2'), 66.5 (CH₂ ar), 61.1 (C6'), 55.3 (CH₃), 54.5 (C2), 51.1 (C3'), 47.9 (C4'), 41.7 (C6), 32.1-22.7 (C5', 14CH₂, 4CH₃), 14.1 (2CH₃). HRMS (ESI⁺) m/z: [M + H]⁺ calcd 826.5787, found 826.5794.

3'-(2"-Aminoethylamino)-(2'R/S)-2'-hydroxypropyl 2-(tert-Butoxycarbonylamino)-2,6-dideoxy-6-(p-methoxybenzyloxycarbonylamino)-3,4-di-O-nonyl- α -D-qlucopyranoside (42a,b). The opening of 31a,b (200 mg, 0.27 mmol) by ethylenediamine (2 mL) was achieved following procedure II. 42a,b: 86% yield (186 mg, 0.23 mmol, colorless paste). ¹H NMR (400 MHz, CDCl₃) δ 7.26 (d, 2H, I = 8.5Hz, H ar), 6.85 (d, 2H, J = 8.6 Hz, H ar), 5.34–4.81 (m, 2H, CH₂ ar, NH), 4.68 (s, 1H, H-1), 3.86-3.27 (m, 15H, H-2, H-3, H-5, H-6, H-1', H-2', CH₂O, CH₂, NH), 3.09-2.41 (m, 7H, H-4, H-3', H-4', H-5'), 1.58–1.38 (m, 13H, CH₂, CH₃), 1.32–1.15 (m, 24H, CH₂), 0.85 (dd, 6H, J = 6.0, 7.0 Hz, CH₃). ¹³C NMR (100 MHz, CDCl₃) δ 159.5 (C ar), 156.6 (COO), 155.5 (COO), 130.0 (CH ar), 128.8 (C ar), 113.9 (CH ar), 98.6, 98.5 (C1), 81.3 (C3), 81.2 (C-O), 79.5 (C4), 73.6, 73.5 (2CH₂O), 71.0 (C1'), 70.3 (C5), 68,5 (C2'), 66.5 (CH₂ ar), 56.4 (C4'),55.3 (CH₃), 54.4 (C2), 52.0 (C3'), 41.7 (C6), 41.3 (C5'), 31.9–22.7 (14CH₂, 4CH₃), 14.1 (2CH₃). HRMS (ESI⁺) m/z: $[M + H]^+$ calcd 811.5791, found 811.5788.

3'-(Tetrahydropyrimidin-1(2H)-yl)-(2'R/S)-2'-hydroxypropyl 2-(tert-Butoxycarbonylamino)-2,6-dideoxy-6-(p-methoxybenzyloxycarbonylamino)-3,4-di-O-nonyl- α -D-glucopyranoside (43a,b). The opening of 31a,b (195 mg, 0.26 mmol) by 1,3-diaminopropane (2 mL) was achieved following procedure II. **43a,b**: 82% yield (178 mg, 0.21 mmol, colorless paste). ¹H NMR (400 MHz, $CDCl_3$) δ 7.26 (d, 2H, J = 8.5 Hz, H ar), 6.84 (d, 2H, J = 8.6 Hz, H ar), 5.37-4.92 (m, 2H, CH₂ ar, NH), 4.66 (s, 1H, H-1), 4.27-3.85 (m, 1H, H2', NH), 3.80-3.20 (m, 17H, H-2, H-3, H-5, H-6, H-1', H-7', CH₂O, CH₃, NH), 3.13-2.49 (m, 7H, H-4, H-3', H-4', H-6'), 1.94-1.73 (m, 2H, H-5'), 1.57-1.35 (m, 13H, CH₂, CH₃), 1.33-1.03 (m, 24H, CH₂), 0.87-0.82 (m, 6H, CH₃). ¹³C NMR (100 MHz, CDCl₃) δ 159.5 (C ar), 156.5 (COO), 155.6 (COO), 129.9 (CH ar), 128.8 (C ar), 113.9 (CH ar), 98.7 (C1), 81.1 (C3), 79.5 (C-O), 79.4 (C4), 73.6, 73.5 (2CH₂O), 70.2 (C1'), 70.0 (C5), 66.9 (C2'), 66.5 (CH₂ ar), 56.7 (NCH₂N), 56.6 (C3'), 56.3 (C7'), 55.3 (CH₃), 54.4 (C2), 51.6 (C4'), 43.3 (C6'), 41.7 (C6), 31.9-22.7 (C5', 14CH₂, 4CH₃), 14.1 (2CH₃). HRMS (ESI⁺) m/z: [M + H]⁺ calcd 825.5947, found 825.5943.

3'-(2"-Hydroxyethylamino)-(2'R/S)-2'-hydroxypropyl 2,6-Diamino-2,6-dideoxy-3,4-di-O-nonyl-α-D-glucopyranoside (44a,b). The deprotection of 40a,b (195 mg, 0.24 mmol) was achieved following procedure I. 44a,b: 91% yield (194 mg, 0.22 mmol, colorless paste). ¹H NMR (400 MHz, CD₃OD) δ 5.13 (d, 1H, *J* = 3.5 Hz, H-1), 4.23–4.14 (m, 1H, H-2'), 3.97–3.66 (m, 8H, H-3, H-5, H-1', H-5', CH₂O), 3.62–3.52 (m, 2H, H-1', CH₂O), 3.39–3.28 (m, 3H, H-2, H-6, H-3'), 3.27–3.07 (m, 5H, H-4, H-6, H-3', H-4'), 1.76–1.53 (m, 4H, CH₂), 1.41–1.23 (m, 24H, CH₂), 0.90(t, 6H, *J* = 7.0 Hz, CH₃). ¹³C NMR (100 MHz, CD₃OD) δ 95.9, 95.8 (C1), 80.1, 80.0 (C4), 77.7, 77.6 (C3), 73.7, 73.2 (CH₂O), 70.0, 69.8 (C1'), 68.5 (C5), 65.5, 65.3 (C2'), 56.3 (C5'), 53.5 (C2), 49.3, 49.2, 49.1 (C3', C4'), 40.1 (C6), 31.7–22.3 (14CH₂), 13.0 (2CH₃). HRMS (ESI⁺) *m/z*: [M + H]⁺ calcd 548.4633, found 548.4634.

3'-(3"-Hydroxypropylamino)-(2'R/S)-2'-hydroxypropyl 2,6-Diamino-2,6-dideoxy-3,4-di-O-nonyl-α-D-glucopyranoside (45a,b). The deprotection of 41a,b (215 mg, 0.29 mmol) was achieved following procedure I. 45a,b: 93% yield (241 mg, 0.27 mmol, colorless paste). ¹H NMR (400 MHz, CD₃OD) δ 5.14 (d, 1H, *J* = 3.5 Hz, H-1), 4.19–4.14 (m, 1H, H-2'), 3.97–3.69 (m, 8H, H-3, H-5, H-1', H-6', CH₂O), 3.64–3.51 (m, 2H, H-1', CH₂O), 3.37–3.09 (m, 8H, H-2, H-4, H-6, H-3', H-4'), 2.00–1.90 (m, 2H, H-5'), 1.73–1.56 (m, 4H, CH₂), 1.42–1.25 (m, 24H, CH₂), 0.94–0.87 (m, 6H, CH₃). ¹³C NMR (100 MHz, CD₃OD) δ 97.1 (C1), 81.5 (C4), 78.9 (C3), 75.1, 74.7 (CH₂O), 71.1 (C1'), 69.9 (C5), 66.8 (C2'), 60.7 (C6'), 54.9 (C2), 50.9 (C3'), 47.8 (C4'), 41.5 (C6), 33.1–23.7 (C5', 14CH₂), 14.4 (2CH₃). HRMS (ESI⁺) *m*/*z*: [M + H]⁺ calcd 562.4790, found 562.4791.

3'-(2"-Aminoethylamino)-(2'R/S)-2'-hydroxypropyl 2,6-Fiamino-2,6-dideoxy-3,4-di-O-nonyl-α-D-glucopyranoside (**46a**,**b**). The deprotection of **42a**,**b** (181 mg, 0.22 mmol) was achieved following procedure I. **46a**,**b**: 89% yield (199 mg, 0.20 mmol, colorless paste). ¹H NMR (400 MHz, CD₃OD) δ 5.13 (d, 1H, *J* = 3.5 Hz, H-1), 4.23– 4.14 (m, 1H, H-2'), 3.96–3.66 (m, 6H, H-3, H-5, H-1', CH₂O), 3.63– 3.51 (m, 2H, H-1', CH₂O), 3.45–3.27 (m, 7H, H-2, H-6, H-3', H-4', H-5'), 3.27–3.16 (m, 2H, H-4, H-3'), 3.11 (dd, J = 9.6, 13.0 Hz, H-6), 1.70–1.56 (m, 4H, CH₂), 1.41–1.22 (m, 24H, CH₂), 0.90 (t, 6H, J = 6.9 Hz, CH₃). ¹³C NMR (100 MHz, CD₃OD) δ 97.1 (C1), 81.5 (C4), 78.9 (C3), 75.0, 74.6 (CH₂O), 71.0 (C1'), 69.8 (C5), 67.1 (C2'), 54.8 (C2), 51.5 (C3'), 46.0 (C4'), 41.5 (C6), 37.0 (C5'), 33.0–23.7 (14CH₂), 14.4 (2CH₃). HRMS (ESI⁺) m/z: [M + H]⁺ calcd 547.4793, found 547.4793.

3'-(3"-Aminopropylamino)-(2'R/S)-2'-hydroxypropyl 2,6-diamino-2,6-dideoxy-3,4-di-O-nonyl-α-D-glucopyranoside (47a,b). The deprotection of 43a,b (100 mg, 119.5 μmol) was achieved following procedure I. 47a,b: 72% yield (88 mg, 86.5 μmol, colorless paste). ¹H NMR (400 MHz, CD₃OD) δ 5.13 (d, 1H, *J* = 3.5 Hz, H-1), 4.21–4.14 (m, 1H, H-2'), 3.97–3.66 (m, 6H, H-3, H-5, H-1', CH₂O), 3.64–3.48 (m, 2H, H-1', CH₂O), 3.40–3.02 (m, 10H, H-2, H-4, H-6, H-3', H-4', H-6'), 2.19–1.05 (m, 2H, H-5'), 1.71–1.56 (m, 4H, CH₂), 1.42–1.22 (m, 24H, CH₂), 0.90 (t, 6H, *J* = 6.9 Hz, CH₃). ¹³C NMR (100 MHz, CD₃OD) δ 97.2 (C1), 81.5 (C4), 79.0 (C3), 75.1, 74.6 (CH₂O), 71.1 (C1'), 69.8 (C5), 66.9 (C2'), 54.9 (C2), 51.2, 51.1 (C3'), 46.0 (C4'), 41.5 (C6), 37.9 (C6'), 33.1–23.7 (C5', 14CH₂), 14.4 (2CH₃). HRMS (ESI⁺) *m*/*z*: [M + H]⁺ calcd 562.4790, found 562.4791.

Biological and Biochemical Assays. *MIC Determination.* All strains were grown overnight at 37 $^{\circ}$ C on trypticase soy agar (TSA) petri dishes (BD Diagnostics, BD, Franklin Lakes, NJ). MICs were determined by microdilution using a fresh culture in cation-adjusted Mueller–Hinton broth (CA-MHB) and a starting inoculum of 10⁶ cells, according to the recommendations of the Clinical and Laboratory Standards Institute (CLSI).¹¹⁰

Assessment of Eukaryotic Cell Viability. Cell viability was assessed by evaluating their metabolic activity using the MTT assay (reduction in mitochondria of 3-[4,5-dimethylthiazol-2-yl]-2,5-diphenyl tetrazolium in purple formazan crystal).^{111,112} Briefly, cells exposed for 24 h to 10 or 30 μ M of compounds were incubated for 1 h with 0.2 mg/mL MTT. The reaction was stopped by addition of dimethyl sulfoxide (DMSO). The OD was measured at 570 nm.

P. aeruginosa Inner Membrane Permeabilization. *Bacterial* Strain and Growth Conditions. Trypticase soy agar (TSA) medium was used to grow *P. aeruginosa* ATCC 27853 overnight at 37 °C. One colony was suspended in cation-adjusted Müller–Hinton Broth (MHB-CA) and incubated overnight at 37 °C on a rotary shaker (130 rpm). The bacterial suspension was diluted 100-fold in MHB-CA and incubated (130 rpm; 37 °C; 4 h) until it reached the mid logarithmic (mid log) phase ($OD_{620} \sim 0.4-0.5$). *Membrane Permeabilization*.²⁹ Permeabilization of the inner

*Membrane Permeabilization.*²⁹ Permeabilization of the inner bacterial membrane was studied with a membrane-impermeable fluorescent dye (propidium iodide). A stock solution of PI (3 mM in pure water) was diluted 10³-fold with the bacterial suspension (OD₆₀₀: 0.05). 3',6-DiNn (7) and 3',4'-diNn (11) neamines and 3,4diNn neosamine **47a,b** in HEPES buffer, at final concentrations ranging from 1 to 10 μ M, were added to the propidium iodidecontaining bacterial suspension in 96-well microplates. The fluorescence intensity was measured with a SpectraMax M3 microplate reader (Molecular Devices, Sunnyvale, CA, USA) at 25 °C after 15 min of stabilization at excitation and emission wavelengths of 540 and 610 nm, respectively. The data were normalized based on the fluorescence intensity measured in the presence of either quaternary ammonium hexadecyltrimethylammonium bromide (CTAB) 150 μ M (positive control, 100%)¹⁰⁷ or imipenem (negative control, 0%). *Resistance Selection Method*.¹⁰⁹ A serial passage method was used

Resistance Selection Method.¹⁰⁹ A serial passage method was used in which *P. aeruginosa* ATCC 27853 were grown in the continuous presence of a drug concentration corresponding to half of the MIC. The bacteria were examined daily for a change in MIC, followed by a corresponding increase of the drug concentration for up to 12 days. For this purpose, an initial inoculum of 2.5×10^6 cfu/mL of each of the original strain was exposed in broth to a range of antibiotic concentrations from 0.1 to 5–10-fold their original MIC (using arithmetic increases). After 24 h at 37 °C, the tubes were examined to determine the minimal drug concentration preventing bacterial growth (this value was equal to the MIC determined on an agar plate for the corresponding strain). Bacteria growing at a drug concentration of half this value were then readjusted at a density of 2.5×10^6 cfu/mL and

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again exposed for 24 h to drug concentrations from 0.1 to 5-10-fold the MIC. This process was repeated each day, looking for growth at drug concentrations larger than the original MIC. If this was observed, the new minimal drug concentration preventing bacterial growth was determined based on visual inspection of the cultures (and samples taken for confirmation of the MIC by agar dilution). Bacteria growing at a concentration corresponding to half of this new value were then used for continuation of the experiment, for a total of 12 days.

ASSOCIATED CONTENT

S Supporting Information

The Supporting Information is available free of charge on the ACS Publications website at DOI: 10.1021/acs.jmed-chem.6b00818.

General information for the synthesis, ¹H and ¹³C NMR spectra, and purities of the evaluated amphiphilic derivatives (PDF)

Molecular formula strings (CSV)

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Notes

The authors declare no competing financial interest.

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ABBREVIATIONS USED

AAC, aminoglycoside N-acetyltransferase; ar, aromatic (NMR); ax, axial (NMR); AG, aminoglycoside; ANT, aminoglycoside O-nucleotidyltransferase; APH, aminoglycoside O-phosphoryltransferase; CA, cationic amphiphiles; eq, equatorial (NMR); MRSA, methicillin resistant *S. aureus*; 2NB, 2-naphthylbutyl; 2NM, 2-naphthylmethylene; 2NP, 2-naphthylpropyl; np, naphthyl ring (NMR); Nn, nonyl; OM, outer membrane; PMBnoc, para-methoxybenzyloxycarbonyl; PI, propidium iodide

REFERENCES

(1) Davies, J.; Gorini, L.; Davis, B. D. Misreading of RNA codewords induced by aminoglycoside antibiotics. *Mol. Pharmacol.* **1965**, *1*, 93–106.

(2) Moazed, D.; Noller, H. F. Interaction of antibiotics with functional sites in 16S ribosomal RNA. *Nature* **1987**, *327*, 389–394.

(3) Purohit, P.; Stern, S. Interaction of a small RNA with antibiotic and RNA ligands of the 30S subunit. *Nature* **1994**, *370*, 659–662.

(4) Recht, M. I.; Fourmy, D.; Blanchard, S. C.; Dahlquist, K. D.; Puglisi, J. D. RNA sequence determinants for aminoglycoside binding to an A-site rRNA model oligonucleotide. *J. Mol. Biol.* **1996**, *262*, 421–436.

(5) Fourmy, D.; Recht, M. L.; Puglisi, J. D. Binding of neomycin-class aminoglycoside antibiotics to the A-site of 16 S rRNA. *J. Mol. Biol.* **1998**, 277, 347–362.

(6) François, B.; Russell, R. J. M.; Murray, J. B.; Aboul-ela, F.; Masquida, B.; Vicens, Q.; Westhof, E. Crystal structures of complexes between aminoglycosides and decoding site oligonucleotides: role of the number of rings and positive charges in the specific binding leading to miscoding. *Nucleic Acids Res.* **2005**, *33*, 5677–5690.

(7) Borovinskaya, M. A.; Pai, R. D.; Zhang, W.; Schuwirth, B. S.; Holton, J. M.; Hirokawa, G.; Kaji, H.; Kaji, A.; Cate, J. H. D. Structural basis for aminoglycoside inhibition of bacterial ribosome recycling. *Nat. Struct. Mol. Biol.* **2007**, *14*, 727–732.

(8) Wright, G. D.; Berghuis, A. M.; Mobashery, S. Aminoglycoside antibiotics: structures, functions, and resistance. *Adv. Exp. Med. Biol.* **1998**, 456, 27–69.

(9) Jana, S.; Deb, J. K. Molecular understanding of aminoglycoside action and resistance. *Appl. Microbiol. Biotechnol.* 2006, 70, 140–150.
(10) Fosso, Y.; Li, Y.; Garneau-Tsodikova, S. New trends in the use

of aminoglycosides. MedChemComm **2014**, 5, 1075–1091.

(11) Chandrika, N. T.; Garneau-Tsodikova, S. A review of patents (2011–2015) towards combating resistance to and toxicity of aminoglycosides. *MedChemComm* **2016**, *7*, 50–68.

(12) Garneau-Tsodikova, S.; Labby, K. J. Mechanisms of resistance to aminoglycoside antibiotics: overview and perspectives. *MedChem*-*Comm* **2016**, *7*, 11–27.

(13) Mingeot-Leclercq, M.-P.; Glupczynski, Y.; Tulkens, P. M. Aminoglycosides: activity and resistance. *Antimicrob. Agents Chemother.* **1999**, 43, 727–737.

(14) Kotra, L.; Haddad, J.; Mobashery, S. Aminoglycosides: perspectives on mechanisms of action and resistance and strategies to counter resistance. *Antimicrob. Agents Chemother.* **2000**, *44*, 3249–3256.

(15) Magnet, S.; Blanchard, J. S. Insights into aminoglycoside action and resistance. *Chem. Rev.* 2005, 105, 477–497.

(16) Ramirez, M. S.; Tolmasky, M. E. Aminoglycoside modifying enzymes. Drug Resist. Updates 2010, 13, 151–171.

(17) Galimand, M.; Sabtcheva, S.; Courvalin, P.; Lambert, T. Worldwide disseminated *armA* aminoglycoside resistance methylase gene is borne by composite transposon Tn1548. *Antimicrob. Agents Chemother.* **2005**, 49, 2949–2953.

(18) Doi, Y.; Arakawa, Y. 16S Ribosomal RNA methylation: emerging resistance mechanism against aminoglycosides. *Clin. Infect. Dis.* **2007**, *45*, 88–94.

(19) Schmitt, E.; Galimand, M.; Panvert, M.; Courvalin, P.; Mechulam, Y. Structural bases for 16 S rRNA methylation catalyzed by ArmA and RmtB methyltransferases. *J. Mol. Biol.* **2009**, *388*, 570–582.

(20) François, B.; Szychowski, J.; Adhikari, S. S.; Pachamuthu, K.; Swayze, E. E.; Griffey, R. H.; Migawa, M. T.; Westhof, E.; Hanessian, S. Antibacterial aminoglycosides with a modified mode of binding to the ribosomal-RNA decoding site. *Angew. Chem., Int. Ed.* **2004**, *43*, 6735–6738.

(21) Hanessian, S.; Szychowski, J.; Adhikari, S. S.; Vasquez, G.; Kandasamy, P.; Swayze, E. E.; Migawa, M. T.; Ranken, R.; Francois, B.; Wirmer-Bartoschek, J.; Kondo, J.; Westhof, E. Structure-based design, synthesis, and A-site rRNA cocrystal complexes of functionally novel aminoglycoside antibiotics: C2["] ether analogues of paromomycin. *J. Med. Chem.* **200**7, *50*, 2352–2369.

(22) Hanessian, S.; Pachamuthu, K.; Szychowski, J.; Giguère, A.; Swayze, E. E.; Migawa, M. T.; François, B.; Kondo, J.; Westhof, E. Structure-based design, synthesis and A-site rRNA co-crystal complexes of novel amphiphilic aminoglycoside antibiotics with new binding modes: a synergistic hydrophobic effect against resistant bacteria. *Bioorg. Med. Chem. Lett.* **2010**, *20*, 7097–7101.

(23) Szychowski, J.; Kondo, J.; Zahr, O.; Auclair, K.; Westhof, E.; Hanessian, S.; Keillor, J. W. Inhibition of aminoglycoside-deactivating enzymes APH(3')-IIIa and AAC(6')-Ii by amphiphilic paromomycin $O_{2^{n-2}}$ ether analogues. *ChemMedChem* **2011**, *6*, 1961–1966.

(24) For review see: Mingeot-Leclercq, M.-P.; Decout, J.-L. Bacterial lipid membranes as promising targets to fight antimicrobial resistance, molecular foundations and illustration through the renewal of aminoglycoside antibiotics and emergence of amphiphilic aminoglycosides. *MedChemComm* **2016**, *7*, 586–611.

(25) Baussanne, I.; Bussière, A.; Halder, S.; Ganem-Elbaz, C.; Ouberai, M.; Riou, M.; Paris, J.-M.; Ennifar, E.; Mingeot-Leclercq, M.-P.; Décout, J.-L. Synthesis and antimicrobial evaluation of amphiphilic neamine derivatives. J. Med. Chem. **2010**, 53, 119–127.

(26) Ouberai, M.; El Garch, F.; Bussière, A.; Riou, M.; Alsteens, D.; Lins, L.; Baussanne, I.; Dufrêne, Y. F.; Brasseur, R.; Décout, J.-L.; Mingeot-Leclercq, M.-P. The *Pseudomonas aeruginosa* membranes: A target for a new amphiphilic aminoglycoside derivative? *Biochim. Biophys. Acta, Biomembr.* **2011**, *1808*, 1716–1727.

(27) Zimmermann, L.; Bussière, A.; Ouberai, M.; Baussanne, I.; Jolivalt, C.; Mingeot-Leclercq, M.-P.; Décout, J.-L. Tuning the antibacterial activity of amphiphilic neamine derivatives and comparison to paromamine homologues. *J. Med. Chem.* **2013**, *56*, 7691–7705.

(28) Sautrey, G.; Zimmermann, L.; Delbar, A.; Deleu, M.; Souza Machado, L.; Jeannot, K.; Van Bambeke, F.; Buyck, J.; Decout, J.-L.; Mingeot-Leclercq, M.-P. New amphiphilic neamine derivatives active against resistant *Pseudomonas aeruginosa*: Interactions with lipopolysaccharides. *Antimicrob. Agents Chemother.* **2014**, *58*, 4420–4430.

(29) Sautrey, G.; El Khoury, M.; Giro dos Santos, A.; Zimmermann, L.; Deleu, M.; Lins, L.; Decout, J.-L.; Mingeot-Leclercq, M.-P. Negatively-charged lipids as potential target for new amphiphilic aminoglycoside antibiotics: a biophysical study. *J. Biol. Chem.* **2016**, 291, 13864–13874.

(30) For review see: Gorityala, B. K.; Guchhait, G.; Schweizer, F. Amphiphilic aminoglycoside antimicrobials in antibacterial discovery. In *Carbohydrates in Drug Design and Discovery (RSC Drug Discovery)*; Jimenez-Barbero, J., Canada, F. J., Martin-Santamaria, S., Eds.; Royal Society of Chemistry: Cambridge, 2015; pp 255–285.

(31) For review see: Herzog, I. M.; Fridman, M. Design and synthesis of membrane-targeting antibiotics: from peptides- to aminosugarbased antimicrobial cationic amphiphiles. *MedChemComm* **2014**, *5*, 1014–1026.

(32) Zhang, J.; Chiang, F.-I.; Wu, L.; Czyryca, P. G.; Li, D.; Chang, C.-W. T. Surprising alteration of antibacterial activity of 5"-modified neomycin against resistant bacteria. *J. Med. Chem.* **2008**, *51*, 7563–7573.

(33) Bera, S.; Zhanel, G. G.; Schweizer, F. Design, synthesis and antibacterial activities of neomycin-lipid conjugates: polycationic lipids with potent gram-positive activity. *J. Med. Chem.* **2008**, *51*, 6160–6164.

(34) Zhang, J.; Keller, K.; Takemoto, J. Y.; Bensaci, M.; Litke, A.; Czyryca, P. G.; Chang, C.-W. T. Synthesis and combinational antibacterial study of 5"-modified neomycin. *J. Antibiot.* **2009**, *62*, 539–544.

(35) Bera, S.; Zhanel, G. G.; Schweizer, F. Antibacterial activities of aminoglycoside antibiotics-derived cationic amphiphiles. Polyol-modified neomycin B-, kanamycin A-, amikacin- and neamine-based amphiphiles with potent broad-spectrum antibacterial activity. *J. Med. Chem.* **2010**, *53*, 3626–3631.

(36) Dhondikubeer, R.; Bera, S.; Zhanel, G. G.; Schweizer, F. Antibacterial activity of amphiphilic tobramycin. *J. Antibiot.* **2012**, *65*, 495–498.

(37) Herzog, I. M.; Green, K. D.; Berkov-Zrihen, Y.; Feldman, M.; Vidavski, R. R.; Eldar-Boock, A.; Satchi-Fainaro, R.; Eldar, A.; Garneau-Tsodikova, S.; Fridman, M. 6"-Thioether tobramycin analogues: towards selective targeting of bacterial membranes. *Angew. Chem., Int. Ed.* **2012**, *51*, 5652–5656.

(38) Udumula, V.; Ham, Y. W.; Fosso, M. Y.; Chan, K. Y.; Rai, R.; Zhang, J.; Li, J.; Chang, C.-W. T. Investigation of antibacterial mode of action for traditional and amphiphilic aminoglycosides. *Bioorg. Med. Chem. Lett.* **2013**, *23*, 1671–1675.

(39) Berkov-Zrihen, Y.; Herzog, I. M.; Benhamou, R. I.; Feldman, M.; Steinbuch, K. B.; Shaul, P.; Lerer, S.; Eldar, A.; Fridman, M. Tobramycin and nebramine as pseudo-oligosaccharide scaffolds for the development of antimicrobial cationic amphiphiles. *Chem. - Eur. J.* **2015**, *21*, 4340–4349.

(40) Benhamou, R. I.; Shaul, P.; Herzog, I. M.; Fridman, M. Di-Nmethylation of anti-gram-positive aminoglycoside-derived membrane disruptors improves antimicrobial potency and broadens spectrum to gram-negative bacteria. *Angew. Chem., Int. Ed.* **2015**, *54*, 13617–13621. (41) Guchhait, G.; Altieri, A.; Gorityala, B.; Yang, X.; Findlay, B.; Zhanel, G. G.; Mookherjee, N.; Schweizer, F. Amphiphilic tobramycins with immunomodulatory properties. *Angew. Chem., Int. Ed.* **2015**, *54*,

6278–6282. (42) Gorityala, B. K.; Guchhait, G.; Fernando, D. M.; Deo, S.; McKenna, S. A.; Zhanel, G. G.; Kumar, A.; Schweizer, F. Adjuvants based on hybrid antibiotics overcome resistance in *Pseudomonas aeruginosa* and enhance fluoroquinolone efficacy. *Angew. Chem., Int. Ed.* **2016**, *55*, 555–559.

(43) Shaul, P.; Benhamou, R. I.; Herzog, I. M.; Louzoun Zada, S.; Ebenstein, Y.; Fridman, M. Synthesis and evaluation of membrane permeabilizing properties of cationic amphiphiles derived from the disaccharide trehalose. *Org. Biomol. Chem.* **2016**, *14*, 3012–3015.

(44) Fosso, M. Y.; Shrestha, S. K.; Green, K. D.; Garneau-Tsodikova, S. Synthesis and bioactivities of kanamycin B-derived cationic amphiphiles. *J. Med. Chem.* **2015**, *58*, 9124–9132.

(45) Chang, C.-W. T.; Takemoto, J. Y. Antifungal amphiphilic aminoglycosides. *MedChemComm* **2014**, *5*, 1048–1057.

(46) Fosso, M.; AlFindee, M. N.; Zhang, Q.; Nziko, V. d. P. N.; Kawasaki, Y.; Shrestha, S. K.; Bearss, J.; Gregory, R.; Takemoto, J. Y.; Chang, C.-W. T. Structure-Activity relationships for antibacterial to antifungal conversion of kanamycin to amphiphilic analogues. *J. Org. Chem.* **2015**, *80*, 4398–4411.

(47) Hurdle, J. G.; O'Neill, A. J.; Chopra, I.; Lee, R. E. Targeting bacterial membrane function: an underexploited mechanism for treating persistent infections. *Nat. Rev. Microbiol.* **2011**, *9*, 62–75.

(48) Findlay, B.; Zhanel, G. G.; Schweizer, F. Cationic amphiphiles, a new generation of antimicrobials inspired by the natural antimicrobial peptide scaffold. *Antimicrob. Agents Chemother.* **2010**, *54*, 4049–4058.

(49) Domalaon, R.; Zhanel, G. G.; Schweizer, F. Short antimicrobial peptides and peptide scaffolds as promising antibacterial agents. *Curr. Top. Med. Chem.* **2016**, *16*, 1217–1230.

(50) Marr, A. K.; Gooderham, W. J.; Hancock, R. E. Antibacterial peptides for therapeutic use: Obstacles and realistic outlook. *Curr. Opin. Pharmacol.* **2006**, *6*, 468–472.

(51) Van Bambeke, F.; Mingeot-Leclercq, M.-P.; Struelens, M. J.; Tulkens, P. M. The bacterial envelope as a target for novel anti-MRSA antibiotics. *Trends Pharmacol. Sci.* **2008**, *29*, 124–134.

(52) Rotem, S.; Mor, A. Antimicrobial peptide mimics for improved therapeutic properties. *Biochim. Biophys. Acta, Biomembr.* **2008**, *1788*, 1582–1592.

(53) Lohner, K. New strategies for novel antibiotics: peptides targeting bacterial cell membranes. *Gen. Physiol. Biophys.* **2009**, *28*, 105–116.

(54) Chen, C.; Pan, F.; Zhang, S.; Hu, J.; Cao, M.; Wang, J.; Xu, H.; Zhao, X.; Lu, J. R. Antibacterial activities of short designer peptides: A link between propensity for nanostructuring and capacity for membrane destabilization. *Biomacromolecules* **2010**, *11*, 402–411.

(55) Giuliani, A.; Rinaldi, A. C. Beyond natural antimicrobial peptides: multimeric peptides and other peptidomimetic approaches. *Cell. Mol. Life Sci.* **2011**, *68*, 2255–2266.

(56) Steinbuch, K. B.; Fridman, M. Mechanisms of resistance to membrane-disrupting antibiotics in Gram-positive and Gram-negative bacteria. *MedChemComm* **2016**, *7*, 86–102.

(57) Gurjar, M. Colistin for lung infection: an update. *J. Intensive Care* **2015**, *3*, 3–12.

(58) Velkov, T.; Roberts, K. D.; Nation, R. L.; Wang, J.; Thompson, P. E.; Li, J. Teaching 'old' polymyxins new tricks: New-generation lipopeptides targeting Gram-negative 'superbugs'. *ACS Chem. Biol.* **2014**, *9*, 1172–1177.

(59) Li, J.; Nation, R. L.; Turnidge, J. D.; Milne, R. W.; Coulthard, K.; Rayner, C. R.; Paterson, D. L. Colistin: The re-emerging antibiotic for multidrug-resistant gram-negative bacterial infections. *Lancet Infect. Dis.* **2006**, *6*, 589–601.

(60) Molina, J.; Cordero, E.; Pachon, J. New information about the polymyxin/colistin class of antibiotics. *Expert Opin. Pharmacother.* **2009**, *10*, 2811–2828.

(61) Michalopoulos, A. S.; Karatza, D. C. Multidrug-resistant Gram negative infections: The use of colistin. *Expert Rev. Anti-Infect. Ther.* **2010**, *8*, 1009–1017.

(62) Velkov, T.; Roberts, K. D.; Nation, R. L.; Thompson, P. E.; Li, J. Pharmacology of polymyxins: new insights into an old' class of antibiotics. *Future Microbiol.* **2013**, *8*, 711–724.

(63) Velkov, T.; Thompson, P. E.; Nation, R. L.; Li, J. Structureactivity relationships of polymyxin antibiotics. *J. Med. Chem.* **2010**, *53*, 1898–1916.

(64) Ma, Z.; Wei, D.; Yan, P.; Zhu, X.; Shan, A.; Bi, Z. Characterization of cell selectivity, physiological stability and endotoxin neutralization capabilities of alpha-helix-based peptide amphiphiles. *Biomaterials* **2015**, *52*, 517–530.

(65) Hevener, K. E.; Cao, S.; Zhu, T.; Su, P.-C.; Mehboob, S.; Johnson, M. E. Chapter Eighteen – Special challenges to the rational design of antibacterial agents. *Annu. Rep. Med. Chem.* **2013**, *48*, 283–298.

(66) Yin, L. M.; Edwards, M. A.; Li, J.; Yip, C. M.; Deber, C. M. Roles of hydrophobicity and charge distribution of cationic antimicrobial peptides in peptide-membrane interactions. *J. Biol. Chem.* **2012**, *287*, 7738–7745.

(67) Su, Y.; Waring, A. J.; Ruchala, P.; Hong, M. Structures of β hairpin antimicrobial protegrin peptides in lipopolysaccharide membranes: mechanism of Gram selectivity obtained from solidstate NMR. *Biochemistry* **2011**, *50*, 2072–2083.

(68) Epand, R. F.; Savage, P. B.; Epand, R. M. Bacterial lipid composition and the antimicrobial efficacy of cationic steroid compounds (Ceragenins). *Biochim. Biophys. Acta, Biomembr.* 2007, 1768, 2500–2509.

(69) Glukhov, E.; Stark, M.; Burrows, L. L.; Deber, C. M. Basis for selectivity of cationic antimicrobial peptides for bacterial versus mammalian membranes. *J. Biol. Chem.* **2005**, *280*, 33960–33967.

(70) Renner, L. D.; Weibel, D. B. Cardiolipin microdomains localize to negatively curved regions of *Escherichia coli* membranes. *Proc. Natl. Acad. Sci. U. S. A.* **2011**, *108*, 6264–6269.

(71) Renner, L. D.; Weibel, D. B. MinD and MinE interact with anionic phospholipids and regulate division plane formation in *Escherichia coli. J. Biol. Chem.* **2012**, *287*, 38835–38844.

(72) Svenson, J.; Brandsdal, B. O.; Stensen, W.; Svendsen, J. S. Albumin binding of short cationic antimicrobial micropeptides and its influence on the *in vitro* bactericidal effect. *J. Med. Chem.* **2007**, *50*, 3334–3339.

(73) Perigolo de Oliveira, M.; Constant, J.-F.; Peuchmaur, M.; Pitta, I.; Décout, J.-L. Antibiotic drugs aminoglycosides cleave DNA at abasic sites: Shedding new light on their toxicity? *Chem. Res. Toxicol.* **2013**, 26, 1710–1719.

(74) Denamur, S.; Tyteca, D.; Marchand-Brynaert, J.; Van Bambeke, F.; Tulkens, P. M.; Courtoy, P. J.; Mingeot-Leclercq, M.-P. Role of oxidative stress in lysosomal membrane permeabilization and apoptosis induced by gentamicin, an aminoglycoside antibiotic. *Free Radical Biol. Med.* **2011**, *51*, 1656–1665.

(75) Tabuchi, K.; Nishimura, B.; Nakamagoe, M.; Hayashi, K.; Nakayama, M.; Hara, A. Ototoxicity: mechanisms of cochlear impairment and its prevention. *Curr. Med. Chem.* **2011**, *18*, 4866–4871.

(76) Mingeot-Leclercq, M.-P.; Tulkens, P. Aminoglycosides: nephrotoxicity. *Antimicrob. Agents Chemother.* **1999**, *43*, 1003–1012.

(77) Kotra, L. P.; Mobashery, S. A renaissance of interest in aminoglycoside antibiotics. *Curr. Org. Chem.* **2001**, *5*, 193–205.

(78) Haddad, J.; Kotra, L. P.; Llano-Sotelo, B.; Kim, C.; Azucena, E. F.; Liu, M.; Vakulenko, S. B.; Chow, C. S.; Mobashery, S. Design of novel antibiotics that bind to the ribosomal acyltransfer site. *J. Am. Chem. Soc.* **2002**, *124*, 3229–3237.

(79) Agnelli, F.; Sucheck, S. J.; Marby, K. A.; Rabuka, D.; Yao, S.-L.; Sears, P. S.; Liang, F.-S; Wong, C.-H. Dimeric aminoglycosides as antibiotics. *Angew. Chem., Int. Ed.* **2004**, *43*, 1562–1566.

(80) Hermann, T. Aminoglycoside antibiotics: old drugs and new therapeutic approaches. *Cell. Mol. Life Sci.* 2007, 64, 1841–1852.

(81) Silva, J. G.; Carvalho, I. New insights into aminoglycoside antibiotics and derivatives. *Curr. Med. Chem.* **2007**, *14*, 1101–1119.

(82) Thomas, J. R.; Hergenrother, P. J. Targeting RNA with small molecules. *Chem. Rev.* **2008**, *108*, 1171–1224.

(83) Zhou, J.; Wang, G.; Zhang, L.-H.; Ye, X.-S. Modifications of aminoglycoside antibiotics targeting RNA. *Med. Res. Rev.* 2007, 27, 279–316.

(84) Dozzo, P.; Moser, H. E. New aminoglycoside antibiotics. *Expert Opin. Ther. Pat.* **2010**, *20*, 1321–1341.

(85) Houghton, J. L.; Green, K. D.; Chen, W.; Garneau-Tsodikova, S. The future of aminoglycosides: The end or renaissance. *ChemBioChem* **2010**, *11*, 880–902.

(86) Guo, L. N.; Wan, Y.; Wang, X.; Wang, P. G.; Zhao, W. Development of aminoglycoside antibiotics by carbohydrate chemistry. *Mini-Rev. Med. Chem.* **2012**, *12*, 1533–1541.

(87) Hevener, K. E.; Cao, S. Y.; Zhu, T.; Su, P. C.; Mehboob, S.; Johnson, M. E. Special challenges to the rational design of antibacterial agents. *Annu. Rep. Med. Chem.* **2013**, *48*, 283–298.

(88) Labby, K. J.; Garneau-Tsodikova, S. Strategies to overcome the action of aminoglycoside-modifying enzymes for treating resistant bacterial infections. *Future Med. Chem.* **2013**, *5*, 1285–1309.

(89) Pinsetta, F. R.; Kawano, D. F.; de Carvalho, M. R.; de Oliveira, J. A. A.; Corrado, A. P.; Hyppolito, M. A.; Carvalho, I. Synthesis of neamine-based pseudodisaccharides as potential vestibulotoxic agents to treat vertigo in Ménière's disease. *Carbohydr. Res.* **2013**, *373*, 97–102.

(90) Riguet, E.; Désiré, J.; Bailly, C.; Décout, J.-L. A route for preparing new neamine derivatives targeting HIV-1 TAR RNA. *Tetrahedron* **2004**, *60*, 8053–8064.

(91) Riguet, E.; Tripathi, S.; Chaubey, B.; Désiré, J.; Pandey, V. N.; Décout, J.-L. A Peptide nucleic acid-neamine conjugate that targets and cleaves HIV-1 TAR RNA inhibits viral replication. *J. Med. Chem.* **2004**, *47*, 4806–4809.

(92) Riguet, E.; Désiré, J.; Boden, O.; Ludwig, V.; Göbel, M.; Bailly, C.; Décout, J.-L. Neamine dimers targeting the HIV-1 TAR RNA. *Bioorg. Med. Chem. Lett.* **2005**, *15*, 4651–4655.

(93) Chaubey, B.; Tripathi, S.; Désiré, J.; Baussanne, I.; Décout, J.-L.; Pandey, V. N. Mechanism of RNA cleavage catalyzed by sequence specific polyamide nucleic acid-neamine conjugate. *Oligonucleotides* **2007**, *17*, 302–313.

(94) Ennifar, E.; Paillart, J.-C.; Bernacchi, S.; Walter, P.; Pale, P.; Décout, J.-L.; Marquet, R.; Dumas, P. A structure-based approach for targeting the HIV-1 genomic RNA dimerization initiation site. *Biochimie* **2007**, *89*, 1195–1203.

(95) Le Gall, T.; Baussanne, I.; Halder, S.; Carmoy, N.; Montier, T.; Lehn, P.; Décout, J.-L. Synthesis and transfection properties of a series of lipidic neamine derivatives. *Bioconjugate Chem.* **2009**, *20*, 2032–2046.

(96) Jackowski, O.; Bussière, A.; Vanhaverbeke, C.; Baussanne, I.; Peyrin, E.; Mingeot-Leclercq, M.-P.; Décout, J.-L. Major increases of the reactivity and selectivity in aminoglycoside O-alkylation due to the presence of fluoride ions. *Tetrahedron* **2012**, *68*, 737–746.

(97) Wong, C.-H.; Hendrix, M.; Manning, D. D.; Rosenbohm, C.; Greenberg, W. A. A library approach to the discovery of small molecules that recognize RNA: use of a 1,3-hydroxyamine motif as core. *J. Am. Chem. Soc.* **1998**, *120*, 8319–8327.

(98) Sucheck, S. J.; Wong, A. L.; Koeller, K. M.; Boehr, D. D.; Draker, K.; Sears, P. S.; Wright, G. D.; Wong, C.-H. Design of bifunctional antibiotics that target bacterial rRNA and inhibit resistance-causing enzymes. *J. Am. Chem. Soc.* **2000**, *122*, 5230–5231.

(99) Wu, B.; Yang, J.; Robinson, D.; Hofstadler, S.; Griffey, R.; Swayze, E. E.; He, Y. Synthesis of linked carbohydrates and evaluation of their binding for 16S RNA by mass spectrometry. *Bioorg. Med. Chem. Lett.* **2003**, *13*, 3915–3918.

(100) Vourloumis, D.; Winters, G. C.; Takahashi, M.; Simonsen, K. B.; Ayida, B. K.; Shandrick, S.; Zhao, Q.; Hermann, T. Novel acyclic deoxystreptamine mimetics targeting the ribosomal decoding site. *ChemBioChem* **2003**, *4*, 879–885.

(101) Das, I.; Désiré, J.; Manvar, D.; Baussanne, I.; Pandey, V. N.; Décout, J.-L. A peptide nucleic acid-aminosugar conjugate targeting transactivation response element of HIV-1 RNA genome shows a high bioavailability in human cells and strongly inhibits Tat-mediated transactivation of HIV-1 transcription. *J. Med. Chem.* **2012**, *55*, 6021– 6032.

(102) Henry, C.; Joly, J.-P.; Chapleur, Y. Efficient syntheses of methyl 2-amino-2-deoxy-3,4,6-tri-O-benzyl- α -D-glucopyranoside and its 2-tert-butoxycarbonylamino- and 2-methylamino derivatives from N-acetyl-D-glucosamine. J. Carbohydr. Chem. **1999**, *18*, 689–695.

(103) Peter, M. G.; Boldt, P.-C.; Petersen, S. Asymmetric induction in the epoxidation of alkenyl glycosides of tri-O-acetyl-N-acetyl- β -Dglucosamine. *Liebigs Ann. Chem.* **1992**, 1992, 1275–1279.

(104) Kasprzycka, A.; Szeja, W.; Grynkiewicz, G. Glycosylation of acid sensitive acceptors. Synthesis of (2,3-epoxy-1-propyl) glycosides. *Carbohydr. Res.* **2005**, 340, 2443–2446.

(105) Viswanadhan, V. N.; Ghose, A. K.; Revankar, G. R.; Robins, R. K. Atomic physicochemical parameters for three dimensional structure directed quantitative structure-activity relationships. 4. Additional parameters for hydrophobic and dispersive interactions and their application for an automated superposition of certain naturally occurring nucleoside antibiotics. *J. Chem. Inf. Model.* **1989**, *29*, 163–172.

(106) Niven, G. W.; Mulholland, F. Cell membrane integrity and lysis in Lactococcus lactis: the detection of a population of permeable cells in post-logarithmic phase cultures. *J. Appl. Microbiol.* **1998**, *84*, 90–96.

(107) Di Pasquale, E.; Salmi-Smail, C.; Brunel, J. M.; Sanchez, P.; Fantini, J.; Maresca, M. Biophysical studies of the interaction of squalamine and other cationic amphiphilic molecules with bacterial and eukaryotic membranes: importance of the distribution coefficient in membrane selectivity. *Chem. Phys. Lipids* **2010**, *163*, 131–140.

(108) Epand, R. F.; Pollard, J. E.; Wright, J. O.; Savage, P. B.; Epand, R. M. Depolarization, bacterial membrane composition, and the antimicrobial action of ceragenins. *Antimicrob. Agents Chemother.* **2010**, *54*, 3708–3713.

(109) Avrain, L.; Garvey, M.; Mesaros, N.; Glupczynski, Y.; Mingeot-Leclercq, M.-P.; Piddock, L. J.; Tulkens, P. M.; Vanhoof, R.; Van Bambeke, F. Selection of quinolone resistance in *Streptococcus pneumoniae* exposed in vitro to subinhibitory drug concentrations. J. *Antimicrob. Chemother.* **2007**, *60*, 965–972.

(110) Performance Standards for Antimicrobial Susceptibility Testing: 225 Informational Supplement; Document M100-S25; Clinical and Laboratory Standards Institute: Wayne, PA, 2015; http://clsi.org/blog/2015/01/08/clsi-publishes-new-antimicrobial-susceptibility-testing-standards/.

(111) Berridge, M. V.; Herst, P. M.; Tan, A. S. Tetrazolium dyes as tools in cell biology: new insights into their cellular reduction. *Biotechnol. Annu. Rev.* **2005**, *11*, 127–152.

(112) Berridge, M. V.; Tan, A. S. Characterization of the cellular reduction of 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bro-

mide (MTT): subcellular localization, substrate dependence, and involvement of mitochondrial electron transport in MTT reduction. *Arch. Biochem. Biophys.* **1993**, *303*, 474–482.

Supporting information

New Broad-spectrum Antibacterial Amphiphilic Aminoglycosides Active Against Resistant Bacteria: From Neamine Derivatives to

Smaller Neosamine Analogues

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Purity determination of the evaluated derivatives prepared

The purity was determined by HPLC for the compounds carrying chromophores (also by ¹H NMR spectrometry and TLC) and by ¹H NMR spectrometry and TLC for the other derivatives (compounds **12**, **37**, **39**, **40**, **43** described in the manuscript and compounds **27**, **38** previously described).

The AAGs carrying chromophores were analyzed by HPLC on an Agilent 1100 series HPLC machine using C18 reversed phase column (Macherey-Nagel, Nucleodur C18 ISIS, 5 μ m particle size, 100 Å pore size, 250 x 4.6 mm) and a diode array detector eluting with mixtures of different solutions:

X: Aqueous solution of trifluroroacetic acid (pH 2.5 at 20°C)

- Y: Aqueous solution of ammonium acetate 10 mM (pH 2.9 at 20 °C)
- Z: Acetonitrile

<u>Method A:</u> Gradient 100% X to 95:5 Z:X over 30 min, 1 mL/min, detection at 281 nm. <u>Method B:</u> Gradient 100% X to 95:5 Z:Y over 30 min, 1 mL/min, detection at 281 nm. Method C: Isocratic 95:5 Z:Y over 30 min, 1 mL/min, detection at 281 nm.

The purity was not determined by elemental analysis since (i) the main difficulty in the purification was the removal of isomers (for example, the 3',6-isomers of the 3',4'-neamine derivatives), (ii) the final compounds are very hygroscopic and (iii) it was not possible to detect by NMR at 400 MHz the presence of two diastereoisomers in compounds **39a,b**, **44a,b**-**47a,b** (expected diastereoisomeric ratio 2:1 from the corresponding epoxides mixture **31a,b** detected in the NMR spectrum).

TLC and ¹H NMR spectrometry allowed to detect the presence of isomers, for instance, the retardation factors of the 3',6-dinonyl derivative 7 and its 3',4'-isomer **11** were 0.5 and 0.3, respectively (eluent: EtOH/H₂O/(NH₃, H₂O) (20%) 80:10:10; TLC visualizations: sulfuric acid spray (5 mL in 100 mL EtOH) and ninhydrin spray (0.3 g, 3 mL AcOH, 100 mL EtOH).

Therefore, the absence or the presence of aminoglycoside isomers and other aminosugars as minor impurities was checked by ¹H NMR spectrometry (integration of the peaks corresponding to anomeric protons) and by TLC on silica gel.

Compounds	Purity (%)	Method
4579	> 05	UDL (NMD [®] (mercianaly, described)
4, 5, 7, 8	> 95	HPLC, NNR [®] (previously described)
10	98	HPLC
11	> 98	NMR ^a
17	96	HPLC, elemental analysis
28	96	HPLC
29	> 95	NMR ^a
34	> 95	NMR ^{a,b}
37	> 95	NMR ^{a,b}
38	> 95	NMR ^{a,b}
39a,b	> 95	NMR ^{a,b}
44a,b	> 95	NMR ^{a,b}
45a,b	> 95	NMR ^{a,b}
46a,b	> 95	NMR ^{a,b}
47a,b	> 95	NMR ^{a,b}

a: determined by ¹H NMR integration of the observed peaks corresponding to anomeric protons b: one diastereoisomer

c: 2:1 mixture of two diastereoisomers, the presence of two diastereoisomers cannot be detected by ¹H NMR at 400 MHz.

Compound **11** (¹H NMR):



Compound **29** (¹H NMR):







Allyl 6-azido-2,6-dideoxy-2-tritylamino-α-D-glucopyranoside (15) ¹H NMR CDCl₃



Allyl 6-azido-2,6-dideoxy-3,4-di-*O*-[(2-naphtyl)methyl]-2-tritylamino-α-D-glucopyranoside (16), ¹H NMR CDCl₃



Allyl 2,6-diamino-2,6-dideoxy-3,4-di-*O*-[(2-naphtyl)methyl]-α-D-glucopyranoside (17) ¹H NMR CDCl₃



Allyl 6-amino-2,6-dideoxy-2-tritylamino-α-D-glucopyranoside (18) ¹H NMR CDCl₃



Allyl 2,6-diamino-2,6-dideoxy-α-D-glucopyranoside (19) ¹H NMR CDCl₃





Compound 21





¹H NMR CDCl₃







Compound 26



Compound 27



Compound 28



Compound 29



¹H NMR CDCl₃



Compound **31a,b**





¹H NMR CDCl₃













¹H NMR CDCl₃



Compound 37

¹H NMR CD₃OD



¹H NMR CD₃OD



Compound **39a,b**



Compound 40a,b



Compound 41a,b

¹H NMR CDCl₃



Compound 42a,b

¹H NMR CDCl₃



Compound 43a,b



Compound 44a,b





Compound 45a,b



Compound 46a,b

¹H NMR CD₃OD



Compound 47a,b

¹H NMR CD₃OD

