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Development of Fragment-Based Inhibitors of the Bacterial Deacetylase LpxC with Low Nanomolar Activity

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INTRODUCTION

isoserine-based amides.

Due to the emergence of pathogenic bacteria being resistant to most or all of the currently available antibiotics and the low success rate of drug development in this field, the need for new molecular frameworks is particularly crucial for antibiotics.^{1–6} Following the golden age of antibiotic research (1940–1960), the development of new antibiotic scaffolds for Gram-negative pathogens dramatically decreased.^{1,7} The target-based high-throughput screening campaigns carried out against a number of bacterial enzymes all failed to deliver candidates, and very low hit rates were obtained in comparison to what was typically observed with nonbacterial targets. Whole-cell high-throughput screening also yielded low hit rates.^{1,8} Alternative approaches and/or novel therapeutic targets are therefore necessary for the generation of novel antibiotics.

The inhibition of the biosynthesis of lipid A is a promising but hitherto clinically unexploited strategy for the development of antibiotics selectively combating Gram-negative bacteria.⁹ Lipid A acts as the hydrophobic membrane anchor of the lipopolysaccharides (LPS), which represent the main component of the outer monolayer of the outer membrane of Gramnegative bacteria, thus being essential for growth and viability of nearly all Gram-negative bacteria.¹⁰

In Gram-negative bacteria, the biosynthesis of Kdo_2 -lipid A comprises a conserved pathway including nine enzymes.¹⁰ Its second step, in *Escherichia coli* the irreversible deacetylation of uridine diphosphate (UDP)-3-O-[(R)-3-hydroxymyristoyl]-N-

acetylglucosamine (1, Figure 1A), is considered as the committed step of lipid A biosynthesis.¹¹ This step is catalyzed by the Zn^{2+} -dependent deacetylase LpxC. The enzyme is present in virtually all Gram-negative bacteria, is highly conserved among them, and possesses no mammalian counterpart, which makes LpxC an excellent target for the development of novel antibiotics.¹²

Structural studies revealed that LpxC displays a " $\beta - \alpha - \alpha - \beta$ sandwich" fold, being formed by two domains with similar topologies.¹⁴ The catalytic Zn²⁺-ion is complexed by one aspartate and two histidine residues at the bottom of a conical active-site cleft, which is located at one side of the sandwich at the interface of the two domains (Figure 1B).¹⁵ A hydrophobic tunnel leads out of the active-site pocket, which binds the 3-*O*-[(*R*)-3-hydroxyacyl] substituent of the enzyme's natural substrate **1** during catalysis.¹⁶

Hitherto, various structural classes of small-molecule LpxC inhibitors have been described.^{22–27} Like the *N*-aroyl-L-threonine derivatives CHIR-090 (Figure S1) and LPC-011 (**3a**, Figure 2),^{16,28} most of these inhibitors exhibit a Zn^{2+} -

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Figure 1. (A) LpxC-catalyzed deacetylation of UDP-3- $O_{-}[(R)$ -3-hydroxymyristoyl]-N-acetylglucosamine (1). (B) Molecular surface (colored according to the hydrophobicity, polar regions are colored magenta, hydrophobic regions are colored green) of *E. coli* LpxC near the deacetylated natural product 2 (PDB ID: 4MDT).¹³ UDP-3- $O_{-}[(R)$ -3-hydroxymyristoyl]-glucosamine (2) is shown as a cyan colored ball and stick model.

Figure 2. Structures of the described LpxC inhibitors targeting the enzyme's UDP-binding site. Reported inhibitory activities toward *E. coli* LpxC (*Ec*LpxC) and *P. aeruginosa* LpxC (*Pa*LpxC) are given.^{17–21}

chelating hydroxamate moiety, which is linked to a lipophilic side chain addressing the enzyme's hydrophobic tunnel. Although extensive structure–activity relationship (SAR) studies around the lipophilic side chain as well as the linker region have been performed, only a few LpxC inhibitors addressing the binding site of the UDP moiety of the enzyme's natural substrate 1 have been reported so far.^{22,23} The uridinebased compound 1–68A (4, Figure 2), which weakly inhibits LpxC, is a rare example of an inhibitor targeting the UDPbinding site of LpxC.¹⁷ Additionally, some attempts were undertaken to expand known inhibitors into the UDP-binding site. For instance, the attachment of a hydroxyphenyl group to the β -carbon atom of the threonyl moiety of LPC-011 (3a) led to a 1.6-fold improvement in inhibitory activity toward LpxC.¹⁸ In case of ether **5a** and amide **6a**, the attachment of suitable substituents also led to an increase in binding affinity over the respective parent compound.^{19,20} Finally, analogues of biphenyl derivative **7a** bearing substituents on the β -amino group retained potent inhibitory activity against LpxC, although they exhibited weaker antimicrobial activity.²¹ Altogether, these examples clearly show the feasibility of such attempts. However, although the fragment-based discovery of LpxC inhibitors containing a nonhydroxamate Zn²⁺-chelating motif has been reported recently,²⁹ no systematic fragment-based studies aiming to find suitable substituents addressing the UDP-binding site have been described in the literature so far.

Figure 3. Structures of the described benzyloxyacetohydroxamic acid-based LpxC inhibitors and the envisaged linked ligands (encircled).^{30–32}

Table 1.	Antibacterial	and LpxC	Inhibitory	Activities	of Reported	and Newly	Synthesized	ł Hydroxamic	Acids

$\langle \rangle$		Ki	[µM]	MIC [µg/mL]		Inhibition zone [mm]	
R = -	compound	<i>E. coli</i> LpxC C63A	<i>P. aeruginosa</i> LpxC	<i>E. coli</i> BL21(DE3)	E. coli D22	<i>E. coli</i> BL21(DE3)	E. coli D22
	(S) -8 ³¹	0.41 ± 0.074	n.d.	32	1.0	15.7 ± 0.6	25.8 ± 1.9
HO	(R) -8 ³¹	0.24 ± 0.046	n.d.	64	2.00	12.3 ± 0.6	22.0 ± 1.3
HO. N. C. C. R.	9 ³¹	1.5 ± 0.37	n.d.	64	1.0	16.5 ± 0.4	20.3 ± 1.0
	(2 <i>S</i> ,3 <i>S</i>)- 10 ³²	0.28 ± 0.21	n.d.	4.0	0.13	11.2 ± 0.8	19.5 ± 1.5
HOTO	(2R,3S) -10 ³²	0.87 ± 0.29	n.d.	>64	4.0	8.5 ± 0.6	13.3 ± 1.7
HO. N L O	(S) -11a	0.606 ± 0.356	0.725 ± 0.335	>32	0.50	≤6	15.4 ± 2.4
	(<i>R</i>)-11a	3.62 ± 0.64	n.d.	>32	2.0	≤6	15.7 ± 1.2
	(S) -12a	4.59 ± 1.26	0.862 ± 0.070	16	0.50	13.0 ± 1.0	24.3 ± 2.1
HN T	(<i>R</i>)-12a	5.90 ± 2.02	n.d.	32	4.0	10.7 ± 1.2	20.0 ± 1.8
HO.N.O.	(S) -37	>2.54	>3.25	>16	16	≤6	7.7 ± 1.2
	(R) -37	>2.54	n.d.	>16	>16	≤6	≤6
HO. N CONTR	(S) -13a	0.149 ± 0.056	0.0582 ± 0.0001	8.0	0.13	15.7 ± 1.6	26.3 ± 1.5
HN CONTRACT	(<i>R</i>)-13a	2.56 ± 1.08	n.d.	>64	4.0	9.7 ± 0.6	18.0 ± 2.0

^{*a*}n.d.: not determined.

Previously, we reported on the design, synthesis, and biological evaluation of a series of benzyloxyacetohydroxamic acid derivatives like glyceric acid derivatives (S)-8 and (R)-8 as well as the α -position-unsubstituted benzyloxyacetic acid derivative 9 (Figure 3, Table 1).^{30,31} While glyceric acid derivatives (S)-8 and (R)-8 were found to exhibit promising LpxC inhibitory and antibacterial activities, removal of the hydroxymethyl group in α -position, leading to compound 9, caused a decrease in inhibitory activity. As these compounds leave the binding pocket for the UDP moiety of the natural substrate of LpxC unoccupied (Figure 4), substituents addressing the UDP-binding pocket should be introduced in the α -position of the hydroxamate moiety in order to gain further favorable interactions with the enzyme, thus improving the affinity of the compounds toward LpxC. To demonstrate

the feasibility of this approach, we previously synthesized aldotetronic acid derivatives $10^{.32}$ Even though the LpxC inhibitory activity of the most potent stereoisomer (2S,3S)-10 only slightly exceeded the one of glyceric acid derivative (S)-8 (Table 1), the observation that the elongation of the substituent in α -position did not diminish LpxC inhibitory activity shows that it is possible to grow our inhibitors into the UDP-binding site of LpxC.

Having shown that the introduction of a substituent addressing the UDP-binding pocket of LpxC in the α -position of the hydroxamate moiety is feasible, in this paper we report how we pursued this attractive strategy to further optimize our benzyloxyacetohydroxamic acid derivatives. In order to generate LpxC inhibitors that address the UDP-binding pocket, various linkers were investigated. Thus, besides the

Figure 4. Docking pose of inhibitor 9 (colored yellow) at the LpxC UDP-binding pocket superimposed with the reaction product UDP-3-O-[(R)-3-hydroxymyristoyl]-glucosamine (2, colored green) crystallized with LpxC (PDB ID: 4MDT).¹³ The molecular surface is displayed and colored according to the polarity. Polar regions are colored magenta, and hydrophobic regions are colored green. The zinc ion is shown as a cyan sphere. The black arrow marks the site where the inhibitors of the current study were substituted to target the UDP-binding pocket.

tetrose-derived ethers 10, glyceric acid-derived ethers 11 as well as isoserine-based amines 12 and amides 13 were synthesized. In parallel, to identify chemical structures capable of binding into the available UDP pocket, we have performed a fragment screen against LpxC in the presence of compound 9. We screened a library of 650 fragments to identify small substructures that bind the protein target through a minimal recognition motif. Fragments, small compounds (MW < 250 Da) with low complexity, typically bind proteins with weak affinities (usually $K_D > 100 \ \mu M$), nevertheless exhibiting high ligand efficiency (binding energy per heavy atom) due to highquality interactions.^{33,34} Saturation transfer difference (STD)-NMR, WaterLogsy, and subsequent NMR-interligand nuclear Overhauser effect (ILOE) experiments³⁵⁻³⁷ were performed to identify fragments that bind into the enzyme's UDP-binding pocket near the methylene group in the α -position of the

Scheme 1. Synthesis of Benzyl Ether (S)-11a^a

hydroxamate moiety of compound 9. After the identification of an optimal linker, substituents inspired by the identified fragments were connected with the benzyloxyacetohydroxamic acids, and the biological activities of the obtained compounds were investigated. Additionally, the effect of the stereochemistry of the newly synthesized compounds on their biological activities was studied.

RESULTS AND DISCUSSION

Synthesis of Compounds with Ether, Amine, and Amide Linkers. In order to find an optimal linker, a phenyl ring as an exemplary substituent should be connected to the scaffold of the benzyloxyacetohydroxamic acid derivatives via structural elements of different structures.

Thus, to vary the length and structure of the linker region compared to the previously investigated aldotetronic acidbased LpxC inhibitors, benzyl ethers (S)-11a and (R)-11a were synthesized. Ether (S)-11a was obtained from (R)glycidol (14, Scheme 1). After the protection of the primary alcohol of 14 with tert-butyldiphenylsilyl chloride, the oxirane ring of the resulting silvl ether 15 was opened with benzyl alcohol in the presence of erbium(III) triflate, yielding secondary alcohol 16.38 Then, a Williamson ether synthesis with 4-iodobenzyl bromide was performed, and the silyl protective group was cleaved with tetrabutylammonium fluoride to give primary alcohol 17. Subsequently, the alcohol was oxidized using an oxidant solution, which was composed of periodic acid and catalytic amounts of chromium(VI) oxide in wet acetonitrile, and the resulting carboxylic acid was esterified with methanol in the presence of sulfuric acid. The thereby obtained ester (S)-18 was subjected to a Sonogashira coupling with 4-(4-ethynylbenzyl)morpholine to yield diphenylacetylene derivative (S)-19. Finally, ester (S)-19 was converted into the desired hydroxamic acid (S)-11a by performing an aminolysis with hydroxylamine.^{39,40}

The enantiomeric ether (R)-11a was accessed in a chiral pool synthesis starting from D-mannitol (**20**, Scheme 2). After the conversion of D-mannitol (**20**) into trisacetonide **21** with acetone in the presence of sulfuric acid, **21** was stirred at 40 °C in 70% aqueous acetic acid to yield 3,4-O-isopropylidene-D-

"Reagents and Conditions: (a) TBDPSCl, imidazole, CH_2Cl_2 , 0 °C \rightarrow rt, 86%; (b) BnOH, $Er(OTf)_3$, rt, 73%; (c) (1) NaH, 4-iodobenzyl bromide, THF, 0 °C \rightarrow rt, (2) TBAF, THF, rt, 71%; (d) (1) CrO_3/H_5IO_6 , ACN, H_2O , 0 °C \rightarrow rt, (2) H_2SO_4 , MeOH, Δ , 48%; (e) 4-(4-ethynylbenzyl)morpholine, [PdCl₂(PPh₃)₂], CuI, diisopropylamine, THF, rt, 88%; (f) aq. NH₂OH, THF/*i*-PrOH (1:1), 0 °C \rightarrow rt, 67%.

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Scheme 2. Synthesis of Benzyl Ether (R)-11a^a

^{*a*}Reagents and Conditions: (a) acetone, H₂SO₄, rt, 32%; (b) 70% HOAc, 40 °C, 69%; (c) (1) Bu₂SnO, toluene, Δ , (2) benzyl bromide, Bu₄NI, toluene, 70 °C, 92%; (d) NaH, 4-iodobenzyl bromide, THF, rt, 85%; (e) 80% TFA, 0 °C, 97%; (f) (1) NaIO₄, MeOH, rt, (2) Br₂, NaHCO₃, MeOH/H₂O (9:1), rt, 69%; (g) 4-(4-ethynylbenzyl)morpholine, [PdCl₂(PPh₃)₂], CuI, diisopropylamine, THF, Δ , then rt, 64%; (h) aq. NH₂OH, THF/*i*-PrOH (1:1), 0 °C \rightarrow rt, 65%.

mannitol (22).^{41,42} Subsequently, the primary alcohols of tetrol 22 were selectively alkylated with benzyl bromide to give diether 23, which was subjected to a Williamson ether synthesis with 4-iodobenzyl bromide, yielding tetraether 24. Then, the remaining acetonide moiety was hydrolyzed, and the C-C bond between C-3 and C-4 of the resulting glycol 25 was cleaved with sodium periodate to yield two identical (*R*)-configured aldehydes, which were directly transformed into glyceric acid ester (*R*)-18 via an oxidation with bromine in a 9:1 mixture of methanol and water in the presence of sodium bicarbonate.⁴³ Like its enantiomer, hydroxamic acid (*R*)-11a was finally obtained from aryl iodide (*R*)-18 via a Sonogashira coupling and a subsequent aminolysis with hydroxylamine.

The envisaged isoserine-derived amines and amides were synthesized from azides (S)-28 and (R)-28. To obtain azide (S)-28, epoxide 15 was subjected to a ring-opening reaction with sodium azide in the presence of ammonium chloride to yield secondary alcohol 26 as the preferred regioisomer (Scheme 3).⁴⁴ A Williamson ether synthesis with 4-iodobenzyl bromide and the subsequent cleavage of the silyl group afforded primary alcohol 27, which was oxidized and thereafter reacted with methanol in the presence of sulfuric acid to give the desired ester (S)-28.

The enantiomeric azide (R)-28 could have been synthesized in the same way starting from commercially available (S)glycidol. However, an alternative synthetic route starting from D-mannitol (20) was established. Thus, the primary alcohols of D-mannitol-derived tetrol 22 were selectively tosylated via the intermediate formation of the respective stannylidene acetals (Scheme 3). The thereby obtained bistosylate 29 was subjected to a nucleophilic substitution with sodium azide to give diazide **30**. Subsequently, the remaining hydroxy groups were alkylated with 4-iodobenzyl bromide yielding diether **31**, followed by an acetal cleavage to obtain glycol **32**. As for the synthesis of the (R)-configured benzyl ethers, diol **32** was subjected to a glycol cleavage with sodium periodate and the obtained identical (R)-configured aldehydes were oxidized to ester (R)-**28** with bromine in a mixture of methanol and water (9:1) buffered with sodium bicarbonate.

While the chiral pool synthesis starting from D-mannitol required more steps and gave (R)-28 in a considerably lower overall yield compared to the synthesis of (S)-28 starting from (R)-glycidol [8.5% for (R)-28 vs 32% for (S)-28], this synthetic route yielded (R)-28 in an enantiomerically pure form [*ee* of (R)-28: 100% vs *ee* of (S)-28: 99.2%].

Starting from azide (S)-28, secondary amine (S)-33 could be obtained via a Staudinger/aza-Wittig reaction, followed by the reduction of the resultant imine intermediate (Scheme 4). Thus, azide (S)-28 was reacted with triethyl phosphite, and the resulting triethoxyiminophosphorane intermediate was converted into an imine with benzaldehyde, which was then reduced with sodium borohydride.45 In contrast, a Staudinger reduction of azide (S)-28 with polymer-bound triphenylphosphine in the presence of water and subsequent reductive alkylation of the resultant primary amine with an excess of benzaldehyde and sodium triacetoxyborohydride yielded tertiary amine (S)-35.⁴⁶ Finally, benzamide (S)-38a could be accessed via a Staudinger-Vilarrasa reaction by reacting azide (S)-28 with benzoic acid in the presence of trimethylphosphane and 2,2'-dithiodipyridine.47 Subsequently, the three isoserine-derived aryl iodides (S)-33, (S)-35, and (S)-38a were subjected to Sonogashira couplings with 4-(4-ethynylbenzyl)-

Scheme 3. Synthesis of Azides (S)-28 and (R)-28^a

^aReagents and Conditions: (a) NaN₃, NH₄Cl, MeOH, H₂O, 65 °C, 70%; (b) (1) NaH, 4-iodobenzyl bromide, THF, 0 °C \rightarrow rt, (2) TBAF, THF, rt, 78%; (c) (1) CrO₃/H₅IO₆, ACN, H₂O, 0 °C \rightarrow rt, (2) H₂SO₄, MeOH, Δ , 65%; (d) (1) Bu₂SnO, toluene, Δ , (2) *p*-TsCl, CHCl₃, 0 °C \rightarrow rt, 94%; (e) NaN₃, DMSO, 80 °C, 89%; (f) NaH, 4-iodobenzyl bromide, THF, rt, 82%; (g) 80% HOAc, 0 °C, 90%; (h) (1) NaIO₄, MeOH, rt, (2) Br₂, NaHCO₃, MeOH/H₂O (9:1), rt, 62%.

morpholine and final aminolyses with hydroxylamine to yield hydroxamic acids (S)-12a, (S)-37, and (S)-13a.

The respective enantiomers (R)-12a, (R)-37, and (R)-13a were obtained in principally the same way starting from azide (R)-28.

Identification of the Most Suitable Linker. In order to determine the inhibitory activities of the compounds of interest toward *E. coli* LpxC, a fluorescence-based enzyme assay was deployed, in which the formed deacetylated natural product UDP-3-O-[(R)-3-hydroxymyristoyl]glucosamine (2) is transformed into a fluorescent isoindole with o-phthaldialdehyde and 2-mercaptoethanol.^{48,49} In the assay, *E. coli* LpxC C63A was used, as this mutant is significantly less susceptible to being inhibited by high concentrations of Zn²⁺ compared to the wild-type enzyme.^{50,51} The $K_{\rm M}$ of *E. coli* LpxC C63A was determined experimentally using a mass spectrometry-based LpxC assay (Figure S8, Supporting Information, vide infra). The $K_{\rm M}$ was found to be 3.6 μ M, thus being in the same range as the reported $K_{\rm M}$ of wild-type *E. coli* LpxC (4.0 μ M).⁵²

In order to identify the optimal linker to connect a phenyl ring with the benzyloxyacetic acid scaffold, the LpxC inhibitory activities of the tetrose-derived ethers (2S,3S)-10 and (2R,3S)-10, glyceric acid-derived ethers (S)-11a and (R)-11a, isoserine-based secondary amines (S)-12a and (R)-12a, tertiary amines (S)-37 and (R)-37, and amides (S)-13a and (R)-13a were compared (Table 1). As reported previously, the inhibitory activity of tetrose-derived ether (2S,3S)-10 slightly exceeded the one of glyceric acid derivative (S)-8.³² And while the

configuration in α -position had only a minor effect on the inhibitory activities of glyceric acid-derived hydroxamic acids (S)-8 and (R)-8,³¹ with a slight superiority of the (R)configured enantiomer, the (2S)-configured aldotetronic acid derivative (2S,3S)-10 exhibited an about 3-fold higher inhibitory activity compared to its (2R)-configured diastereomer (2R,3S)-10.³² The newly synthesized glyceric acidderived ether (S)-11a was found to be a slightly less potent LpxC inhibitor than glyceric acid derivative (S)-8 and tetrosederived ether (2S,3S)-10. In case of ethers (S)-11a and (R)-**11a**, the inhibitory activity of the (S)-configured stereoisomer exceeds the one of its (R)-configured enantiomer by a factor of 6. While the isoserine-based amines (S)-12a, (R)-12a, (S)-37, and (R)-37 showed considerably reduced LpxC inhibitory activities, the isoserine-based amide (S)-13a exhibited a promising K_i value of 0.15 μ M, thus exceeding the inhibitory activities of glyceric acid derivative (S)-8 and tetrose-derived ether (2S,3S)-10. Additionally, in case of benzamides (S)-13a and (R)-13a, the eudysmic ratio was the highest in the examined series of compounds, with (S)-13a being a 17-fold more potent LpxC inhibitor than its enantiomer (R)-13a.

Therefore, the amide linker turned out to be the most favorable with respect to inhibitory activity toward LpxC.

Besides measuring the inhibitory activities toward *E. coli* LpxC C63A, the antibacterial activities of the synthesized hydroxamic acids were evaluated by determining their minimal inhibitory concentrations (MIC) in broth dilution tests as well as by performing disc diffusion assays (Table 1). Thus, the

Scheme 4. Synthesis of Hydroxamic Acids (S)-12a, (S)-37, and (S)-13a-n^a

^aReagents and Conditions: (a) (1) P(OEt)₃, toluene, 0 °C \rightarrow rt, (2) benzaldehyde, 0 °C \rightarrow rt, (3) NaBH₄, MeOH, 0 °C \rightarrow rt, 67%; (b) 4-(4ethynylbenzyl)morpholine, [PdCl₂(PPh₃)₂], CuI, diisopropylamine, THF, rt, 68%; (c) aq. NH₂OH, THF/*i*-PrOH (1:1), 0 °C \rightarrow rt, 70%; (d) (1) PS-PPh₃, THF, H₂O, 40 °C, (2) benzaldehyde, NaBH(OAc)₃, dichloroethane, 0 °C \rightarrow rt, (3) H₂SO₄, MeOH, 80 °C, 50%; (e) 4-(4ethynylbenzyl)morpholine, [PdCl₂(PPh₃)₂], CuI, diisopropylamine, THF, rt, 87%; (f) aq. NH₂OH, THF/*i*-PrOH (1:1), 0 °C \rightarrow rt, 47%; (g) P(CH₃)₃, R-CO₂H, 2,2'-dithiodipyridine, toluene, 0 °C \rightarrow rt, (S)-38a 83%, (S)-38b 74%, (S)-38c 76%, (S)-38d 77%, (S)-38e 92%, (S)-38f 82%, (S)-38g 76%, (S)-38h 65%, (S)-38i 64%, (S)-38j 34%, (S)-38k 33%, (S)-38l 50%, (S)-38m 66%, (S)-38m 83%; (h) 4-(4-ethynylbenzyl)morpholine, [PdCl₂(PPh₃)₂], CuI, diisopropylamine, THF, rt, (S)-39a 94%, (S)-39b 50%, (S)-39c 88%, (S)-39d 90%, (S)-39e 99%, (S)-39f 78%, (S)-39g 87%, (S)-39h 98%, (S)-39i 64%, (S)-39j 83%, (S)-39k 78%, (S)-39l 78%, (S)-39m 88%, (S)-39n 87%; (i) aq. NH₂OH, THF/*i*-PrOH (1:1), 0 °C \rightarrow rt, (S)-13a 41%, (S)-13b 82%, (S)-13c 70%, (S)-13d 77%, (S)-13e 68%, (S)-13f 79%, (S)-13g 82%, (S)-13h 67%, (S)-13i 63%, (S)-13j 86%, (S)-13k 75%, (S)-13l 48%, (S)-13m 76%, (S)-13n 84%.

compounds were tested against *E. coli* BL21(DE3) ($lpxC^+$) and the defective *E. coli* D22 (lpxC101) strain,⁵³ exhibiting reduced LpxC activity.

A comparison of the benzyloxyacetohydroxamic acids bearing a phenyl substituent connected via different linker regions revealed that the tetrose-derived ether (2S,3S)-10 as well as the isoserine-based amide (S)-13a exhibit the lowest MIC values against the two investigated *E. coli* strains, which agrees with the low K_i values of the two compounds. Surprisingly, although showing only negligible inhibitory activity against *E. coli* LpxC C63A, notable antibacterial activity was found for secondary amine (S)-12a.

Fragment Screening Using NMR. In order to find fragments specifically addressing the UDP-binding pocket, which can be linked to the LpxC inhibitors, a fragment screen was performed against LpxC bound to 9. First, 650 fragments in cocktails of five were tested for binding to the LpxC–9 complex using STD-NMR and WaterLogsy experiments. Then, fragment hits identified as potential binders were tested one by

one against the LpxC-9 complex by STD-NMR and WaterLogsy experiments performed under identical experimental conditions. A total of 97 fragment hits were identified as binders, leading to a hit rate of 15%.

High hit rates are typically observed in fragment screening due to the low complexity and small size of the fragments as well as artifacts. Also, the highly hydrophobic nature of the fragments (mostly aromatic) gives them a high ability to bind to protein pockets. Such fragments can exhibit multiple binding modes when binding to proteins, which give rise to STD factors that are averaged and can therefore have similar values for each proton of the fragment.⁵⁴ By contrast, fragments that bind proteins with a privileged binding mode should exhibit a STD-based epitope mapping,^{54,55} where the STD factor value for each proton reflects the proximity between the fragment proton to the protein protons. Here, due to the high rate of fragment hits, we decided to select fragments that displayed a STD-based epitope mapping. We acknowledge that this approach does not guarantee a unique binding mode for the fragments, and that some interesting fragments can be missed. We considered that this strategy would help in the selection of the most promising ligands. Finally, 19 fragment hits were selected as potentially interesting LpxC binders, which correspond to a hit rate of 2.9% (Table 2 and Figure 5A).

Then, NMR-ILOE experiments were performed with selected fragment hits in the presence of LpxC and the unsubstituted hydroxamic acid 9. This allowed the identification of fragments that bind in the enzyme's UDP-binding pocket near the methylene group in the α -position of the

Table 2. Fragment Hits Identified as Binders of the LpxC-9 Complex

Fragment code	Chemical structure of fragments identified as binders using STD and WaterLogsy experiments	Inter-Ligand NOESY peaks between the fragment and 9
F1	HJCO	yes
F2	но-	yes
F3	<u>(</u> کې	yes
F4		yes
F5	ĊĊţ	yes
F6		yes
F7		no
F8		no
F9	HO	no
F10	$(\mathbf{Q}_{\mathbf{g}}\mathbf{Q})$	no
F11	H,N	yes
F12		no
F13	С С На	no
F14		yes
F15	HO CON NH	no
F16		no
F17	H ₂ N C	no
F18	CL.	no
F19		yes

inhibitor's hydroxamate moiety. Weak interligand NOESY peaks were observed between inhibitor 9 and nine fragments. By contrast, no interligand NOESY peaks were observed between compound 9 and the other ten fragments. The nine fragments exhibited weak interligand NOESY peaks between protons of the fragment and protons of the CH_2 groups near the hydroxamate moiety as well as the aromatic resonances of 9 (see, for example, Figure 5B, showing the NOESY spectrum between 9 and F1). Unexpectedly, we also observed interligand NOESY between the fragment protons and protons of the morpholino moiety.

Using the data from the STD, WaterLogsy, and NOESY experiments, we identified nine fragments that were capable of binding LpxC in the UDP-binding pocket near the inhibitor's hydroxamate moiety (Table 2). The interpretation of the NOESY was rather ambiguous, and no further structural information (such as the relative orientation of the fragment and compound 9) was inferred from these experiments. Thus, from these chemical structures, we selected four substructures (biphenyl, indole, quinoline, and pyrazole) to be linked to the benzyloxyacetic acid scaffold of hydroxamic acid 9.

Synthesis and LpxC Inhibitory Activity of Linked Compounds. As the amide linker had turned out to be the most favorable with respect to inhibitory activity toward LpxC and the Staudinger–Vilarrasa reaction provided convenient access to the desired amides, the selected biphenyl-, indole-, quinoline-, and pyrazole-based substituents were linked via an amide linker to the scaffold of hydroxamic acid 9. Additionally, to broaden the SAR, diphenyl ether-, pyrrole-, and isoxazole-based substituents were introduced. Thus, azide (S)-28 was ligated with the respective carboxylic acids to finally obtain hydroxamic acids (S)-13b–n (Scheme 4). The enantiomeric amides (R)-13b–h,j,m,n were obtained in principally the same way starting from azide (R)-28.

The synthesized amide-based hydroxamic acids **13** were tested for inhibitory activity toward *E. coli* LpxC C63A. The results are reported in Table 3.

Generally, as observed for benzamides (*S*)-13a and (*R*)-13a, the (S)-configured compound was found to be the eutomer of all investigated pairs of enantiomeric amides. In case of the synthesized benzamide derivatives, the introduction of another phenyl ring in position 4 of the benzoyl moiety, leading to biphenyl derivative (S)-13b, was found to be detrimental to inhibitory activity. While the introduction of a 4-phenoxy substituent, leading to diphenyl ether (S)-13d, caused only a slight reduction of inhibitory activity compared to (S)-13a, the introduction of a 3-phenyl substituent led to a pronounced increase in inhibitory activity. Thus, the relatively bent biphenyl derivative (S)-13c was found to exhibit a K_i value of 42 nM. The investigated acylamides containing a fivemembered heterocycle, namely pyrrole (S)-13k, pyrazole derivatives (S)-13l and (S)-13m, as well as isoxazole (S)-13n, exhibited inhibitory activities being slightly superior to the one of benzamide (S)-13a. In case of the synthesized acylamides containing benzannelated heterocycles, quinoline derivatives (S)-13f and (S)-13e were found to be equally or slightly less active compared to (S)-13a. However, among the investigated indole derivatives, superior inhibitory activities were observed. While indole-2-carboxamide (S)-13g was found to be the least potent indole derivative with a K_i value of 87 nM, indole-3-carboxamide (S)-13h, indole-4-carboxamide (S)-13i, and indole-5-carboxamide (S)-13j exhibited K_i values

Figure 5. NMR experiments for the identification of fragments bound to the complex LpxC-9. (A) STD spectrum of fragment F3 in the presence of the complex LpxC-9. Arrows highlight two resonances of fragment F3 exhibiting the lowest STD factors, indicating that the corresponding protons are more solvent-exposed than the other protons. (B) Nuclear Overhauser effect spectroscopy (NOESY) spectrum showing transferred NOESY peaks observed for compounds 9 and F1 bound to LpxC. The largest interligand NOESY peak between 9 and F1 is shown in the square.

between 9 and 15 nM, thus representing the most potent LpxC inhibitors of the synthesized series of acylamides.

As Pseudomonas aeruginosa is one of the so-called ESKAPE pathogens,⁵⁶ being the leading cause of nosocomial infections throughout the world, and carbapenem-resistant P. aeruginosa strains were assigned the highest priority (priority 1: critical) within the "WHO priority pathogens list for R&D of new antibiotics" (WHO 2017),⁵⁷ new antibacterial agents combating these bacteria are urgently required. Thus, our newly developed LpxC inhibitors should be tested for inhibitory activity toward P. aeruginosa LpxC, which shares 57% sequence identity with *E. coli* LpxC.²⁸ Therefore, a liquid chromatography with tandem mass spectrometry (LC-MS/MS)-based P. aeruginosa LpxC assay was established. Even though UDP-3-O-[(R)-3-hydroxydecanoyl]-N-acetylglucosamine is the natural substrate of P. aeruginosa LpxC,⁵⁸ the E. coli LpxC substrate UDP-3-O-[(R)-3-hydroxymyristoyl]-N-acetylglucosamine (1) was employed, as LpxC had been found to be relatively nonspecific with respect to the acyl chain length of the substrate.⁵⁹⁻⁶¹ After incubation and subsequent termination of the enzymatic reaction, the samples were subjected to LC-MS/MS analysis. Following the chromatographic separation of substrate 1 and deacetylated product 2, the eluted compounds were ionized and subjected to fragmentation. Three multiple reaction monitoring (MRM) transitions were detected per compound [precursor ions: m/z (substrate 1) 832; m/z(deacetylated product 2) 790; product ions: m/z (product 1) 385, collision energy = -60 V; m/z (product 2) 159, collision energy = -80 V; m/z (product 3) 79, collision energy = -140V]. The MRM transitions $832 \rightarrow 79$ of substrate 1 and $790 \rightarrow$ 79 of product 2 were used as quantifiers; the other mass transitions were used as qualifiers.

Generally, the LC–MS/MS-based method represents an alternative to analyze compounds, which cannot be tested in the fluorescence-based assay, like primary amines and intrinsically fluorescent compounds. Additionally, it offers the advantage that lower substrate concentrations (10%) are used in the established protocol compared to the fluorescence-based assay, in which the detection of respectively low product concentrations would be difficult to detect due to inherently high background fluorescence. Thus, the conditions of the established LC–MS/MS-based enzyme assay were employed

to determine the $K_{\rm M}$ value for the *P. aeruginosa* LpxC-catalyzed deacetylation of UDP-3-O-[(*R*)-3-hydroxymyristoyl]-*N*-acetyl-glucosamine (1) (Figure S9, Supporting Information), which was found to be 4.7 μ M. Additionally, the $K_{\rm M}$ of *E. coli* LpxC C63A could be determined using the LC-MS/MS-based method (Figure S8, Supporting Information, vide supra).

With respect to their inhibitory activities toward P. aeruginosa LpxC, the investigated amides showed the same trends as previously observed against E. coli LpxC C63A (Table 3). Remarkably, the determined K_i values against P. aeruginosa LpxC were generally lower by a factor of around 1.3 to 4.5 than the ones against E. coli LpxC C63A. Only biphenyl derivative (S)-13c was found to exhibit a higher K_i value against P. aeruginosa LpxC than against E. coli LpxC C63A. Again, indole-3-carboxamide (S)-13h, indole-4-carboxamide (S)-13i, and indole-5-carboxamide (S)-13j were found to be the most potent LpxC inhibitors of the investigated series of acylamides, with compounds (S)-13h and (S)-13j exhibiting K_i values against P. aeruginosa LpxC in the single-digit nanomolar range. As in accordance with the results of the E. coli LpxC C63A assay, the (R)-configured indole-3-carboxamide (R)-13h showed considerably lower inhibitory activity toward P. aeruginosa LpxC than its enantiomer (S)-13h, the investigation of the other (R)-configured acylamides in the P. aeruginosa LpxC assay was omitted.

The isoserine-based amides were additionally tested for antibacterial activity (Table 3). Also in case of these compounds, the most potent inhibitors of E. coli LpxC C63A, namely indole derivatives (S)-13h, (S)-13i, and (S)-13j, were found to exhibit the lowest MIC values against E. coli BL21(DE3) (1-4 μ g/mL) and E. coli D22 (0.031 μ g/mL). Among the moderately active LpxC inhibitors (S)-13c, (S)-13g, (S)-13k, (S)-13l, and (S)-13m, biphenyl derivative (S)-13c showed no antibacterial activity against *E. coli* BL21(DE3). In contrast, indole-2-carboxamide (S)-13g, pyrrole (S)-13k, as well as pyrazole derivatives (S)-13l and (S)-13m were found to be only one serial dilution step less active against E. coli D22 $(0.063 \ \mu g/mL)$ and about one to two serial dilution steps less active against E. coli BL21(DE3) (4-8 μ g/mL) compared to indole derivatives (S)-13h, (S)-13i, and (S)-13j. Particularly, pyrrole derivative (S)-13k and pyrazole derivative (S)-13l

Table 3. Antibacterial and LpxC Inhibitory Activities of Isoserine-Based Hydroxamic Acids^a

compound		<i>K</i> i[r	nM]	міс [µg/mL]	Inhibition zone [mm]					
		E. coli LpxC C63A	P. aeruginosa LpxC	<i>E. coli</i> BL21(DE3)	E. coli D22	E. coli BL21(DE3)	E. coli D22	E. coli TOP 10	E. coli ATCC 35218	K. pneumoniae ATCC 700603	P. aeruginosa ATCC 27853
(S) -13a	(N ¹	149 ± 56	58.2 ± 0.1	8.0	0.13	15.7 ± 1.6	26.3 ± 1.5	23.3 ± 0.6	19.3 ± 0.6	7.0 ± 1.0	7.0 ± 1.0
(<i>R</i>)-13a		2560 ± 1080	n.d.	>16	4.0	9.7 ± 0.6	18.0 ± 2.0	n.d.	n.d.	n.d.	n.d.
(S) -13b		>2540	n.d.	>16	0.13	≤6	11.2 ± 2.9	<6	<6	<6	<6
(<i>R</i>)-13b		>2540	n.d.	>16	>16	≤6	≤6	n.d.	n.d.	n.d.	n.d.
(S)-13c		41.5 ± 5.7	149 ± 14	>16	0.13	≤6	19.0 ± 1.5	8.0 ± 0	7.7 ± 1.5	<6	6.3 ± 0.6
(<i>R</i>)-13c		>2540	n.d.	>16	>16	≤6	≤6	n.d.	n.d.	n.d.	n.d.
(S)-13d		323 ± 126	196 ± 18	>16	0.25	≤6	9.3 ± 1.6	<6	7.0 ± 1.0	<6	6.3 ± 0.6
(<i>R</i>)-13d		>2540	n.d.	>16	>16	≤6	≤6	n.d.	n.d.	n.d.	n.d.
(S)- 13e		377 ± 83	168 ± 8	>16	0.25	≤6	16.0 ± 2.5	8.0 ± 0	7.7 ± 1.5	<6	6.3 ± 0.6
(<i>R</i>)-13e		>2540	n.d.	>16	8.0	≤6	12.7 ± 1.6	n.d.	n.d.	n.d.	n.d.
(S) -13f	r r n 1	117 ± 21	91.1 ± 17.5	>16	0.25	11.5 ± 2.1	29.0 ± 1.0	18.0 ± 6.0	15.0 ± 0	<6	<6
(R)- 13f		>2540	n.d.	>16	>16	≤6	10.3 ± 1.2	n.d.	n.d.	n.d.	n.d.
(S)- 13g		87.0 ± 58.5	46.0 ± 16.4	8.0	0.063	10.0 ± 1.0	23.0 ± 1.0	16.0 ± 6.1	10.7 ± 0.6	<6	<6
(<i>R</i>)-13g		>2540	n.d.	>16	16	≤6	9.0 ± 1.0	n.d.	n.d.	n.d.	n.d.
(S)-13h		10.7 ± 1.2	5.91 ± 1.47	1.0	0.031	19.7 ± 1.6	29.3 ± 1.6	18.7 ± 5.8	19.0 ± 1.0	6.7 ± 1.2	7.0 ± 1.0
(<i>R</i>)-13h	HNJ	1580 ± 216	631 ± 198	>16	4.0	≤6	13.3 ± 1.0	n.d.	n.d.	n.d.	n.d.
(S)- 13 i	HN	15.3 ± 3.6	10.3 ± 2.0	4.0	0.031	20.4 ± 1.6	>30	23.0 ± 1.0	17.0 ± 3.5	7.3 ± 1.2	6.3 ± 0.6
(S)- 13 j		9.45 ± 1.01	5.58 ± 0.53	2.0	0.031	20.0 ± 1.0	29.0 ± 1.6	22.3 ± 3.8	21.0 ± 1.0	8.0 ± 1.7	8.3 ± 0.6
(R)- 13 j	NH NH	1150 ± 317	n.d.	>16	16	≤6	11.7 ± 1.2	n.d.	n.d.	n.d.	n.d.
(S) -13k	() NH	97.1 ± 9.4	29.0 ± 7.6	4.0	0.063	23.3 ± 0.6	>30	27.0 ± 2.8	23.0 ± 1.4	9.5 ± 0.7	9.0 ± 1.4
(S)-13I	HN-N	80.9 ± 9.0	18.0 ± 2.9	4.0	0.063	24.7 ± 0.6	>30	30.5 ± 0.7	25.5 ± 0.7	11.5 ± 0.7	12.0 ± 0
(S) -13m		85.2 ± 21.7	41.9 ± 18.5	8.0	0.063	11.3 ± 1.6	24.5 ± 1.8	14.7 ± 0.6	13.0 ± 1.7	<6	<6
(<i>R</i>)-13m		>2540	n.d.	>16	>16	≤6	9.7 ± 1.3	n.d.	n.d.	n.d.	n.d.
(S)-13n	H ₃ C Th	127 ± 40	80.4 ± 9.7	16	0.13	20.8 ± 1.1	>30	24.0 ± 0	20.0 ± 1.7	7.0 ± 1.0	<6
(<i>R</i>)-13n	CH3	>2540	n.d.	>16	8.0	≤6	16.3 ± 1.6	n.d.	n.d.	n.d.	n.d.

^an.d.: not determined.

caused the largest zones of growth inhibition in the performed disc diffusion assays against *E. coli* BL21(DE3) and *E. coli* D22.

As the (S)-configured isoserine-based amides exhibited higher antibacterial activities than their (R)-configured enantiomers, these compounds were further characterized by testing them against *E. coli* TOP10 and a series of Gramnegative wild-type strains representing clinically relevant species, namely *E. coli* ATCC 35218, *Klebsiella pneumoniae* ATCC 700603, and *P. aeruginosa* ATCC 27853. In the performed disc diffusion assays, indole derivatives (S)-13h,

Figure 6. Docking poses of potent LpxC inhibitors from the current study (PDB ID 4MQY). (A) (S)-13a, (B) (S)-13c, (C) (S)-13j, (D) (S)-13h. Inhibitors are shown in ball-and-stick mode. The zinc ion is shown as a cyan sphere (van der Waals radius), and hydrogen bonds are shown as redcolored dashed lines.

(S)-13i, and (S)-13j caused promising halos of growth inhibition. However, again the largest zones of growth inhibition were found for pyrrole derivative (S)-13k and pyrazole derivative (S)-13l, which showed considerable activity against K. pneumoniae ATCC 700603 and P. aeruginosa ATCC 27853.

Molecular Docking Studies. To rationalize the SAR of the inhibitors synthesized, molecular docking into the E. coli and P. aeruginosa LpxC structures (PDB ID: 4MQY and 5VWM) was carried out using the program Glide (Figure 6, Table S1, for details see Computational Methods). The docked inhibitors showed a similar binding mode in both enzymes, which resembled the commonly observed binding mode of known LpxC inhibitors, such as CHIR-090 (Figure S1) and LPC-011 (3a, Figure 2). Since more in vitro data were obtained for the E. coli LpxC orthologue and since the observed inhibition data for both forms show high correlation, only the docking results observed for the E. coli orthologue are discussed here. Generally, the hydroxamic acid moiety of the inhibitors coordinates to the Zn^{2+} -ion in a bidentate fashion. The hydroxamic acid group further exhibits hydrogen bond interactions with E78 and T191. The lipophilic side chain is located in a hydrophobic tunnel comprising L18, L62, I198, C207, F212, and V217. The substituents introduced at the α -

carbon atom of hydroxamic acid 9 (indicated by the arrow in Figure 4) protrude into the UDP pocket of LpxC and can be used for adding pocket-filling substituents. From the first series of synthesized inhibitors (Table 1), the (2S)-configured isoserine-based amide (S)-13a demonstrated the most favorable LpxC inhibitory activity. The docking pose of (S)-13a showed an additional hydrogen bond between the amide group of the side chain and the backbone CO of F192, whereas the terminal phenyl group is involved in aromatic interactions with the side chain of F192 (Figure 6A). These interactions might contribute to the enhanced inhibitory activity compared to the analogues that do not show this hydrogen bond [e.g., ether (S)-11a]. Due to the favorable hydrogen bond of the amide linker, it was retained for further optimization of the derivatives (Table 3). The size of the aromatic substituent is another option to improve the binding in the UDP pocket, as can be seen in the interaction of (S)-13c (Figure 6B). The additional aromatic substituent in the meta-position forms aromatic interactions with F192 and F194, which is not the case with the para-substituted isomer (*S*)-13b. The phenyl ring located in the para-position of (S)-13b does not fit properly into the UDP-binding pocket and results in a loss of the correct bidentate chelation of the hydroxamic acid group.

In case of the most active inhibitor (S)-13j, the terminal indole ring is involved in a hydrogen bond with D197 (Figure 6C). In case of the (*R*)-configured isomers [e.g., (*R*)-13j, (*R*)-13c], the binding is less favorable, and in all cases, a hydrogen bond between the amide and the backbone CO of F192 was not observed. In addition, the interaction of the aromatic substituents is less favorable indicated by weaker docking scores. For the second and third most potent inhibitors (*S*)-13h and (*S*)-13i, a direct hydrogen bond between the indole and D197 was not observed. However, the distance is short enough that a water molecule could mediate a hydrogen bond. The aromatic interaction with F192 is similar to that observed for (*S*)-13j (shown exemplarily for (*S*)-13h in Figure 6D).

In order to test the stability and the dynamic behavior of potent inhibitors and their less active enantiomers, we selected the two pairs (S)-13c/(R)-13c and (S)-13j/(R)-13j and analyzed them by means of molecular dynamics (MD) simulations. For comparison, we also simulated the crystal structure of LpxC in complex with the inhibitor LPC-138 (Figure S1, PDB structure 4MQY).⁶² The MD simulations of the crystal structure (PDB ID: 4MQY) and the two potent inhibitors (S)-13c and (S)-13j showed that the complexes and the interactions of the inhibitors are very stable (Figures S2-S7). This is reflected in small root-mean-square deviation (RMSD) changes for the ligand, the chelation of the zinc ion, and the stable hydrogen bonds. In contrast, simulations of the less active stereoisomers (R)-13c and (R)-13j showed that although the protein structure was stable, the inhibitors showed significant changes in the interactions, including loss of chelation in case of (R)-13c.

Further Biological Evaluation. Inhibition of the Virulence Factor LasB from P. aeruginosa. The elastase LasB of P. aeruginosa is a secreted metalloprotease, which actively controls the infection process and is considered an important virulence factor in this pathogen.^{63,64} Thus, inhibitors of LasB are described to reduce pathogenicity without being bactericidal.^{65,66} These inhibitors could be applied together with a standard-of-care antibiotic in combination therapy. Hence, an additional antivirulence effect via LasB inhibition could be beneficial for the LpxC inhibitors, leading to, e.g., a dual LpxC/LasB inhibitor.

As aldotetronic acid derivative (2S,3S)-10 as well as its stereoisomer (2S,3R)-10 had been found to exhibit promising inhibitory activities against LasB,³² the newly synthesized hydroxamic acids were also evaluated for inhibitory activity toward the zinc metalloprotease (Table 4). As in case of the aldotetronic acid derivatives (2S)-configuration had been found to be crucial for inhibitory activity toward LasB, generally only the (S)-enantiomers of the newly synthesized compounds were investigated.

In case of benzyl ether (S)-11a and benzylamine (S)-12a, IC₅₀ values of 14.3 and 21.9 μ M were found, respectively. These were in the same range as those determined for aldotetronic acid derivatives (2S,3S)-10 and (2S,3R)-10. In contrast, tertiary amine (S)-37 exhibited no inhibitory activity.

While benzamide (S)-13a showed moderate inhibition of the enzymatic activity of LasB at a concentration of 20 μ M, being similarly potent as indole derivatives (S)-13h and (S)-13i as well as pyrrole derivative (S)-13k (20–30% inhibition at 20 μ M), little to no inhibitory activity was found for biphenyl derivatives (S)-13b and (S)-13c, diphenyl ether (S)-13d, quinoline derivatives (S)-13e and (S)-13f, indole derivatives (S)-13g and (S)-13j, as well as pyrazole derivatives (S)-13l and

Table 4.	In	Vitro	Activities	of the	Newly	Synthesized
Hydroxa	mic	Acids	s against I	LasB ^a		

com	npound	LasB				
HO.N.C.		Inhibition at 20 μΜ [%]	/C₅₀ [µM]			
(2S,3S) -10 ³²		61.0 ± 0.6	17.7 ± 0.8			
(2 <i>S</i> ,3 <i>R</i>)-10 ³²		64.9 ± 3.2	14.8 ± 0.4			
(2 <i>R</i> ,3 <i>S</i>)-10 ³²	ножол	n.i.				
(2 <i>R</i> ,3 <i>R</i>) -10 ³²		n.i.				
(2 <i>S</i> ,3 <i>S</i>) -40 ³²	HO + O + O O O O O O O O O O O O O O O O	n.i.				
(S) -11a		62.3 ± 3.2	14.3 ± 0.9			
(S)- 12a	HN TH	57.0 ± 2.0	21.9 ± 1.2			
(S) -37		n.i.				
(S) -13a	HN TO	20.1 ± 3.9				
(<i>S</i>)-13b	HN TO	n.i.				
(S) -13c	HR HR	n.i.				
(S)-13d		n.i.				
(S)- 13e		11.6 ± 1.9				
(S)-13f		n.i.				
(S) -13g		n.i.				
(S)-13h		25.2 ± 5.6				
(S)- 13 i	HN HN	27.6 ± 2.9				
(S)- 13j		n.i.				
(S)-13k	HN CNH	22.4 ± 3.1				
(S)-13I	HN-N	11.6 ± 0.7				
(S) -13m		n.i.				
(S)-13n	H ₃ C HN	92.7 ± 0.7	1.80 ± 0.15			
(<i>R</i>)-13n	№́Д ^с о	n.i.				

^an.i.: <10% inhibition.

(S)-13m. While in case of the previously investigated aldotetronic acid derivatives, the replacement of the benzyl group by a (3,5-dimethylisoxazol-4-yl)methyl substituent was

found to be detrimental for LasB inhibition [cf. (2S,3S)-40], the exchange of the benzoyl moiety of isoserine-based benzamide (S)-13a by a 3,5-dimethylisoxazole-4-carbonyl group, leading to hydroxamic acid (S)-13n, caused a considerable increase in inhibitory activity. With an IC₅₀ value of 1.8 μ M, isoxazole (S)-13n is the most potent LasB inhibitor of the investigated series of hydroxamic acids and represents a promising starting point for further optimization steps.

To confirm the superiority of the stereoisomers with (S)configuration in the α -position of the hydroxamate moiety, isoxazole derivative (R)-13n, the enantiomer of (S)-13n, was evaluated for inhibitory activity toward LasB. Just like in case of the aldotetronic acid derivatives, the inversion of configuration in the α -position of isoserine-based amide (S)-13n was detrimental for LasB inhibition, with enantiomer (R)-13n exhibiting no inhibitory activity.

In Vitro ADMET Properties and Off-Targets. As indole derivatives (S)-13h and (S)-13j as well as pyrazole derivative (S)-13l were found to exhibit promising antibacterial as well as LpxC inhibitory activities, the three frontrunner compounds were further investigated regarding their in vitro absorption—distribution—metabolism—excretion—toxicity (ADMET) profile (Table 5).

Table 5. In Vitro ADMET Profile of (S)-13h, (S)-13j, and (S)-13l

	(S) -13h	(S) -13j	(S) -13
	Port HN	$\langle D \rangle^{*}$	KN-N
microsomal metabolic stability			
mouse			
tup [min]	stable stable		53 3
the finite	> 60	> 60	00.0
Clist [u] /min/ma protein]	stable	stable	26.0
	< 23	< 23	20.0
human			
tua [min]	stable	stable	stable
tuz [rinii]	> 60	> 60	> 60
Cliet [u] /min/ma protein]	stable	stable	stable
	< 23	< 23	< 23
metabolic stability in mouse liver			
S9 fractions			
t _{1/2} [min]	15.5 ± 1.0	25.5 ± 2.7	64.0 ± 5.1
Clint [µL/min/mg protein]	44.8 ± 2.9	27.6 ± 2.7	10.9 ± 0.9
mouse plasma stability			
t _{1/2} [min]	22.9 ± 4.8	43 ± 13	24.2 ± 5.0
plasma protein binding			
mouse [%]	94.59 ± 1.3	92.05 ± 1.7	79.42 ± 2.5
human [%]	98.25 ± 0.2	97.95 ± 0.3	97.76 ± 1.4
logD _{7.4}	3.63	3.33	2.43
Kinetic solubility (1% DMSO/PBS)	535+19	46 + 11	>100
[µM]	00.0 ± 1.0	40 ± 11	- 100
HepG2 cytotoxicity			
/C ₅₀ [μM]	15.1 ± 1.7	8.1 ± 1.4	30.2 ± 1.9

Metabolism with liver microsomes did show that the compounds were generally stable in humans and mice, with pyrazole derivative (S)-13l being the least stable with a half-life of around 53 min in the presence of mouse liver microsomes. In contrast to the metabolism with liver microsomes, the metabolism in liver S9 fractions revealed lower stability of indole derivative (S)-13h with a clearance of around 45 μ L/min/mg protein, whereas indole derivative (S)-13j had a significantly lower clearance similar to the one observed in microsomes. Superior metabolic stability was found for the pyrazole-substituted compound (S)-13l with a moderate half-

life of more than 60 min. Also, the plasma stability of the compounds in mouse plasma was generally low with half-lives between 22.9 min [(S)-13h] and 43.1 min [(S)-13j].

In general, plasma protein binding in humans was high with the compounds having a plasma protein binding above 97%. Interestingly, the compounds exhibited lower plasma protein binding in mice compared to humans, which was considerably pronounced in case of pyrazole derivative (S)-13l (<80%), resulting in a much higher unbound fraction.

With a log $D_{7.4}$ of 2.43, the pyrazole-substituted compound (*S*)-13l is also the least lipophilic of the three compounds. And while the two indole derivatives showed similar kinetic solubility around 50 μ M, pyrazole-substituted (*S*)-13l was found to be soluble up to 100 μ M. This compound was also the least cytotoxic among the ones tested with an IC₅₀ value of 30.2 μ M (vs 15.1 and 8.1 μ M).

Since LpxC and LasB are zinc-dependent enzymes, we investigated potential off-target activity of the three frontrunners on mammalian zinc-dependent enzymes. Our panel comprised three representative matrix-metalloproteases (MMP1–3) and tumor necrosis factor- α converting enzyme (TACE, also known as ADAM17) (Table 6). MMP inhibition

Table 6. In Vitro Activities of Hydroxamic Acids (S)-13l, (S)-13h, and (S)-13j against MMPs 1-3 and TACE (ADAM-17)^{*a*}

	MMP1	MMP2	ММР3	TACE
	(% inhibition	(% inhibition	(% inhibition	(% inhibition
	at 100 µM)	at 100 µM)	at 100 µM)	at 20 µM)
(S) -13h	15 ± 7	15 ± 3	n.i.	n.i.
(S) -13j	n.i.	47 ± 3	16 ± 1	38 ± 5
(0) 421	11 ± 6	65 ± 4	19 ± 2	81 ± 2
(3)-131		$/C_{50} = 67 \pm 4 \ \mu M$		$IC_{50} = 5.4 \pm 0.9 \ \mu M$

^{*a*}n.i.: <10% inhibition.

was mostly below 20% at 100 μ M, except for MMP2: here, (S)-13j resulted in ~50% inhibition at 100 μ M and (S)-13l gave an IC₅₀ value of 67 μ M. Both compounds were also active against TACE, again with stronger inhibition by (S)-13l [IC₅₀ 5.4 μ M vs 38% inhibition at 100 μ M by (S)-13j]. Considering the potent activity on LpxC, this compound still shows excellent to moderate selectivity (selectivity factor 105 for MMP2 and 8.4 for TACE).

CONCLUSIONS

In this study, we aimed to generate LpxC inhibitors that address the UDP-binding pocket, which is located in close proximity to the Zn^{2+} -chelating hydroxamate moiety of our previously reported inhibitors.

Thus, fragment screening was utilized as the strategy to optimize our benzyloxyacetohydroxamic acid derivatives. We used STD-NMR, WaterLogsy, and subsequent NMR-ILOE experiments to identify nine out of 650 fragments, which bind into the UDP-binding pocket of LpxC near the methylene group in the α -position of the hydroxamate moiety of LpxC inhibitor 9, and four substructures (biphenyl, indole, quinoline, and pyrazole) were selected to be linked to the benzyloxyacetic acid scaffold of hydroxamic acid 9.

In parallel, in order to find an optimal linker to connect the structures of interest with the benzyloxyacetohydroxamic acid derivatives, a phenyl ring as an exemplary substituent was connected to the scaffold of hydroxamic acid **9** via structural elements of different lengths and structures. The biological evaluation of the investigated tetrose- and glyceric acid-derived ethers as well as the isoserine-based amides and secondary and tertiary amines revealed the amide linker of isoserine derivative (*S*)-**13a** to be the most favorable with respect to inhibitory activity toward LpxC (Table 1). Besides holding the introduced fragment in an appropriate distance to the benzyloxyacetohydroxamic acid scaffold, molecular docking studies indicate that the amide linker undergoes favorable hydrogen bonding with the enzyme (backbone CO of F192).

Among the synthesized amides exhibiting fragment-derived substituents as well as further acyl residues, the (S)-configured indole derivatives (S)-13h, (S)-13i, and (S)-13j as well as pyrazole derivatives (S)-13l and (S)-13m exhibited highest inhibitory activities toward E. coli LpxC C63A as well as P. aeruginosa LpxC (Table 3). Thus, linking these two substructures, which were identified through fragment screening using NMR spectroscopy, to the scaffold of hydroxamic acid 9 resulted in more potent LpxC inhibitors than linking the other investigated residues. Consequently, by appending a privileged fragment via a suitable linker to the scaffold of hydroxamic acid 9, this micromolar LpxC inhibitor could be converted into the one-digit nanomolar LpxC inhibitor (S)-13j $[K_i (EcLpxC C63A) = 9.5 nM; K_i (PaLpxC): 5.6 nM].$ Therefore, NMR-based fragment screening proved to be an appropriate strategy for the fragment-based drug discovery of LpxC inhibitors specifically addressing the enzyme's UDPbinding site.

Molecular docking into an *E. coli* LpxC structure indicated that the substituents introduced at the α -carbon atom of hydroxamic acid 9 protrude into the UDP pocket of LpxC with the amide linker of the (*S*)-configured isomers undergoing hydrogen bonding with the backbone CO of F192. The aromatic substituents enter the substrate-ribose region of the UDP-binding pocket through the amide linker (Figure 4). In case of the most active inhibitor (*S*)-13*j*, besides undergoing aromatic interactions with the side chain of F192, the indole ring is involved in a hydrogen bond with D197 being indicative of its high inhibitory activity (Figure 6).

Indole derivatives (S)-13h, (S)-13i, and (S)-13j, the most potent LpxC inhibitors of the investigated isoserine-based amides, were found to exhibit the lowest MIC values against *E.* coli BL21(DE3) (1-4 μ g/mL) and *E.* coli D22 (0.031 μ g/mL) (Table 3). In the performed disc diffusion assays against the Gram-negative wild-type strains *E.* coli ATCC 35218, *K.* pneumoniae ATCC 700603, and *P.* aeruginosa ATCC 27853, indole derivatives (S)-13h, (S)-13i, and (S)-13j caused promising halos of growth inhibition. However, the largest zones of growth inhibition were found for pyrrole derivative (S)-13k and pyrazole derivative (S)-13l.

As our previously reported benzyloxyacetohydroxamic acid derivatives had shown inhibitory activity toward the elastase LasB of *P. aeruginosa*,³² the newly developed isoserine-based amides were also tested against this important virulence factor (Table 4). While indole derivatives (S)-13g, (S)-13h, (S)-13i, and (S)-13j as well as pyrazole derivatives (S)-13l and (S)-13m exhibited little to no inhibitory activity toward LasB, the (S)-configured 3,5-dimethylisoxazol-4-yl derivative (S)-13n was found to be the most potent LasB inhibitor of the investigated series of hydroxamic acids, exhibiting an IC₅₀ value of 1.8 μ M. Thus, the isoserine-based amides represent a promising starting point for further optimization steps, as depending on the introduced acyl residue, the compounds can be developed into selective or dual inhibitors of the two Zn²⁺-dependent enzymes.

Additionally, the three frontrunner compounds (S)-13h, (S)-13j, and (S)-13l were tested for inhibitory activity toward several mammalian zinc-dependent enzymes (MMP1-3 and TACE), representing potential off-targets (Table 6). The compounds exhibited good selectivities for LpxC, with pyrazole derivative (S)-13l being the least selective compound, particularly with respect to MMP2 and TACE inhibition.

The three frontrunner compounds, exhibiting promising antibacterial as well as LpxC inhibitory activities, were additionally investigated regarding their in vitro ADMET profile (Table 5). While the metabolic stability of the compounds in the presence of human and mouse liver microsomes was generally high, particularly the stability of indole derivative (S)-13h in the presence of mouse liver S9 fractions, including phase I and phase II reactions, was low. These findings indicate, that the metabolism of these compounds might not be mainly CYP-mediated, as S9 fractions harbor additional non-CYP-enzymes, or might be a cause of phase II metabolism.^{67,68} As plasma stability of the compounds in mouse plasma was also found to be low, their stability needs to be improved in further optimization steps.

Among the three frontrunner compounds, the pyrazolesubstituted compound (S)-13l was the least lipophilic with a $\log D_{7.4}$ of 2.43. This is in line with the higher kinetic solubility and the considerably lower plasma protein binding in mice of the compound compared to the two indole derivatives. In comparison to marketed Gram-negative antibacterials, which exhibit an average $c \log D$ of -2.8,⁶⁹ all of our compounds are substantially more lipophilic. However, the whole class of LpxC inhibitors proved to be an exception to the trend that primarily hydrophilic compounds exhibit potent P. aeruginosa and E. coli activity.⁷⁰ Nonetheless, as the UDP-binding site of LpxC is solvent-exposed, in further optimization steps of the developed isoserine-based amides, additional polar functional groups could be introduced at the substituent, addressing the UDP-binding site to reduce the lipophilicity of the inhibitors. Additionally, having found compounds tightly binding to the UDP-binding site, their lipophilic side chain could be shortened to reduce the impact of this hydrophobic moiety on the polarity of the compounds.

To date, a lot of preclinical work has been done on LpxC inhibitors with promising results, showing the feasibility of developing hydroxamate-based LpxC inhibitors into clinical candidates, which in contrast to ACHN-975 (Figure S1), the first LpxC inhibitor to reach human clinical trials, do not exhibit cardiovascular toxicity.^{22–24,71,72} Altogether, our initial screening revealed some ADMET parameters that need to be considered during the further development of our newly developed isoserine-based amides, paving the way for multiparameter optimization with the aim to merge compound features favorable for activity with those improving the in vitro ADMET profile as well as selectivity.

EXPERIMENTAL SECTION

Chemistry, General. All experiments involving water- or airsensitive compounds were carried out under anhydrous conditions (N_2 atmosphere). Reagents were purchased from various suppliers

and were used without further purification unless otherwise noted. Anhydrous solvents were purchased from Acros Organics (extra dry over molecular sieves). Solvents for flash column chromatography were purchased in technical grade and distilled prior to use. Ultrapure water for reversed-phase chromatography was purified using a Sartorius Arium pro system (Sartopore 0.2 μ m, UV). Acetonitrile (ACN) for reversed-phase chromatography was purchased from VWR (HPLC grade). Flash column chromatography on silica gel was performed using Macherey Nagel silica gel 60 M (0.040-0.063 mm). Parentheses include the diameter of the column, fraction size, eluent, and R_f value. Thin-layer chromatography was performed on Macherey Nagel precoated TLC sheets (ALUGRAM Xtra SIL G/UV₂₅₄). Visualization was achieved by heat-staining using a cerium molybdate dipping bath $[Ce(SO_4)_2 (1.8 \text{ g}), (NH_4)_6Mo_7O_{24} \times 4H_2O (45 \text{ g}),$ conc. H₂SO₄ (45 g), H₂O (900 mL)]. Automatic reversed-phase flash column chromatography was performed using Biotage SNAP Ultra C18 columns on an Isolera One (Biotage). Product-containing fractions were combined and lyophilized using a Christ Alpha 2-4 LDplus freeze-dryer. Automatic normal-phase flash column chromatography was performed using Biotage SNAP Ultra HP-Sphere columns on an Interchim puriFlash XS 420 system. Productcontaining fractions were combined, and the solvent was removed in vacuo. Melting points were measured with a Büchi Melting Point M-565 and are uncorrected. Optical rotation α [deg] was determined with a P8000 polarimeter (A. Krüss Optronic GmbH); path length 1 dm, wavelength 589 nm (sodium D line); the unit of the specific rotation $[\alpha]_D^{20}$ (deg mL dm⁻¹ g⁻¹) is omitted; the concentration of the sample $c (mg mL^{-1})$ and the solvent used are given in brackets. IR spectra were recorded on a Bruker Alpha FT-IR Platinum ATR spectrophotometer. NMR spectra were recorded at ambient temperature on Bruker Avance I 400, DRX 500, and Avance III 600 instruments. High-resolution mass spectrometry was performed using an Agilent 6224 ESI-TOF instrument via flow injection analysis in 50:50 water + 0.1% formic acid/acetonitrile + 0.1% formic acid at a flow rate of 0.3 mL/min via electrospray ionization. HPLC methods for the determination of product purity: method 1: VWR Hitachi equipment; UV/vis detector: 5420; autosampler: 5260; pump: 5160; column: LiChrospher 60 RP-select B (5 µm); LiChroCART 250-4 mm cartridge; flow rate: 1.00 mL/min; injection volume: 5.0 μ L; detection at $\lambda = 210$ nm for 30 min; solvents: (A) water with 0.05% (V/V) trifluoroacetic acid, (B) acetonitrile with 0.05% (V/V)trifluoroacetic acid; gradient elution: (A %): 0-4 min: 90%, 4-29 min: gradient from 90 to 0%, 29-31 min: 0%, 31-31.5 min: gradient from 0 to 90%, 31.5-40 min: 90%; data were collected and evaluated by Chromaster software. Method 2: VWR Hitachi equipment; UV/vis detector: 5420; autosampler: 5260; pump: 5160; column: Phenomenex Gemini 5 μ m C6-Phenyl 110 Å; LC Column 250 × 4.6 mm; flow rate: 1.00 mL/min; injection volume: 5.0 μ L; detection at λ = 254 nm for 20 min; solvents: (A) acetonitrile/10 mM ammonium formate = 10:90 with 0.1% formic acid, (B) acetonitrile/10 mM ammonium formate = 90:10 with 0.1% formic acid; gradient elution: (A %): 0-5 min: 100%, 5-12 min: gradient from 100 to 0%, 12-20 min: 0%, 20-22 min: gradient from 0 to 100%, 22-30 min: 100%; data were collected and evaluated by Chromaster software. Method 3: VWR Hitachi equipment; UV/vis detector: 5420; autosampler: 5260; pump: 5160; column: LiChrospher 60 RP-select B (5 μ m); LiChroCART 250-4 mm cartridge; flow rate: 1.00 mL/min; injection volume: 5.0 μ L; detection at λ = 210 nm for 40 min; solvents: (A) water with 0.05% (V/V) trifluoroacetic acid, (B) acetonitrile with 0.05% (V/V) trifluoroacetic acid: gradient elution: (A %): 0-4 min: 90%, 4-29 min: gradient from 90 to 0%, 29-41 min: 0%, 41-41.5 min: gradient from 0 to 90%, 41.5-50 min: 90%; data were collected and evaluated by Chromaster software. Method 4: KNAUER equipment; UV/vis detector: Azura UVD 2.1S; pump: Azura P4.1S; column: Daicel Chiralpak IA; flow rate: 1.00 mL/min; injection: manual, injection valve; injection volume: 300 μ L; detection at λ = 230 nm for 20 min; solvent: isohexane/isopropanol = 97.5:2.5; data were collected and evaluated by the software ClarityChrom Preparativ Version 5.0.5.98. The purity of all test compounds was \geq 95%.

Synthetic Procedures. (S)-tert-Butyl(oxiran-2-ylmethoxy)diphenylsilane (15). Under a N2 atmosphere, tert-butyldiphenylsilyl chloride (9.5 mL, 10 g, 37 mmol) was added to an ice-cooled solution of (R)-glycidol (2.0 mL, 2.3 g, 31 mmol) and imidazole (3.0 g, 43 mmol) in dry dichloromethane (60 mL). After stirring the reaction mixture for 10 min at 0 °C, the ice-bath was removed and the mixture was stirred for 24 h at ambient temperature. Afterward, the mixture was washed with water and brine. The organic layer was dried (Na_2SO_4) , filtered, and the solvent was removed in vacuo. The residue was purified by flash column chromatography ($\emptyset = 6 \text{ cm}, h =$ 22 cm, V = 50 mL, petroleum ether/ethyl acetate = 10:1, $R_f = 0.59$) to give 15 as a colorless oil (8.3 g, 27 mmol, 86%). $[\alpha]_D^{20} = +2.2$ (1.9, methanol); ¹H NMR (DMSO- d_6): δ [ppm] = 1.00 (s, 9H, $SiC(CH_3)_3$, 2.57 (dd, J = 5.2/2.7 Hz, 1H, CH_2CHCH_2OSi), 2.71 (dd, J = 5.2/4.2 Hz, 1H, CH₂CHCH₂OSi), 3.10-3.17 (m, 1H, CH₂CHCH₂OSi), 3.59 (dd, J = 11.9/5.3 Hz, 1H, CH₂CHCH₂OSi), 3.91 (dd, J = 11.9/2.7 Hz, 1H, CH₂CHCH₂OSi), 7.39–7.50 (m, 6H, 3'-H_{diphenylsilyb} 4'-H_{diphenylsilyb} 5'-H_{diphenylsilyl}), and 7.60–7.67 (m, 4H, 2'-H_{diphenylsilyb} 6'-H_{diphenylsilyl}); ¹³C NMR (DMSO- d_6): δ [ppm] = 18.8 (1C, SiC(CH₃)₃), 26.5 (3C, SiC(CH₃)₃), 43.4 (1C, CH₂CHCH₂OSi), 51.7 (1C, CH₂CHCH₂OSi), 64.4 (1C, CH₂CHCH₂OSi), 127.9 (4C, C-3'_{diphenylsilyl}, C-5'_{diphenylsilyl}), 129.9 (2C, C-4'_{diphenylsilyl}), 132.76 (1C, C-1'_{diphenylsilyl}), 132.79 (1C, C-1'_{diphenylsilyl}), 127.90 (1C, C-1'_{diphenylsilyl}), 132.79 (1C, C-1'_{diphenylsilyl}), 127.90 (1C, C-1'_{diphenylsilyl}), 127 $1'_{diphenylsilyl}$, 135.0 (2C, C-2'_{diphenylsilyl}, C-6'_{diphenylsilyl}), and 135.1 (2C, C-2'_{diphenylsilyl}, C-6'_{diphenylsilyl}); IR (neat): $\tilde{\nu}$ [cm⁻¹] = 3071, 3050, 2998, 2958, 2930, 2893, 2857, 1472, 1427, 1390, 1361, 1254, 1159, 1106, 1089, 980, 917, 823, 738, 699, 612, 503, 486, and 425; HRMS (m/z): $[M + Na]^+$ calcd for $C_{19}H_{24}NaO_2Si$, 335.1438; found, 335.1461; HPLC (method 3): $t_{\rm R} = 28.3$ min, purity 99.5%.

(S)-1-(Benzyloxy)-3-[(tert-butyldiphenylsilyl)oxy]propan-2-ol (16). A mixture of 15 (380 mg, 1.2 mmol), erbium(III) triflate (75 mg, 0.12 mmol), and benzyl alcohol (0.15 mL, 160 mg, 1.4 mmol) was stirred at ambient temperature for 24 h. Then, a saturated aqueous solution of NaHCO₃ (25 mL) was added, and the mixture was extracted with dichloromethane $(3\times)$. The combined organic layers were dried (Na₂SO₄), filtered, and the solvent was removed in vacuo. The residue was purified by flash column chromatography (Ø = 4 cm, h = 24 cm, V = 30 mL, petroleum ether/ethyl acetate = 4:1, R_{f} = 0.53) to give 16 as a colorless oil (380 mg, 0.89 mmol, 73%). $[\alpha]_{D}^{-2}$ = +4.1 (2.2, methanol); ¹H NMR (DMSO- d_6): δ [ppm] = 0.97 (s, 9H, SiC(CH₃)₃), 3.46 (dd, J = 9.7/5.6 Hz, 1H, CH₂CHCH₂OSi), 3.57 (dd, J = 9.7/4.8 Hz, 1H, CH₂CHCH₂OSi), 3.60 (dd, J = 10.1/5.2 Hz, 1H, CH_2CHCH_2OSi), 3.63 (dd, J = 10.1/5.8 Hz, 1H, CH2CHCH2OSi), 3.73-3.79 (m, 1H, CH2CHCH2OSi), 4.49 (s, 2H, OCH₂Ph), 4.87 (d, J = 5.3 Hz, 1H, CHOH), 7.26-7.29 (m, 1H, 4'-Hbenzyl), 7.29-7.35 (m, 4H, 2'-Hbenzyl 3'-Hbenzyl 5'-Hbenzyl 6'-Hbenzyl), 7.37–7.43 (m, 4H, 3"-H_{diphenylsilyl}, 5"-H_{diphenylsilyl}), 7.43–7.47 (m, 2H, 4"-H_{diphenylsilyl}), and 7.60–7.65 (m, 4H, 2"-H_{diphenylsilyl}, 6"-H_{diphenylsilyl}); ¹³C NMR (DMSO- d_6): δ [ppm] = 18.8 (1C, SiĆ(ĆH₃)₃), 26.6 (3C, SiC(CH₃)₃), 65.1 (1C, CH₂CHCH₂OSi), 70.0 (1C, CH₂CHCH₂OSi), 71.3 (1C, CH₂CHCH₂OSi), 72.3 (1C, OCH₂Ph), 127.3 (1C, C-4'_{benzyl}), 127.4 (2C, C-2'_{benzyl}, C-6'_{benzyl}), UCH₂Pn), 127.3 (1C, C-4 benzyl), 127.4 (2C, C-2 benzyl), C-6 benzyl), 127.80 (2C, C-3" diphenylsilyl), C-5" diphenylsilyl), 127.81 (2C, C-3" diphenylsilyl) C-5" diphenylsilyl), 128.2 (2C, C-3' benzyl), 129.75 (1C, C-4" diphenylsilyl), 133.08 (1C, C-1" diphenylsilyl), 133.10 (1C, C-1" diphenylsilyl), 135.06 (2C, C-2" diphenylsilyl), 6" diphenylsilyl), 135.06 (2C, C-2" diphenylsilyl), 138.55 (1C, C-1' benzyl); IR (neat): $\tilde{\nu}$ [cm⁻¹] = 3070, 2930, 2857, 1472, 1454, 1427, 1390, 1361, 1105, 1028, 998, 823, 738, 697, 611, 503, and 487. 1427, 1390, 1361, 1105, 1028, 998, 823, 738, 697, 611, 503, and 487; HRMS (m/z): $[M + Na]^+$ calcd for C₂₆H₃₂NaO₃Si, 443.2013; found, 443.2013; HPLC (method 3): $t_{\rm R}$ = 28.8 min, purity 98.3%.

(*R*)-3-(*Benzyloxy*)-2-[(4-iodobenzyl)oxy]propan-1-ol (17). Under a N₂ atmosphere, sodium hydride (60% suspension in paraffin oil, 700 mg, 18 mmol) was added to an ice-cooled solution of 16 (1.8 g, 4.2 mmol) in anhydrous THF (40 mL). After stirring the reaction mixture for 20 min at 0 °C, 4-iodobenzyl bromide (1.5 g, 5.0 mmol) was added, and the reaction mixture was stirred for an additional 15 min at 0 °C. Then, the ice-bath was removed, and the mixture was stirred for 24 h at ambient temperature. Afterward, methanol was added under ice-cooling, and the solvent was removed in vacuo. The

residue was dissolved in THF (60 mL), tetrabutylammonium fluoride trihydrate (1.8 g, 5.8 mmol) was added, and the reaction mixture was stirred for 6 h at ambient temperature. Then, the solvent was removed in vacuo, water was added, and the mixture was extracted with ethyl acetate (3×). The combined organic layers were dried (Na_2SO_4) , filtered, and the solvent was removed in vacuo. The residue was purified by flash column chromatography ($\emptyset = 6 \text{ cm}, h = 23 \text{ cm}, V =$ 50 mL, petroleum ether/ethyl acetate = 2:1, $R_f = 0.24$) to give 17 as a colorless oil (1.2 g, 2.9 mmol, 71%). $[\alpha]_{D}^{20} = -2.1$ (2.0, methanol); ¹H NMR (DMSO- d_6): δ [ppm] = 3.44-3.61 (m, 5H, CH_2CHCH_2OH), 4.48 (s, 2H, CH_2OCH_2Ph), 4.55 (d, J = 13.0Hz, 1H, CHOCH₂Ar), 4.59 (d, J = 13.0 Hz, 1H, CHOCH₂Ar), 4.68 (t, J = 5.5 Hz, 1H, CH₂OH), 7.13–7.19 (m, 2H, 2"-H_{4-iodophenyl}, 6"-CH₂Ar), 72.3 (1C, CH₂OCH₂Ph), 79.1 (1C, CH₂CHCH₂OH), 93.1 (1C, C-4"_{4-iodophenyl}), 127.38 (1C, C-4'_{phenyl}), 127.40 (2C, C-2'_{phenyl}) C-6' phenyl), 128.2 (2C, C-3' phenyl), C-5' phenyl), 129.6 (2C, C-2'' 4-iodophenyl), C-6'' 4-iodophenyl), 136.9 (2C, C-3'' 4-iodophenyl), C-5'' 4-iodophenyl), 138.5 (1C, C-1'' phenyl), and 139.0 (1C, C-1'' 4-iodophenyl); IR (neat): $\tilde{\nu}$ [cm⁻¹] = 4.12, 2.62, 1.552, 1.052 3413, 2862, 1589, 1483, 1453, 1401, 1364, 1205, 1058, 1006, 799, 735, 696, 609, and 469; HRMS (m/z): $[M + Na]^+$ calcd for $C_{17}H_{19}INaO_3$, 421.0271; found, 421.0263; HPLC (method 1): $t_R =$ 23.8 min, purity 98.6%.

(4R,4'R,4"R,5'R)-2,2,2',2',2",2"-Hexamethyl-4,4':5',4"-ter(1,3-dioxolane) (21). The compound was synthesized according to the literature with minor variations:⁴¹ Concentrated sulfuric acid (1 mL) was added to a suspension of D-mannitol (10 g, 55 mmol) in acetone (250 mL), and the reaction mixture was stirred for 48 h at ambient temperature. Then, the mixture was neutralized with a 25% aqueous solution of NH₄OH (4.7 mL) and Na₂CO₃ (6.2 g). The solvent was removed in vacuo, and the residue was recrystallized from ethanol to give 21 as a colorless solid (5.3 g, 18 mmol, 32%). mp 70 °C; $[\alpha]_D^{20} =$ +16.3 (6.7, methanol); ¹H NMR (DMSO- d_6): δ [ppm] = 1.27 (s, 6H, $CH_2OC(CH_3)_2$, 1.32 (s, 6H, $(CHO)_2OC(CH_3)_2$), 1.33 (s, 6H, $CH_2OC(CH_3)_2$, 3.83 (dd, J = 8.3/5.7 Hz, 2H, OCH_2CHCH), 3.85– 3.90 (m, 2H, HOCH₂CHCH), 4.03 (dd, $J = \tilde{8.3}/6.6$ Hz, 2H, OCH₂CHCH), and 4.11-4.18 (m, 2H, OCH₂CHCH); ¹³C NMR $(DMSO-d_6): \delta [ppm] = 25.2 (2C, CH_2OC(CH_3)_2), 26.3 (2C, CH_2OC(CH_3)_2))$ $CH_2OC(CH_3)_2$), 27.2 (2C, (CHO)₂C(CH₃)₂), 65.4 (2C, OCH₂CHCH), 75.7 (2C, OCH₂CHCH), 78.6 (2C, OCH₂CHCH), 108.7 (2C, CH₂OC(CH₃)₂), and 109.4 (1C, (CHO)₂C(CH₃)₂); IR (neat): $\tilde{\nu}$ [cm⁻¹] = 2991, 2958, 2936, 2880, 1368, 1257, 1211, 1147, 1063, 969, 844, 787, 509, and 408; HRMS (m/z): $[M + Na]^+$ calcd for C15H26NaO6, 325.1622; found, 325.1613.

(1R,1'R)-1,1'-[(4R,5R)-2,2-Dimethyl-1,3-dioxolane-4,5-diyl]bis-(ethane-1,2-diol) (22). The compound was synthesized according to the literature with minor variations:^{41,42} After stirring 21 (1.0 g, 3.3 mmol) in 70% aqueous acetic acid (20 mL) for 90 min at 40 °C, the solvent was removed in vacuo and the residue was suspended in acetone (30 mL). The suspension was sonicated for 2 min, filtered, and the solvent was removed in vacuo. The residue was purified by flash column chromatography ($\emptyset = 3 \text{ cm}, h = 20 \text{ cm}, V = 20 \text{ mL},$ dichloromethane/methanol = 20:1, $R_f = 0.13$) to give 22 as a colorless solid (510 mg, 2.3 mmol, 69%). mp 86 °C; $[\alpha]_D^{20} = +27.0$ (5.7, methanol); ¹H NMR (DMSO- d_6): δ [ppm] = 1.28 (s, 6H, C(CH₃)₂), 3.32-3.39 (m, 2H, HOCH2CHCH), 3.44-3.51 (m, 2H, HOCH₂CHCH), 3.54 (ddd, J = 11.1/5.6/3.2 Hz, 2H, HOCH₂CHCH), 3.84-3.88 (m, 2H, HOCH₂CHCH), 4.44 (t, J = 5.7 Hz, 2H, CH₂OH), and 5.06 (d, J = 4.5 Hz, 2H, CHOH); ¹³C NMR (DMSO- d_6): δ [ppm] = 27.2 (2C, C(CH_3)_2), 63.0 (2C, HOCH₂CHCH), 72.9 (2C, HOCH₂CHCH), 79.1 (2C, HOCH₂CHCH), and 108.3 (1C, $C(CH_3)_2$); IR (neat): $\tilde{\nu}$ [cm⁻¹] = 3312, 2936, 2884, 1433, 1371, 1335, 1221, 1192, 1166, 1116, 1068, 1035, 1011, 978, 875, 712, and 505; HRMS (*m*/*z*): [M + Na]⁺ calcd for C₉H₁₈NaO₆, 245.0996; found, 245.0996.

(1R,1'R)-1,1'-[(4R,5R)-2,2-Dimethyl-1,3-dioxolane-4,5-diyl]bis[2-(benzyloxy)ethan-1-ol] (23). A mixture of 22 (400 mg, 1.8 mmol)

and dibutyltin oxide (970 mg, 3.9 mmol) was heated to reflux in toluene (20 mL) using a Dean–Stark trap for 16 h. Then, the solvent was removed in vacuo, and the residue was suspended in toluene (10 mL). Benzyl bromide (0.85 mL, 1.2 g, 7.2 mmol) and tetrabutylammonium iodide (660 mg, 1.8 mmol) were added and the mixture was heated to 70 °C for 36 h. Then, the solvent was removed in vacuo and the residue was purified by flash column chromatography (\emptyset = 6 cm, h = 18 cm, V = 50 mL, petroleum ether/ ethyl acetate = 2:1, R_f = 0.29) to give 23 as a colorless oil (670 mg, 1.7 mmol, 92%). $[\alpha]_D^{20} = +26.4$ (4.9, methanol); ¹H NMR (DMSO- d_6): δ [ppm] = 1.28 (s, 6H, C(CH₃)₂), 3.41 (dd, J = 10.1/6.5 Hz, 2H, OCH_2CHCH), 3.59 (dd, J = 10.1/3.2 Hz, 2H, OCH_2CHCH), 3.66– 3.75 (m, 2H, OCH₂CHCH), 3.89–3.94 (m, 2H, OCH₂CHCH), 4.48 (d, J = 12.3 Hz, 2H, OCH₂Ph), 4.52 (d, J = 12.3 Hz, 2H, OCH₂Ph), 5.24 (d, J = 5.0 Hz, 2H, CHOH), and 7.24-7.37 (m, 10H, 2'-H_{phenyb} 3'- H_{phenyb} 4'- H_{phenyb} 5'- H_{phenyb} 6'- H_{phenyb}); ¹³C NMR (DMSO- d_6): δ [ppm] = 27.2 (2C, C(CH_3)_2), 70.9 (2C, OCH_2CHCH), 71.9 (2C, OCH₂CHCH), 72.3 (2C, OCH₂Ph), 79.3 (2C, OCH₂CHCH), 108.5 (1C, C(CH₃)₂), 127.3 (2C, C-4'_{phenyl}), 127.4 (4C, C-2'_{phenyl}, C-6' $_{\rm phenyl}$), 128.2 (4C, C-3' $_{\rm phenyl}$, C-5' $_{\rm phenyl}$), and 138.6 (2C, C-1' $_{\rm phenyl}$); IR (neat): $\tilde{\nu}$ [cm^{-1}] = 3363, 2985, 2912, 2866, 1496, 1453, 1370, 1239, 1211, 1166, 1066, 1027, 907, 872, 734, 696, 595, 509, and 466; HRMS (m/z): $[M + Na]^+$ calcd for $C_{23}H_{30}NaO_{6}$, 425.1935; found, 425.1929; HPLC (method 1): $t_{\rm R}$ = 22.9 min, purity 96.8%.

(4R,5R)-4,5-Bis{(R)-2-(benzyloxy)-1-[(4-iodobenzyl)oxy]ethyl}-2,2-dimethyl-1,3-dioxolane (24). Under a N2 atmosphere and icecooling, sodium hydride (60% suspension in paraffin oil, 390 mg, 9.7 mmol) was added to a solution of 23 (620 mg, 1.5 mmol) in THF (15 mL). After stirring the mixture for 10 min at 0 °C, stirring was continued for 1 h at ambient temperature. Then, 4-iodobenzyl bromide (1.4 g, 4.6 mmol) was added, and the reaction mixture was stirred for 72 h at ambient temperature. Afterward, methanol and water were added, and the mixture was extracted with ethyl acetate $(3\times)$. The combined organic layers were dried (Na_2SO_4) , filtered, and the solvent was removed in vacuo. The residue was purified by flash column chromatography ($\emptyset = 6 \text{ cm}, h = 17 \text{ cm}, V = 50 \text{ mL},$ petroleum ether/ethyl acetate = 10:1, $R_f = 0.27$) to give 24 as a colorless oil (1.1 g, 1.3 mmol, 85%). $[\alpha]_D^{20} = +18.2$ (10.4, acetonitrile); ¹H NMR (DMSO- d_6): δ [ppm] = 1.28 (s, 6H, $C(CH_3)_2$, 3.51 (dd, J = 10.5/5.7 Hz, 2H, OCH₂CHCH), 3.63-3.74 (m, 4H, OCH₂CHCH, OCH₂CHCH (2H)), 4.08-4.14 (m, 2H, OCH₂CHCH), 4.43 (s, 4H, CH₂OCH₂Ph), 4.48 (d, J = 12.1 Hz, 2H, CHOCH₂Ar), 4.58 (d, J = 12.1 Hz, 2H, CHOCH₂Ar), 7.04–7.10 (m, 4H, 2"-H_{4-iodophenyl}, 6"-H_{4-iodophenyl}), 7.24–7.35 (m, 10H, 2'-H_{phenyl}, 3'-H_{phenyl}, 4'-H_{phenyl}, 5'-H_{phenyl}), and 7.59–7.65 (m, 4H, 3"-H_{4-iodophenyl}), ¹³C NMR (DMSO- d_6): δ [ppm] = 27.1 (2C, C(CH₃)₂), 69.6 (2C, OCH₂CHCH), 70.9 (2C, CHOCH₂Ar), 72.3 (2C, CH₂OCH₂Ph), 77.6 (2C, OCH₂CHCH), 79.0 (2C, 1209, 1073, 1006, 872, 798, 733, 696, 610, and 470; HRMS (m/z): $[M + Na]^+$ calcd for $C_{37}H_{40}I_2NaO_6$, 857.0806; found, 857.0780; HPLC (method 3): $t_{\rm R} = 32.1$ min, purity 95.6%.

(2R,3S,4S,5R)-1,6-Bis(benzyloxy)-2,5-bis[(4-iodobenzyl)oxy]hexane-3,4-diol (25). After stirring 24 (870 mg, 1.0 mmol) in 80% aqueous trifluoroacetic acid (5 mL) for 2 h at 0 °C, the reaction mixture was diluted with toluene, and the solvent was removed in vacuo. The residue was purified by flash column chromatography (\emptyset = 3 cm, *h* = 17 cm, *V* = 20 mL, petroleum ether/ethyl acetate = 2:1, *R_f* = 0.31) to give 25 as a colorless solid (800 mg, 1.0 mmol, 97%). mp 80 °C; [α]_D²⁰ = +23.4 (5.9, methanol); ¹H NMR (DMSO-*d*₆): δ [ppm] = 3.59–3.68 (m, 4H, OCH₂CHCHOH, OCH₂CHCHOH (2H)), 3.71–3.79 (m, 2H, OCH₂CHCHOH), 3.82–3.90 (m, 2H, OCH₂CHCHOH), 4.50 (d, *J* = 12.0 Hz, 2H, CHOCH₂Ar), 4.51 (s, 4H, CH₂OCH₂Ph), 4.53–4.61 (m, 2H, OCH₂CHCHOH), 4.64 (d, *J* = 12.0 Hz, 2H, CHOCH₂Ar), 7.08–7.15 (m, 4H, 2″-H_{4-iodophenyl}, δ' -H_{4-iodophenyl}), 7.24–7.36 (m, 10H, 2'-H_{phenyl}, 3'-H_{phenyl}, 4'-H_{phenyl}, 5'- H_{phenyl} 6'-H_{phenyl}), and 7.59–7.66 (m, 4H, 3"-H_{4-iodophenyl}, 5"-H_{4-iodophenyl}); ¹³C NMR (DMSO-*d*₆): δ [ppm] = 68.1 (2C, OCH₂CHCHOH), 70.6 (2C, OCH₂CHCHOH), 71.0 (2C, CHO-CH₂Ar), 72.4 (2C, CH₂OCH₂Ph), 78.6 (2C, OCH₂CHCHOH), 93.0 (2C, C-4"_{4-iodophenyl}), 127.28 (2C, C-4'_{phenyl}), 127.31 (4C, C-2'_{phenyl}) C-6'_{phenyl}), 128.2 (4C, C-3'_{phenyl}) C-5'_{phenyl}), 129.6 (4C, C-2"_{4-iodophenyl}), 138.7 (2C, C-1'_{phenyl}), 136.8 (4C, C-3"_{4-iodophenyl}), 138.7 (2C, C-1'_{phenyl}), and 138.9 (2C, C-1"_{4-iodophenyl}); IR (neat): $\tilde{\nu}$ [cm⁻¹] = 3420, 3261, 2929, 2866, 1484, 1453, 1401, 1373, 1344, 1318, 1298, 1215, 1100, 1077, 1056, 1006, 875, 794, 748, 732, 693, 476, and 456; HRMS (*m*/*z*): [M + Na]⁺ calcd for C₃₄H₃₆I₂NaO₆, 817.0493; found, 817.0495; HPLC (method 3): *t*_R = 29.4 min, purity 96.4%.

Methyl (S)-3-(benzyloxy)-2-[(4-iodobenzyl)oxy]propanoate ((S)-18). An oxidant solution was prepared by dissolving H_5IO_6 (11.4 g, 50 mmol) and CrO₃ (23 mg, 0.23 mmol) in wet acetonitrile (114 mL, 0.75% water V/V) overnight. Under ice-cooling, the oxidant solution (8 mL) was added to a solution of 17 (590 mg, 1.5 mmol) in acetonitrile (10 mL). After stirring the reaction mixture for 30 min at 0 °C, stirring was continued for 16 h at ambient temperature. Then, the solvent was concentrated in vacuo, water was added, and the mixture was extracted with diethyl ether $(3\times)$. The combined organic layers were dried (Na₂SO₄), filtered, and the solvent was removed in vacuo. The residue was dissolved in methanol (15 mL) and concentrated sulfuric acid (0.1 mL) was added. After heating the reaction mixture to reflux for 16 h, the solvent was concentrated in vacuo. Then, the mixture was diluted with ethyl acetate and washed with ice-cold water, a saturated aqueous solution of NaHCO₃, and brine. The combined organic layers were dried (Na_2SO_4) , filtered, and the solvent was removed in vacuo. The residue was purified by flash column chromatography ($\emptyset = 4 \text{ cm}, h = 24 \text{ cm}, V = 30 \text{ mL},$ petroleum ether/ethyl acetate = 6:1, $R_f = 0.31$) to give (S)-18 as a colorless oil (300 mg, 0.71 mmol, 48%). $[\alpha]_D^{20} = -25.6$ (1.3, methanol); HPLC (method 1): $t_{\rm R}$ = 26.0 min, purity 94.5%.

Methyl (R)-3-(benzyloxy)-2-[(4-iodobenzyl)oxy]propanoate ((R)-18). NaIO₄ (320 mg, 1.5 mmol) was added to a solution of 25 (770 mg, 0.97 mmol) in methanol (40 mL), and the mixture was stirred at ambient temperature for 48 h. Then, the solvent was concentrated in vacuo, brine was added, and the mixture was extracted with ethyl acetate (3×). The combined organic layers were dried (Na_2SO_4) , filtered, and the solvent was removed in vacuo. The residue was dissolved in a mixture of methanol and water (9:1, 40 mL) and NaHCO₃ (2.5 g, 29 mmol) was added. Then, Br₂ (0.15 mL, 0.46 g, 2.9 mmol) was added, and the reaction mixture was stirred at ambient temperature for 48 h in a flask protected from ordinary lighting. Afterward, sodium thiosulfate and water were added and the mixture was extracted with ethyl acetate $(3\times)$. The combined organic layers were dried (Na_2SO_4) , filtered, and the solvent was removed in vacuo. The residue was purified by flash column chromatography ($\emptyset = 6$ cm, h = 19 cm, V = 50 mL, petroleum ether/ethyl acetate = 4:1, $R_f = 0.51$) to give (R)-18 as a colorless oil (570 mg, 1.3 mmol, 69%). $[\alpha]_{\rm D}^{20} =$ +29.8 (8.4, methanol); HPLC (method 1): $t_{\rm R} = 26.1$ min, purity 97.8%.

Spectroscopic Data of (S)-18 and (R)-18. ¹H NMR (DMSO-*d*₆): δ [ppm] = 3.67 (s, 3H, CO₂CH₃), 3.71 (dd, J = 10.7/3.8 Hz, 1H, OCHCH₂O), 3.73 (dd, J = 10.7/5.1 Hz, 1H, OCHCH₂O), 4.29 (dd, J = 5.0/4.0 Hz, 1H, OCHCH₂O), 4.41 (d, J = 12.1 Hz, 1H, CHOCH₂Ar), 4.48 (d, J = 12.2 Hz, 1H, CH₂OCH₂Ph), 4.53 (d, J = 12.2 Hz, 1H, CH₂OCH₂Ph), 4.60 (d, J = 12.1 Hz, 1H, CHOCH₂Ar), 7.15–7.18 (m, 2H, 2"-H_{4-iodophenyl}, 6"-H_{4-iodophenyl}), 7.26–7.32 (m, 3H, 2'-H_{phenyl}, 4'-H_{phenyl}, 6'-H_{phenyl}), 7.32–7.37 (m, 2H, 3'-H_{phenyl}, s'-H_{phenyl}), and 7.69–7.73 (m, 2H, 3"-H_{4-iodophenyl}), 5"-H_{4-iodophenyl}); ¹³C NMR (DMSO-*d*₆): δ [ppm] = 51.8 (1C, CO₂CH₃), 70.1 (1C, OCHCH₂O), 70.8 (1C, CHOCH₂Ar), 72.3 (1C, CH₂OCH₂Ph), 77.7 (1C, OCHCH₂O), 93.6 (1C, C-4"_{4-iodophenyl}), 127.47 (2C, C-2'_{phenyl}), 129.8 (2C, C-2"_{4-iodophenyl}), C-6"_{4-iodophenyl}), 137.0 (2C, C-3"_{4-iodophenyl}), 137.5 (1C, CO₂CH₃); IR (neat): $\tilde{\nu}$ [cm⁻¹] = 3070, 2930, 2857, 1472, 1427, 1390, 1361, 1105, 1028, 857, 738, 697, 611, 503, and 487; HRMS (m/z): $[M + H]^+$ calcd for $C_{18}H_{19}INaO_4$, 449.0220; found, 449.0219.

Methyl (S)-3-(benzyloxy)-2-[(4-{[4-(morpholinomethyl)phenyl]ethynyl}benzyl)oxy]propanoate ((S)-19). Under a N₂ atmosphere, copper(I) iodide (11 mg, 0.058 mmol), bis(triphenylphosphine)palladium(II) chloride (38 mg, 0.054 mmol), and diisopropylamine (3 mL) were added to a solution of (S)-18 (200 mg, 0.48 mmol) in dry THF (10 mL) at ambient temperature, and the mixture was stirred for 20 min. Then, 4-(4-ethynylbenzyl)morpholine (220 mg, 1.1 mmol) was added in two portions at an interval of 30 min. After stirring the reaction mixture for 24 h at ambient temperature, the solvent was removed in vacuo. The residue was dissolved in a mixture of petroleum ether and ethyl acetate (1:1) and filtered through a short silica gel column. The solvent was removed in vacuo, and the residue was purified by flash column chromatography ($\emptyset = 4 \text{ cm}, h = 11 \text{ cm},$ V = 20 mL, petroleum ether/ethyl acetate = 1:1, $R_f = 0.17$) to give (S)-19 as a yellow oil (210 mg, 0.42 mmol, 88%). $[\alpha]_{\rm D}^{20} = -21.3$ (1.2, methanol); HPLC (method 1): $t_{\rm R} = 21.7$ min, purity 93.2%.

Methyl (R)-3-(benzyloxy)-2-[(4-{[4-(morpholinomethyl)phenyl]ethynyl]benzyl)oxy]propanoate ((R)-19). Under a N₂ atmosphere, copper(I) iodide (52 mg, 0.27 mmol), bis(triphenylphosphine)palladium(II) chloride (140 mg, 0.20 mmol), and diisopropylamine (10 mL) were added to a solution of (R)-18 (430 mg, 1.0 mmol) in dry THF (5 mL) at ambient temperature, and the mixture was stirred for 10 min. Then, 4-(4-ethynylbenzyl)morpholine (300 mg, 1.5 mmol) was added, and the reaction mixture was heated to reflux for 24 h. After stirring the reaction mixture for an additional 12 h at ambient temperature, the solvent was removed in vacuo. The residue was dissolved in a mixture of petroleum ether and ethyl acetate (1:1) and filtered through a short silica gel column. The solvent was removed in vacuo, and the residue was purified by flash column chromatography (\emptyset = 4 cm, h = 10 cm, V = 30 mL, petroleum ether/ ethyl acetate = 1:1, $R_f = 0.17$) to give (R)-19 as a yellow oil (320 mg, 0.65 mmol, 64%). $[\alpha]_D^{20} = +23.9$ (8.0, methanol); HPLC (method 1): $t_{\rm R} = 21.8$ min, purity 95.5%.

Spectroscopic Data of (S)-19 and (R)-19. ¹H NMR (DMSO-d₆): δ [ppm] = 2.31-2.40 (m, 4H, N(CH₂CH₂)₂O), 3.49 (s, 2H, NCH₂Ar), 3.55-3.60 (m, 4H, N(CH₂CH₂)₂O), 3.68 (s, 3H, CO_2CH_3 , 3.73 (dd, J = 10.7/4.0 Hz, 1H, OCHCH₂O), 3.75 (dd, J= 10.7/5.1 Hz, 1H, OCHCH₂O), 4.32 (dd, J = 5.0/4.1 Hz, 1H, OCHCH₂O), 4.49 (d, J = 12.2 Hz, 1H, CH₂OCH₂Ph), 4.52-4.56 (m, 2H, CHOCH₂Ar (1H), CH₂OCH₂Ph (1H)), 4.68 (d, J = 12.2Hz, 1H, CHOCH₂Ar), 7.27–7.31 (m, 3H, 2'-H_{phenyl}, 4'-H_{phenyl}, 6'-H_{phenyl}), 7.33–7.38 (m, 4H, 3'-H_{phenyl}, 5'-H_{phenyl}, 3'''-H₄(morpholinomethyl)phenyl), 7.38–7.42 (m, 4H, 3H) 2H, 2"-H_{4-{[4-(morpholinomethyl)phenyl]ethynyl}phenyl, 6"-} $H_{4-\{[4-(morpholinomethyl)phenyl]ethynyl\}phenyl})$, and 7.49–7.54 (m, 4H, 3"- $\begin{array}{l} H_{4-\{[4-(m \ or \ p \ holinomethyl]phenyl]ethyl]phenyl phenyl] eth y nyl \} phenyl], 5''-H_{4-\{[4-(m \ or \ p \ holinomethyl]phenyl]ethyl]phenyl 2'''-H_{4-(m \ or \ p \ holinomethyl]phenyl], 5'''-H_{4-(m \ or \ p \ holinomethyl]phenyl), 1''C NMR (DMSO-d_{6}): \delta [ppm] = 51.8 (1C, CO_{2}CH_{3}), 53.2 (2C, N(CH_{2}CH_{2})_{2}O), 62.0 (1C, ArCH_{2}N), (1C, ArCH_{2}N), (1C, CO_{2}CH_{3}), 53.2 (2C, N(CH_{2}CH_{2})_{2}O), 62.0 (1C, ArCH_{2}N), (1C, ArCH_{2}N),$ 66.2 (2C, N(CH₂CH₂)₂O), 70.1 (1C, OCHCH₂O), 71.1 (1C, CHOCH₂Ar), 72.3 (1C, CH₂OCH₂Ph), 77.8 (1C, OCHCH₂O), 89.0 $(1C, C \equiv C), 89.3 (1C, C \equiv C), 120.8 (1C, C = C)$ $\begin{array}{c} 1 & "' & 4 - (m \ or \ p \ ho \ li \ n \ om \ e \ th \ y \) \ p \ hen \ y \), \ 1 \ 2 \ 1 \ . \ 4 \ (1 \ C, \ C \ - 4'' \ 4 + [[4 - (m \ or \ p \ ho \]) \ phen \ y \]), \ 1 \ 2 \ 7 \ . \ 4 \ (2 \ C, \ C \ - 2'' \ phen \ y \]), \ 1 \ 2 \ 7 \ . \ 4 \ (2 \ C, \ C \ - 2'' \ phen \ y \]), \ 1 \ 2 \ 7 \ . \ 5 \ (2 \ C, \ C \ - 2'' \ phen \ y \]), \ 1 \ 2 \ 7 \ . \ 5 \ (2 \ C, \ C \ - 2'' \ phen \ y \]), \ 1 \ 2 \ 7 \ . \ 5 \ (2 \ C, \ C \ - 2'' \ phen \ y \]), \ 1 \ 2 \ 7 \ . \ 5 \ (2 \ C, \ C \ - 2'' \ phen \ y \]), \ 1 \ 2 \ 7 \ . \ 5 \ (2 \ C, \ C \ - 2'' \ phen \ y \]), \ 1 \ 2 \ 7 \ . \ 5$ $2''_{4-\{[4-(morpholinomethyl)phenyl], 12/.8}$ (2C, C- $2''_{4-\{[4-(morpholinomethyl]phenyl]phenyl], 128.3}$ (2C, C-3'phenyl), C- $5''_{phenyl}$), 129.2 (2C, C-3'''_{4-(morpholinomethyl)phenyl), C- $5'''_{4-((morpholinomethyl)phenyl), 121.2}$ (4C) 4-(morpholinomethyl)phenyl), 131.2 (4C, $\begin{array}{c} 5 & 4 - (morpholinomethyl)phenyl), & 151.2 & (TC, C) \\ 3'' & 4 - \{[4 - (morpholinomethyl)phenyl]ethynyl]ethynyl]ethynyl}phenyl, \\ C - 5'' & 4 - ([4 - (morpholinomethyl)phenyl]ethynyl]phenyl), \\ C - 2''' & (-(morpholinomethyl)phenyl), \\ 138.0 & (1C, C - 2''' & (-(morpholinomethyl)phenyl), \\ C - 3'' & 4 - ([4 - (morpholinomethyl)phenyl]ethynyl]phenyl), \\ 138.9 & (1C, C - 2''' & (-(morpholinomethyl)phenyl), \\ C - 4''' & (-(morpholinomethyl)phenyl), \\ C - 2''' & (-(morpholinomethyl)phenyl), \\ C - 2'''' & (-(morpholinomethyl)phenyl), \\ C - 2''' & (-(morpholinomethyl)$ 1204, 1114, 1069, 1007, 913, 865, 819, 794, 736, 697, 540, and 515; HRMS (m/z): $[M + H]^+$ calcd for C₃₁H₃₄NO₅, 500.2431; found, 500.2449.

(S)-3-(Benzyloxy)-N-hydroxy-2-[(4-{[4-(morpholinomethyl)phenyl]ethynyl]benzyl)oxy]propanamide ((S)-11a). Under ice-cooling, an aqueous solution of hydroxylamine (50 wt %, 2.5 mL) was added to a solution of (S)-19 (120 mg, 0.24 mmol) in a mixture of THF (4 mL) and isopropanol (4 mL). After stirring the reaction mixture for 5 min at 0 °C, stirring was continued for 36 h at ambient temperature. Then, the solvent was removed in vacuo, and the residue was purified by automatic flash column chromatography using a Biotage Isolera One system (10% \rightarrow 80% ACN in H₂O, Biotage SNAP Ultra C18 30 g). Fractions containing the desired product were combined and subjected to lyophilization to give (S)-11a as a colorless solid (80 mg, 0.16 mmol, 67%). mp 65 °C; $[\alpha]_D^{20} = -22.5$ (1.5, methanol); HPLC (method 2): $t_R = 13.7$ min, purity 100%.

(1.5, methanol); HPLC (method 2): $t_R = 13.7$ min, purity 100%. (R)-3-(Benzyloxy)-N-hydroxy-2-[(4-{[4-(morpholinomethyl)phenyl]ethynyl]benzyl)oxy]propanamide ((R)-11a). Under ice-cooling, an aqueous solution of hydroxylamine (50 wt %, 6 mL) was added to a solution of (R)-19 (240 mg, 0.48 mmol) in a mixture of THF (5 mL) and isopropanol (5 mL). After stirring the reaction mixture for 5 min at 0 °C, stirring was continued for 72 h at ambient temperature. Then, the solvent was removed in vacuo, water was added, and the mixture was extracted with ethyl acetate $(3\times)$. The combined organic layers were dried (Na2SO4), filtered, and the solvent was removed in vacuo. The residue was purified by automatic flash column chromatography using a Biotage Isolera One system $(10\% \rightarrow 100\%$ ACN in H₂O, Biotage SNAP Ultra C18 12 g). Fractions containing the desired product were combined and subjected to lyophilization to give (R)-11a as a colorless solid (160 mg, 0.32 mmol, 65%). mp 65 °C; $[\alpha]_{\rm D}^{20} = +22.6$ (1.9, methanol); HPLC (method 2): $t_{\rm R} = 13.7$ min, purity 97.1%.

Spectroscopic Data of (S)-11a and (R)-11a. ¹H NMR (DMSO d_6 : δ [ppm] = 2.30-2.40 (m, 4H, N(CH₂CH₂)₂O), 3.49 (s, 2H, NCH₂Ar), 3.55–3.61 (m, 4H, N(CH₂CH₂)₂O), 3.65 (d, J = 5.3 Hz, 2H, OCHCH₂O), 4.01 (t, J = 5.3 Hz, 1H, OCHCH₂O), 4.46-4.53 (m, 3H, CHOC H_2 Ar (1H), CH₂OC H_2 Ph), 4.62 (d, J = 12.6 Hz, 1H, CHOCH₂Ar), 7.27–7.33 (m, 3H, 2'-H_{phenyl}, 4'-H_{phenyl}, 6'-H_{phenyl}), 7.33–7.38 (m, 4H, 3'-H_{phenyl}, 5'-H_{phenyl}, 3^{'''}-H₄(morpholinomethyl)phenyl 5^{'''}-H₄(morpholinomethyl)phenyl), 7.39–7.43 (m, 2H, 2^{''}- $\begin{array}{l} H_{4-\{[4-(morpholinomethyl)phenyl]ethyn)phenyl} \\ H_{4-\{[4-(morpholinomethyl)phenyl]ethynylphenyl} \\ 2'''-H_{4-(morpholinomethyl)phenyl], \\ 8.96 (s br, 1H, CONHOH), and 10.77 (s br, 1H, CONHOH), \\ 13C NMR (DMSO-d_6): \\ \delta [ppm] = 53.2 (2C, CONHOH), \\ 10C CONHOH (context) \\ 10C CONHOH (context), \\ 10C CONHOH (c$ N(CH₂CH₂)₂O), 62.0 (1C, ArCH₂N), 66.2 (2C, N(CH₂CH₂)₂O), 70.3 (1C, OCHCH₂O), 70.7 (1C, CHOCH₂Ar), 72.2 (1C, CH₂OCH₂Ph), 77.7 (1C, OCHCH₂O), 89.1 (1C, C≡C), 89.3 $(1C, C \equiv C)$, 120.8 (1C, C-1^{*m*}₄(morpholinomethyl)phenyl</sub>), 121.4 (1C, C-4^{*m*}₄([4(morpholinomethyl)phenyl]), 127.45 (1C, C-4^{*i*}_{phenyl}), 127.50 (2C, C-2^{*i*}_{phenyl}), C-6^{*i*}_{phenyl}), 127.8 (2C, C-2^{*m*}_{phenyl}), C-6^{*i*}_{phenyl}), 127.8 (2C, C-2^{*m*}_{phenyl}), C-6^{*i*}_{phenyl}), 127.8 (2C, C-2^{*m*}_{phenyl}), C-6^{*i*}_{phenyl}), 127.8 (2C, C-2^{*m*}_{phenyl}), C-6^{*m*}_{phenyl}), C-6^{*m*}_{phenyl} 1"_{4-{[4-(morpholinomethyl)phenyl]}, 138.8 (1C, C-4""_{4-(morpholinomethyl)phenyl)}, and 165.7 (1C, CONHOH); IR (neat): $\tilde{\nu}$ $[cm^{-1}] = 3209, 3030, 2857, 2810, 1664, 1517, 1453, 1350, 1113,$ 1006, 914, 864, 820, 792, 736, 697, 539, and 514; HRMS (*m*/*z*): [M + H]⁺ calcd for $C_{30}H_{33}N_2O_5$, 501.2384; found, 501.2395.

(5)-1-Azido-3-[(tert-butyldiphenylsilyl)oxy]propan-2-ol (26). Sodium azide (2.5 g, 38 mmol) was added to a solution of 15 (2.4 g, 7.6 mmol) and ammonium chloride (890 mg, 17 mmol) in a mixture of methanol (120 mL) and water (15 mL). After stirring the reaction mixture at 65 °C for 16 h, the mixture was diluted with diethyl ether (120 mL), a saturated aqueous solution of NaHCO₃ (100 mL) was added, and the mixture was extracted with diethyl ether (3×). The combined organic layers were washed with brine, dried (Na₂SO₄), filtered, and the solvent was removed in vacuo. The residue was purified by automatic flash column chromatography using an Interchim puriFlash XS 420 system (0% \rightarrow 10% ethyl acetate in petroleum ether, Biotage SNAP Ultra HP-Sphere 50 g) to give **26** as a colorless oil (1.9 g, 5.3 mmol, 70%). $R_f = 0.36$ (petroleum ether/ethyl acetate = 10:1); $[\alpha]_D^{20} = -15.6$ (2.3, methanol); ¹H NMR (DMSO- d_6): δ [ppm] = 0.99 (s, 9H, SiC(CH₃)₃), 3.32 (dd, J = 12.6/6.3 Hz, 1H, N₃CH₂CHCH₂), 3.41 (dd, J = 12.6/3.6 Hz, 1H, N₃CH₂CHCH₂), 3.55 (dd, J = 10.0/7.0 Hz, 1H, N₃CH₂CHCH₂), 3.55 (dd, J = 10.0/7.0 Hz, 1H, N₃CH₂CHCH₂), 3.60 (dd, J = 10.0/5.0 Hz, 1H, N₃CH₂CHCH₂), 3.73–3.85 (m, 1H, N₃CH₂CHCH₂), 5.27 (d, J = 5.2 Hz, 1H, CHOH), 7.39–7.51 (m, 6H, 3'-H_{diphenylsilyl}, 4'-H_{diphenylsilyl});

¹³C NMR (DMSO- d_6): δ [ppm] = 18.8 (1C, SiC(CH₃)₃), 26.6 (3C, SiC(CH₃)₃), 53.2 (1C, N₃CH₂CHCH₂), 65.1 (1C, N₃CH₂CHCH₂), 70.2 (1C, N₃CH₂CHCH₂), 127.9 (4C, C-3' diphenylsilyl), 129.9 (2C, C-4' diphenylsilyl), 132.8 (1C, C-1' diphenylsilyl), 132.9 (1C, C-1' diphenylsilyl), 135.0 (2C, C-2' diphenylsilyl), C-6' diphenylsilyl), and 135.1 (2C, C-2' diphenylsilyl), C-6' diphenylsilyl); IR (neat): $\tilde{\nu}$ [cm⁻¹] = 3430, 3071, 2930, 2858, 2098, 1472, 1427, 1288, 1106, 998, 937, 823, 800, 740, 699, 613, 503, and 486; HRMS (m/z): [M + Na]⁺ calcd for C₁₉H₂₅N₃NaO₂Si, 378.1608; found, 378.1574; HPLC (method 1): $t_{\rm R}$ = 27.6 min, purity 99.8%.

(S)-3-Azido-2-[(4-iodobenzyl)oxy]propan-1-ol (27). Under a N₂ atmosphere, sodium hydride (60% suspension in paraffin oil, 970 mg, 24 mmol) was added to an ice-cooled solution of 26 (1.8 g, 5.1 mmol) in anhydrous THF (75 mL). After stirring the reaction mixture for 15 min at 0 °C, 4-iodobenzyl bromide (1.9 g, 6.5 mmol) was added, and the reaction mixture was stirred for additional 15 min at 0 °C. Then, the ice-bath was removed, and the mixture was stirred for 72 h at ambient temperature. Afterward, methanol was added under ice-cooling, the solvent was removed in vacuo, water was added, and the mixture was extracted with ethyl acetate $(3\times)$. The combined organic layers were dried (Na₂SO₄), filtered, and the solvent was removed in vacuo. The residue was dissolved in THF (80 mL), tetrabutylammonium fluoride trihydrate (3.5 g, 11 mmol) was added, and the reaction mixture was stirred for 36 h at ambient temperature. Then, the solvent was removed in vacuo, diethyl ether was added, and the mixture was washed with a saturated aqueous solution of ammonium chloride, water, and brine. The organic layer was dried (Na_2SO_4) , filtered, and the solvent was removed in vacuo. The residue was purified by automatic flash column chromatography using an Interchim puriFlash XS 420 system (20% \rightarrow 33% ethyl acetate in petroleum ether, Biotage SNAP Ultra HP-Sphere 50 g) to give 27 as a yellow oil (1.3 g, 3.9 mmol, 78%). $R_f = 0.38$ (petroleum ether/ethyl acetate = 2:1); $[\alpha]_{D}^{20} = +7.8$ (2.6, methanol); ¹H NMR $(DMSO-d_6)$: δ [ppm] = 3.33 (dd, J = 13.1/6.1 Hz, 1H, N₃CH₂CHCH₂OH), 3.41-3.61 (m, 4H, N₃CH₂CHCH₂OH (1H), N₃CH₂CHCH₂OH, N₃CH₂CHCH₂OH), 4.55 (d, J = 12.2 Hz, 1H, OCH_2Ar), 4.61 (d, J = 12.2 Hz, 1H, OCH_2Ar), 4.81 (t, J = 5.5 Hz, 1H, CH₂OH), 7.14–7.21 (m, 2H, 2'-H_{4-iodophenyl}, 6'-H_{4-iodophenyl}), and 7.67–7.74 (m, 2H, 3'-H_{4-iodophenyl}, 5'-H_{4-iodophenyl}); ¹³C NMR (DMSO- d_6): δ [ppm] = 50.9 (1C, N₃CH₂CHCH₂OH), 60.3 (1C, $N_3CH_2CHCH_2OH)$, 70.1 (1C, OCH_2Ar), 78.9 (1C, $N_3CH_2CHCH_2OH)$, 93.3 (1C, C-4'_{4-iodophenyl}), 129.7 (2C, C- $2'_{4-iodophenyl}$, C- $6'_{4-iodophenyl}$), 136.9 (2C, C- $3'_{4-iodophenyl}$, C- $5'_{4-iodophenyl}$), and 138.4 (1C, C- $1'_{4-iodophenyl}$); IR (neat): $\tilde{\nu}$ [cm⁻¹] = 3399, 2927, 2871, 2093, 1590, 1484, 1402, 1344, 1273, 1099, 1057, 1006, 798, 627, 554, and 470; HRMS (m/z): $[M + Na]^+$ calcd for C10H12IN3NaO2, 355.9866; found, 355.9875; HPLC (method 1): t_R = 21.7 min, purity 99.9%.

(2R,2'R)-[(4R,5R)-2,2-Dimethyl-1,3-dioxolane-4,5-diyl]bis(2-hydroxyethane-2,1-diyl) bis(4-methylbenzenesulfonate) (29). A mixture of 22 (1.2 g, 5.3 mmol) and dibutyltin oxide (2.8 g, 11 mmol) was heated to reflux in toluene (250 mL) using a Dean–Stark trap for 48 h. Then, the solvent was removed in vacuo and the residue was suspended in chloroform (50 mL). Under ice-cooling, *p*-toluenesulfonyl chloride (2.1 g, 11 mmol) was added, and the reaction mixture was stirred for 5 min at 0 °C and subsequently for 7 d at ambient temperature. Then, the solvent was removed in vacuo and the residue was purified by flash column chromatography ($\emptyset = 6$ cm, h = 21 cm, V = 50 mL, petroleum ether/ethyl acetate = 1:1, $R_f = 0.49$) to give 29 as a colorless oil (2.6 g, 5.0 mmol, 94%). [α]²⁰_D = +26.0 (2.9, methanol); ¹H NMR (DMSO-*d*₆): δ [ppm] = 1.14 (s, 6H, C(CH₃)₂), 2.42 (s, 6H, ArCH₃), 3.62–3.68 (m, 2H, OCH₂CHCH), 3.75–3.79 (m, 2H, OCH₂CHCH), 3.89 (dd, *J* = 10.3/6.6 Hz, 2H, OCH₂CHCH), 4.05 (dd, *J* = 10.3/2.5 Hz, 2H, OCH₂CHCH), 5.58 (d, *J* = 5.5 Hz, 2H, CHOH), 7.45–7.49 (m, 4H, 3'-H_{4-methylbenzenesulfonate, 5'-H_{4-methylbenzenesulfonate}), and 7.75–7.79 (m, 4H, 2'-H_{4-methylbenzenesulfonate}); ¹³C NMR (DMSO-*d*₆): δ [ppm] = 21.1 (2C, ArCH₃), 27.2 (2C, C(CH₃)₂), 69.3 (2C, OCH₂CHCH), 72.0 (2C, OCH₂CHCH), 78.3 (2C, OCH₂CHCH), 109.3 (1C, C(CH₃)₂), 127.7 (4C, C-2'_{4-methylbenzenesulfonate}, C-6'_{4-methylbenzenesulfonate}), 130.1 (4C, C-3'_{4-methylbenzenesulfonate}, C-5'_{4-methylbenzenesulfonate}), 132.2 (2C, C-1'_{4-methylbenzenesulfonate}), and 144.9 (2C, C-4'_{4-methylbenzenesulfonate}); IR (neat): $\tilde{\nu}$ [cm⁻¹] = 3391, 2987, 1735, 1598, 1453, 1356, 1242, 1172, 1075, 974, 937, 902, 812, 664, and 551; HRMS (*m*/*z*): [M + H]⁺ calcd for C₂₃H₃₁O₁₀S₂, 531.1353; found, 531.1348; HPLC (method 1): t_R = 23.6 min, purity 99.7%.}

(1R,1'R)-1,1'-[(4R,5R)-2,2-Dimethyl-1,3-dioxolane-4,5-diyl]bis(2azidoethan-1-ol) (30). Sodium azide (2.2 g, 34 mmol) was added to a solution of 29 (5.7 g, 11 mmol) in DMSO (150 mL), and the reaction mixture was heated to 80 °C for 24 h. Under ice-cooling, water was added and the mixture was extracted with ethyl acetate $(3\times)$. The combined organic layers were dried (Na_2SO_4) , filtered, and the solvent was removed in vacuo. The residue was purified by flash column chromatography (\emptyset = 6 cm, h = 24 cm, V = 50 mL, petroleum ether/ethyl acetate = 1:1, $R_f = 0.78$) to give 30 as a colorless oil (2.6 g, 9.5 mmol, 89%). $[\alpha]_D^{20} = +39.9$ (6.3, methanol); ¹H NMR (DMSO- d_6): δ [ppm] = 1.29 (s, 6H, C(CH₃)₂), 3.28 (dd, J = 12.8/6.7 Hz, 2H, N₃CH₂CHCH), 3.35 (dd, J = 12.8/3.0 Hz, 2H, N₃CH₂CHCH), 3.66-3.74 (m, 2H, N₃CH₂CHCH), 3.85-3.91 (m, 2H, N₃CH₂CHCH), and 5.64 (d, J = 5.3 Hz, 2H, CHOH); ¹³C NMR $(DMSO-d_6): \delta [ppm] = 27.3 (2C, C(CH_3)_2), 53.4 (2C, C(CH_3)_2)$ N₃CH₂CHCH), 71.2 (2C, N₃CH₂CHCH), 79.5 (2C, N₃CH₂CHCH), and 109.1 (1C, C(CH₃)₂); IR (neat): $\tilde{\nu}$ [cm⁻¹] = 3354, 2989, 2934, 2096, 1441, 1373, 1214, 1164, 1069, 870, 656, 555, and 504; HRMS (m/z): $[M + Na]^+$ calcd for C₉H₁₆N₆NaO₄, 295.1125; found, 295.1118.

(4R,5R)-4,5-bis{(R)-2-Azido-1-[(4-iodobenzyl)oxy]ethyl}-2,2-dimethyl-1,3-dioxolane (31). Under a N₂ atmosphere and ice-cooling, sodium hydride (60% suspension in paraffin oil, 210 mg, 5.3 mmol) was added to a solution of 30 (270 mg, 0.98 mmol) in THF (15 mL). After stirring the mixture for 10 min at 0 °C, stirring was continued for 1 h at ambient temperature. Then, 4-iodobenzyl bromide (870 mg, 2.9 mmol) was added, and the reaction mixture was stirred for 72 h at ambient temperature. Afterward, methanol and water were added, and the mixture was extracted with ethyl acetate $(3\times)$. The combined organic layers were dried (Na2SO4), filtered, and the solvent was removed in vacuo. The residue was purified by flash column chromatography ($\emptyset = 4 \text{ cm}, h = 12 \text{ cm}, V = 30 \text{ mL}, \text{ petroleum}$ ether/ethyl acetate = 4:1, R_f = 0.49) to give 31 as a colorless oil (570 mg, 0.81 mmol, 82%). $[\alpha]_{D}^{20} = +17.7$ (6.8, methanol); ¹H NMR (DMSO- d_6): δ [ppm] = 1.31 (s, 6H, C(CH_3)_2), 3.29–3.35 (m, 2H, N₃CH₂CHCH), 3.66-3.73 (m, 4H, N₃CH₂CHCH (2H), N_3CH_2CHCH), 4.09–4.13 (m, 2H, N_3CH_2CHCH), 4.53 (d, J = 11.8 Hz, 2H, OCH₂Ar), 4.58 (d, J = 11.8 Hz, 2H, OCH₂Ar), 7.08-7.13 (m, 4H, 2'-H_{4-iodophenyl}, 6'-H_{4-iodophenyl}), and 7.63-7.67 (m, 4H, $3'-H_{4-iodophenyl}$, $5'-H_{4-iodophenyl}$); ^{13}C NMR (DMSO- d_6): δ [ppm] = 27.1 (2C, C(CH_3)_2), 49.7 (2C, N_3CH_2CHCH), 70.5 (2C, OCH_2Ar), 77.4 (2C, N₃CH₂CHCH), 78.9 (2C, N₃CH₂CHCH), 93.5 (2C, C-4'_{4-iodophenyl}), 109.5 (1C, C(CH₃)₂), 129.8 (4C, C-2'_{4-iodophenyl}, C- $6'_{4-iodophenyl}$), 137.0 (4C, C-3'_{4-iodophenyl}), C-5'_{4-iodophenyl}), and 137.7 (2C, C-1'_{4-iodophenyl}); IR (neat): $\tilde{\nu}$ [cm⁻¹] = 2986, 2933, 2869, 2095, 1590, 1484, 1371, 1238, 1210, 1081, 1006, 867, 828, 797, 656, 629, and 471; HRMS (*m*/*z*): [M + Na]⁺ calcd for C₂₃H₂₆I₂N₆NaO₄, 726.9997; found, 726.9993; HPLC (method 1): $t_{\rm R}$ = 30.1 min, purity 97.0%.

(2R,3S,4S,5R)-1,6-Diazido-2,5-bis[(4-iodobenzyl)oxy]hexane-3,4diol (32). After stirring 31 (4.4 g, 6.3 mmol) in 80% aqueous acetic acid (40 mL) for 5 h at 0 °C, the reaction mixture was diluted with toluene and the solvent was removed in vacuo. The residue was purified by flash column chromatography ($\emptyset = 6$ cm, h = 12 cm, V =50 mL, petroleum ether/ethyl acetate = 1:1, $R_f = 0.51$) to give 32 as a colorless oil (3.8 g, 5.6 mmol, 90%). $[\alpha]_D^{20} = +50.5$ (6.5, methanol); ¹H NMR (DMSO- d_6): δ [ppm] = 3.40 (dd, J = 13.1/4.3 Hz, 2H, N₃CH₂CHCH), 3.61–3.80 (m, 6H, N₃CH₂CHCH (2H), N₃CH₂CHCH, N₃CH₂CHCH), 4.51 (d, J = 11.7 Hz, 2H, OCH₂Ar), 4.62 (d, J = 11.7 Hz, 2H, OCH₂Ar), 4.85 (d, J = 7.9Hz, 2H, CHOH), 7.12–7.19 (m, 4H, 2'-H_{4-iodopheny}, 4.85 (d, J = 7.9Hz, 2H, CHOH), 7.12–7.19 (m, 4H, 2'-H_{4-iodopheny}), ¹³C NMR (DMSO- d_6): δ [ppm] = 50.5 (2C, N₃CH₂CHCH), 68.2 (2C, N₃CH₂CHCH), 70.7 (2C, OCH₂Ar), 78.2 (2C, N₃CH₂CHCH), 93.4 (2C, C-4'_{4-iodopheny}), 129.8 (4C, C-2'_{4-iodopheny}), C-6'_{4-iodopheny}), 137.0 (4C, C-3'_{4-iodopheny}) C-5'_{4-iodopheny}), and 138.2 (2C, C-1'_{4-iodopheny}), 137.0 (4C, C-3'_{4-iodopheny}) C-5'_{4-iodopheny}), and 138.2 (2C, C-1'_{4-iodopheny}), 137.0 (4C, S-3'_{4-iodopheny}) C-5'_{4-iodopheny}), and 138.2 (2C, C-1'_{4-iodopheny}), 137.0 (4C, C-3'_{4-iodopheny}) C-5'_{4-iodopheny}), and 138.2 (2C, C-1'_{4-iodopheny}), 127.0 (4C, C-3'_{4-iodopheny}), C-5'_{4-iodopheny}), and 138.2 (2C, C-1'_{4-iodopheny}), 137.0 (4C, C-3'_{4-iodopheny}) C-5'_{4-iodopheny}), and 138.2 (2C, C-1'_{4-iodopheny}), 137.0 (4C, C-3'_{4-iodopheny}), C-5'_{4-iodopheny}), and 138.2 (2C, C-1'_{4-iodopheny}), 137.0 (4C, C-3'_{4-iodopheny}), C-5'_{4-iodopheny}), 137.0 (4C, C-3'_{4-iodopheny}), C-5'_{4-iodopheny}), C-5'_{4-iodopheny}), C-5'_{4-iodopheny}), 137.0 (4C, C-3'_{4-iodopheny}), C-5'_{4-iodopheny}), C-5'_{4-iodopheny}), C-5'_{4-iodopheny}), C-5'_{4-iodophe}

Methyl (\$)-3-azido-2-[(4-iodobenzyl)oxy]propanoate ((S)-28). An oxidant solution was prepared by dissolving H₅IO₆ (11.4 g, 50 mmol) and CrO₃ (23 mg, 0.23 mmol) in wet acetonitrile (114 mL, 0.75% water V/V) overnight. Under ice-cooling, the oxidant solution (21 mL) was added to a solution of 27 (1.2 g, 3.6 mmol) in acetonitrile (20 mL). After stirring the reaction mixture for 30 min at 0 °C, stirring was continued for 16 h at ambient temperature. Then, Na₂HPO₄ (1.1 g, 7.6 mmol) and water (20 mL) were added under ice-cooling. Afterward, the mixture was diluted with diethyl ether and stirred for 15 min at ambient temperature. Then, 1.0 M HCl (8 mL) was added and the mixture was extracted with diethyl ether $(3\times)$. The combined organic layers were dried (Na2SO4), filtered, and the solvent was removed in vacuo. The residue was dissolved in methanol (60 mL) and concentrated sulfuric acid (0.2 mL) was added. After heating the reaction mixture to reflux for 24 h, the solvent was concentrated in vacuo. Then, the mixture was diluted with dichloromethane and washed with ice-cold water, a saturated aqueous solution of NaHCO₃, and brine. The combined organic layers were dried (Na₂SO₄), filtered, and the solvent was removed in vacuo. The residue was purified by automatic flash column chromatography using an Interchim puriFlash XS 420 system (20% \rightarrow 33% ethyl acetate in petroleum ether, Biotage SNAP Ultra HP-Sphere 50 g) to give (S)-28 as a colorless oil (850 mg, 2.4 mmol, 65%). $R_f = 0.47$ (petroleum ether/ethyl acetate = 4:1; $[\alpha]_{D}^{20} = -47.2$ (7.1, methanol); enantiomeric ratio (HPLC method 4): $t_{\rm R} = 11.2 \text{ min}$, (S)/(R) = 99.6/0.4; HPLC (method 1): $t_{\rm R}$ = 24.3 min, purity 98.9%.

Methyl (R)-3-azido-2-[(4-iodobenzyl)oxy]propanoate ((R)-28). $NaIO_4$ (130 mg, 0.62 mmol) was added to a solution of 32 (250 mg, 0.37 mmol) in methanol (15 mL), and the mixture was stirred at ambient temperature for 16 h. Then, the solvent was concentrated in vacuo, brine was added, and the mixture was extracted with ethyl acetate (3×). The combined organic layers were dried (Na_2SO_4) , filtered, and the solvent was removed in vacuo. The residue was dissolved in a mixture of methanol and water (9:1, 40 mL) and NaHCO₃ (960 mg, 11 mmol) was added. Then, Br₂ (0.06 mL, 190 mg, 1.2 mmol) was added, and the reaction mixture was stirred at ambient temperature for 16 h in a flask protected from ordinary lighting. Afterward, sodium thiosulfate and water were added and the mixture was extracted with ethyl acetate $(3\times)$. The combined organic layers were dried (Na₂SO₄), filtered, and the solvent was removed in vacuo. The residue was purified by flash column chromatography (Ø = 3 cm, h = 20 cm, V = 20 mL, petroleum ether/ethyl acetate = 4:1, R_f = 0.47) to give (R)-28 as a colorless oil (160 mg, 0.45 mmol, 62%). $[\alpha]_{D}^{20}$ = +46.9 (8.2, methanol); enantiomeric ratio (HPLC method 4): $t_{\rm R} = 10.2 \text{ min}, (R)/(S) = 100/0; \text{ HPLC (method 1): } t_{\rm R} = 24.4 \text{ min},$ purity 95.1%.

Spectroscopic Data of (S)-28 and (R)-28. ¹H NMR (DMSO- d_6): δ [ppm] = 3.53 (dd, J = 13.3/5.9 Hz, 1H, CHCH₂N₃), 3.66 (dd, J = 13.3/3.5 Hz, 1H, CHCH₂N₃), 3.70 (s, 3H, CO₂CH₃), 4.36 (dd, J = 5.9/3.5 Hz, 1H, CHCH₂N₃), 4.52 (d, J = 12.0 Hz, 1H, OCH₂Ar), 4.66 (d, J = 12.0 Hz, 1H, OCH₂Ar), 7.15–7.22 (m, 2H, 2'-H_{4-iodophenyl}), 6'-H_{4-iodophenyl}), and 7.70–7.76 (m, 2H, 3'-H_{4-iodophenyl}), 5'-H_{4-iodophenyl}); ¹³C NMR (DMSO- d_6): δ [ppm] = 51.7 (1C, CHCH₂N₃), 52.1 (1C, CO₂CH₃), 71.0 (1C, OCH₂Ar), 77.1 (1C, CHCH₂N₃), 93.7 (1C, C-4'_{4-iodophenyl}), 129.8 (2C, C-2'_{4-iodophenyl}), C-6'_{4-iodophenyl}), 137.0 (2C, C-3'_{4-iodophenyl}), 137.4 (1C, C-1'_{4-iodophenyl}), and 170.0 (1C, CO₂CH₃); IR (neat): $\tilde{\nu}$ [cm⁻¹] = 2952, 2871, 2098, 1748, 1590, 1484, 1436, 1264, 1203, 1122, 1006, 798, 649, 556, and 474; HRMS (*m*/*z*): [M + Na]⁺ calcd for C₁₁H₁₂IN₃NaO₃, 383.9816; found, 383.9819.

Methyl (S)-3-(benzylamino)-2-[(4-iodobenzyl)oxy]propanoate ((S)-33). Under ice-cooling, triethyl phosphite (0.22 mL, 220 mg, 1.3 mmol) was added to a solution of (S)-28 (440 mg, 1.2 mmol) in toluene (2 mL). After stirring the mixture at ambient temperature overnight, freshly distilled benzaldehyde (0.15 mL, 160 mg, 1.5 mmol) was added slowly under ice-cooling. After stirring the reaction mixture at ambient temperature for 72 h, the solvent was removed in high vacuum. The residue was taken up in methanol (2 mL). Under ice-cooling, NaBH₄ (59 mg, 1.6 mmol) was added, and the mixture was stirred at ambient temperature overnight. Then, water was added and the mixture was extracted with ethyl acetate $(3\times)$. The combined organic layers were dried (Na2SO4), filtered, and the solvent was removed in vacuo. The residue was purified by flash column chromatography (\emptyset = 4 cm, h = 26 cm, V = 30 mL, ethyl acetate, $R_f = 0.60$) to give (S)-33 as a colorless oil (350 mg, 0.82 mmol, 67%). $[\alpha]_{D}^{20} = -45.7$ (2.5, methanol); HPLC (method 1): $t_{\rm R} = 20.4$ min, purity 99.3%.

Methyl (R)-3-(benzylamino)-2-[(4-iodobenzyl)oxy]propanoate ((R)-33). Under ice-cooling, triethyl phosphite (0.13 mL, 130 mg, 0.76 mmol) was added to a solution of (R)-28 (270 mg, 0.75 mmol) in toluene (1 mL). After stirring the mixture at ambient temperature overnight, freshly distilled benzaldehyde (0.1 mL, 100 mg, 0.98 mmol) was added slowly under ice-cooling. After stirring the reaction mixture at ambient temperature for 72 h, the solvent was removed in high vacuum. The residue was taken up in methanol (1 mL). Under ice-cooling, NaBH₄ (38 mg, 1.0 mmol) was added, and the mixture was stirred at ambient temperature overnight. Then, water was added and the mixture was extracted with ethyl acetate $(3\times)$. The combined organic layers were dried (Na2SO4), filtered, and the solvent was removed in vacuo. The residue was purified by flash column chromatography ($\emptyset = 4 \text{ cm}, h = 22 \text{ cm}, V = 30 \text{ mL},$ ethyl acetate, $R_f = 0.60$) to give (*R*)-33 as a colorless oil (220 mg, 0.52 mmol, 69%). $[\alpha]_{D}^{20} = +46.7$ (1.2, methanol); HPLC (method 1): $t_{R} = 20.4$ min, purity 97.1%.

Spectroscopic Data of (S)-33 and (R)-33. ¹H NMR (DMSO-d₆): δ [ppm] = 2.23 (s br, 1H, OCHCH₂NH), 2.77 (dd, J = 12.6/6.5 Hz, 1H, OCHCH₂NH), 2.81 (dd, J = 12.6/4.4 Hz, 1H, OCHCH₂NH), 3.65 (s, 3H, CO₂CH₃), 3.66 (d, J = 13.7 Hz, 1H, NHCH₂Ph), 3.69 $(d, J = 13.7 \text{ Hz}, 1\text{H}, \text{NHCH}_2\text{Ph}), 4.15 (dd, J = 6.4/4.4 \text{ Hz}, 1\text{H}, 1\text{H})$ OCHCH₂NH), 4.41 (d, J = 12.1 Hz, 1H, OCH₂Ar), 4.57 (d, J = 12.1 Hz, 1H, OCH₂Ar), 7.15-7.19 (m, 2H, 2'-H_{4-iodophenyl}, 6'-H_{4-iodophenyl}), 7.19-7.23 (m, 1H, 4"-H_{phenyl}), 7.25-7.32 (m, 4H, 2"-H_{phenyl}, 3"- $\begin{array}{l} H_{\text{phenyl}} \; 5'' \cdot H_{\text{phenyl}} \; 6'' \cdot H_{\text{phenyl}} \;), \; \text{and} \; 7.69 - 7.72 \; (m, 2H, 3' \cdot H_{4\text{-iodophenyl}} \;); \; 5' \cdot H_{4\text{-iodophenyl}} \;); \; 1^3 \text{C} \; \text{NMR} \; (\text{DMSO-} d_6): \; \delta \; [\text{ppm}] \; = \; 50.1 \; (1C, 1C) \; (1C) \; (1C)$ OCHCH₂NH), 51.6 (1C, CO₂CH₃), 52.4 (1C, NHCH₂Ph), 70.8 (1C, OCH₂Ar), 77.8 (1C, OCHCH₂NH), 93.4 (1C, C-4'_{4-iodophenyl}), 126.6 (1C, C-4" phenyl), 127.8 (2C, Carom.), 128.1 (2C, Carom.), 129.9 (2C, C-2'_{4-iodophenyl}, C-6'_{4-iodophenyl}), 137.0 (2C, C-3'_{4-iodophenyl}, C-5'_{4-iodophenyl}), 137.7 (1C, C-1'_{4-iodophenyl}), 140.5 (1C, C-1"_{phenyl}), and 171.6 (1C, CO₂CH₃); IR (neat): $\tilde{\nu}$ [cm⁻¹] = 3026, 2949, 2843, 1746, 1590, 1484, 1453, 1435, 1272, 1199, 1120, 1006, 798, 734, 698, and 472; HRMS (m/z): $[M + H]^+$ calcd for C₁₈H₂₁INO₃, 426.0561; found, 426.0549.

Methyl (5)-3-(benzylamino)-2-[(4-{[4-(morpholinomethyl)phenyl]ethynyl]benzyl)oxy]propanoate ((5)-**34**). Under a N_2 atmosphere, copper(I) iodide (9 mg, 0.047 mmol), bis-(triphenylphosphine)palladium(II) chloride (35 mg, 0.050 mmol), and diisopropylamine (2.5 mL) were added to a solution of (S)-**33** (200 mg, 0.46 mmol) in dry THF (12 mL) at ambient temperature, and the mixture was stirred for 20 min. Then, 4-(4-ethynylbenzyl)morpholine (180 mg, 0.87 mmol) was added in two portions at an interval of 30 min. After stirring the reaction mixture for 24 h at ambient temperature, the solvent was removed in vacuo. The residue was dissolved in ethyl acetate and filtered through a short silica gel column. The solvent was removed in vacuo, and the residue was purified by flash column chromatography ($\emptyset = 3 \text{ cm}$, h = 15 cm, V = 20 mL, ethyl acetate, $R_f = 0.22$) to give (S)-34 as a colorless oil (160 mg, 0.32 mmol, 68%). [α]_D²⁰ = -22.0 (1.0, methanol); HPLC (method 1): $t_R = 16.9$ min, purity 96.5%.

Methyl (R)-3-(benzylamino)-2-[(4-{[4-(morpholinomethyl)phenyl]ethynyl]benzyl)oxy]propanoate ((R)-34). Under a N₂ atmosphere, copper(I) iodide (8 mg, 0.042 mmol), bis-(triphenylphosphine)palladium(II) chloride (22 mg, 0.031 mmol), and diisopropylamine (2.5 mL) were added to a solution of (R)-33 (130 mg, 0.30 mmol) in dry THF (10 mL) at ambient temperature, and the mixture was stirred for 20 min. Then, 4-(4-ethynylbenzyl)morpholine (120 mg, 0.60 mmol) was added in two portions at an interval of 30 min. After stirring the reaction mixture for 24 h at ambient temperature, the solvent was removed in vacuo. The residue was dissolved in ethyl acetate and filtered through a short silica gel column. The solvent was removed in vacuo and the residue was purified by flash column chromatography ($\emptyset = 3 \text{ cm}, h = 16 \text{ cm}, V =$ 20 mL, ethyl acetate, $R_f = 0.22$) to give (R)-34 as a colorless oil (100 mg, 0.21 mmol, 69%). $[\alpha]_D^{20} = +23.6$ (1.2, methanol); HPLC (method 1): $t_{\rm R} = 16.7$ min, purity 99.1%.

Spectroscopic Data of (S)-34 and (R)-34. ¹H NMR (DMSO-d₆): δ [ppm] = 2.24 (s br, 1H, OCHCH₂NH), 2.31-2.39 (m, 4H, $N(CH_2CH_2)_2O$, 2.79 (dd, J = 12.6/6.4 Hz, 1H, OCHCH₂NH), 2.83 $(dd, J = 12.6/4.4 Hz, 1H, OCHCH_2NH), 3.49 (s, 2H, NCH_2Ar),$ 3.55–3.60 (m, 4H, N(CH₂CH₂)₂O), 3.66 (s, 3H, CO₂CH₃), 3.67 (d, *J* = 13.8 Hz, 1H, NHCH₂Ph), 3.71 (d, *J* = 13.8 Hz, 1H, NHCH₂Ph), 4.18 (dd, J = 6.4/4.4 Hz, 1H, OCHCH₂NH), 4.49 (d, J = 12.3 Hz, 1H, OCH₂Ar), 4.65 (d, J = 12.3 Hz, 1H, OCH₂Ar), 7.20–7.24 (m, 1H, 4'''-H_{phenyl}), 7.26–7.32 (m, 4H, 2'''-H_{phenyl} 3'''-H_{phenyl} 5'''-H_{phenyl} 5'''-H_{phenyl}, 5'''-H_{phenyl}, 7.34-7.38 (m, 2H, 3''-H₄-(morpholinomethyl)phenyl), 7.39–7.43 (m, 2H, 2'- $\begin{array}{l} H_{4-\{[4-(m \mbox{ or } p \mbox{ hol } n \mbox{ or } m \mbox{ et } h \mbox{ y }] \mbox{ p h } n \mbox{ y } 1] \mbox{ et } h \mbox{ y } n \mbox{ y } 1] \mbox{ et } h \mbox{ y } n \mbox{ y } 1] \mbox{ et } h \mbox{ y } n \mbox{ y } 1] \mbox{ p h } n \mbox{ y } 1] \mbox{$ $\begin{array}{l} H_{4-\{[4-(morpholinomethyl]phenyl]ethynylphenyl)} & f'-H_{4-\{[morpholinomethyl]phenyl]ethynylphenyl} & f'-H_{4-(morpholinomethyl]phenyl)} & f'-H_{4-(morpholinomethyl]phenyl) & f'-H_{4-(morpholinomethyl]phenyl)} & f'-H_{4-(morpholinomethyl]phenyl)} & f'-H_{4-(morpholinomethyl]phenyl)} & f'-H_{4-(morpholinomethyl]phenyl) & f'-H_{4-(morpholinomethyl]phenyl)} & f'-H_{4-(morpholinomethyl]phenyl) & f'-H_{4-(morpholinomethyl]phenyl)} & f'-H_{4-(morpholinomethyl]phenyl) & f'-H_{4-(morpholinome$ (1C, OCHCH₂NH), 51.6 (1C, CO₂CH₃), 52.4 (1C, NHCH₂Ph), 53.2 (2C, N(CH₂CH₂)₂O), 62.0 (1C, ArCH₂N), 66.2 (2C, N(CH₂CH₂)₂O), 71.0 (1C, OCH₂Ar), 77.9 (1C, OCHCH₂NH), 89.0 (1C, C \equiv C), 89.3 (1C, C \equiv C), 120.8 (1C, C-1"4-(morpholinomethyl)phenyl), 121.4 (1C, C-4 - (morpholinomethyl)phenyl)phenyl), 121.7 (1C, C 4'+[[4-(morpholinomethyl)phenyl]phenyl), 126.6 (1C, C-4'''_phenyl), 127.9 (4C, C-2' 4-{[4-(morpholinomethyl)phenyl]phenyl]phenyl]phenyl, C-6'+{[4-(morpholinomethyl)phenyl]phenyl, C-2'''_phenyl, C-6'''_phenyl), 128.1 (2C, C-3'''_phenyl, C-5'''_phenyl), 129.2 (2C, C-3''_4-(morpholinomethyl)phenyl) 5''_4-(morpholinomethyl)phenyl), 131.23 (2C, C_arom,), 131.24 (2C, C_arom), 138.7 (1C, C-1') 138.7 (1C, C-1'_{4-{[4-(morpholinomethyl)phenyl]ethynyl}phenyl), 138.9 (1C, C-} $4''_{4-(morpholinomethyl)phenyl}$, 140.6 (1C, C-1'''_phenyl), and 171.6 (1C, CO₂CH₃); IR (neat): $\tilde{\nu}$ [cm⁻¹] = 2951, 2852, 2808, 1748, 1517, 1453, 1349, 1201, 1114, 1007, 914, 866, 819, 794, 735, 698, 541, and 515; HRMS (m/z): $[M + H]^+$ calcd for C₃₁H₃₅N₂O₄, 499.2591; found, 499.2605.

(S)-3-(Benzylamino)-N-hydroxy-2-[(4-{[4-(morpholinomethyl)phenyl]ethynyl}benzyl)oxy]propanamide ((S)-12a). Under ice-cooling, an aqueous solution of hydroxylamine (50 wt %, 2 mL) was added to a solution of (S)-34 (93 mg, 0.19 mmol) in a mixture of THF (4 mL) and isopropanol (4 mL). After stirring the reaction mixture for 5 min at 0 °C, stirring was continued for 48 h at ambient temperature. Then, the solvent was removed in vacuo and the residue was purified by automatic flash column chromatography using a Biotage Isolera One system (20% \rightarrow 100% ACN in H₂O, Biotage SNAP Ultra C18 30 g). Fractions containing the desired product were combined and subjected to lyophilization to give (S)-12a as a colorless solid (65 mg, 0.13 mmol, 70%). mp 59 °C; $[a]_D^{20} = -29.3$ (1.5, methanol); HPLC (method 2): $t_R = 12.6$ min, purity 98.8%.

(R)-3-(Benzylamino)-N-hydroxy-2-[(4-{[4-(morpholinomethyl)phenyl]ethynyl}benzyl)oxy]propanamide ((R)-12a). Under ice-cooling, an aqueous solution of hydroxylamine (50 wt %, 2 mL) was added to a solution of (R)-34 (58 mg, 0.12 mmol) in a mixture of THF (4 mL) and isopropanol (4 mL). After stirring the reaction mixture for 5 min at 0 °C, stirring was continued for 48 h at ambient temperature. Then, the solvent was removed in vacuo and the residue was purified by automatic flash column chromatography using a Biotage Isolera One system (20% \rightarrow 100% ACN in H₂O, Biotage SNAP Ultra C18 30 g). Fractions containing the desired product were combined and subjected to lyophilization to give (*R*)-**12a** as a colorless solid (36 mg, 0.072 mmol, 62%). mp 59 °C; $[\alpha]_{D}^{20}$ = +32.5 (2.0, methanol); HPLC (method 2): t_{R} = 12.6 min, purity 99.2%.

Spectroscopic Data of (S)-12a and (R)-12a. ¹H NMR (DMSO d_6 : δ [ppm] = 2.30-2.40 (m, 4H, N(CH₂CH₂)₂O), 2.72 (dd, J = 12.5/4.9 Hz, 1H, OCHCH₂NH), 2.77 (dd, J = 12.5/6.9 Hz, 1H, OCHCH₂NH), 3.49 (s, 2H, NCH₂Ar), 3.55-3.61 (m, 4H, $N(CH_2CH_2)_2O)$, 3.68 (s, 2H, NHCH₂Ph), 3.89 (dd, J = 6.8/4.9Hz, 1H, OCHCH₂NH), 4.43 (d, J = 12.5 Hz, 1H, OCH₂Ar), 4.59 (d, $\begin{array}{l} J = 12.5 \text{ Hz}, 1H, \text{ OCH}_2\text{Ar}), 7.17-7.26 \text{ (m, 1H, 4'''-H_{phenyl})}, 7.26-7.33 \text{ (m, 4H, 2'''-H_{phenyl})}, 3'''-H_{phenyl}, 5'''-H_{phenyl}, 6'''-H_{phenyl}), 7.33-7.39 \text{ (m, 2H, 3''-H_4-(morpholinomethyl)phenyl)}, 5'''-H_4-(morpholinomethyl)phenyl)}, 7.39-7.39 \text{ (m, 2H, 3''-H_4-(morpholinomethyl)phenyl)}, 7.39-7.39 \text{ (m, 2H, 3''-H_4-(mor$ 7.45 (m, 2H, $2'-H_{4-\{[4-(morpholinomethyl)phenyl]ethynyl\}phenyl}$, 6'-H_{4-{[4-(morpholinomethyl)phenyl]ethynyl}phenyl)}, 7.48-7.55 (m, 4H, 3'- $\begin{array}{l} H_{4-\{[4-(morpholinomethyl)phenyl] ethynyl \} phenyl \}} & 5' \\ H_{4+\{[4-(morpholinomethyl)phenyl] ethynyl \} phenyl } & 2'' \\ H_{4-(morpholinomethyl)phenyl } & 6'' \\ \end{array}$ H4-(morpholinomethyl)phenyl), and 8.88 (s br, 1H, CONHOH), the signals for OCHCH₂NH and CONHOH cannot be observed in the spectrum; ^{13}C NMR (DMSO- d_6): δ [ppm] = 50.5 (1C, OCHCH₂NH), 52.6 (1C, NHCH₂Ph), 53.2 (2C, N(CH₂CH₂)₂O), 62.0 (1C, ArCH₂N), 66.2 (2C, N(CH₂CH₂)₂O), 70.6 (1C, OCH₂Ar), 77.8 (1C, OCHCH₂NH), 89.1 (1C, C≡C), 89.3 (1C, C≡C), 120.8 (1C, C-1"_{4-(morpholinomethyl)phenyl}), 121.4 (1C, C-4'_{4-{[4-(morpholinomethyl)phenyl}), 126.6 (1C, C-4""_{phenyl}), 127.85 (2C, C_{arom}), 127.90 (2C, C_{arom}), 128.1 (2C, C-3""_{phenyl}), C-5""_{phenyl}), 129.2 (2C, C 2" (129.2 (2C, C-3"_{4-(morpholinomethyl)phenyl)}, C-5"_{4-(morpholinomethyl)phenyl)}, 131.18 (2C, C_{arom.}), 131.23 (2C, C_{arom.}), 138.78 (1C, C_{arom.}), 138.84 (1C, C_{arom.}), 140.6 (1C, C-1""_{phenyl}), and 167.0 (1C, CONUCL) $\mathbb{P} = \begin{bmatrix} -1 \\ 2 \end{bmatrix} = 2170 - 2202 - 2022 - 2$ CONHOH); IR (neat): $\tilde{\nu}$ [cm⁻¹] = 3179, 3028, 2853, 2808, 1660, 1517, 1453, 1349, 1332, 1308, 1291, 1114, 1007, 914, 865, 820, 792, 741, 698, 539, and 517; HRMS (m/z): $[M + H]^+$ calcd for C30H34N3O4, 500.2544; found, 500.2555.

Methyl (S)-3-(dibenzylamino)-2-[(4-iodobenzyl)oxy]propanoate ((S)-35). Under a N₂ atmosphere, polymer-bound triphenylphosphine (~3 mmol/g triphenylphosphine loading, 360 mg, 1.1 mmol) was added to a solution of (S)-28 (200 mg, 0.55 mmol) in a mixture of tetrahydrofuran (12 mL) and water (8 mL). After stirring the reaction mixture for 24 h at 40 °C, the mixture was filtered and the solvent was removed under high vacuum. The residue was taken up in 1,2dichloroethane (10 mL) and freshly distilled benzaldehyde (0.12 mL, 130 mg, 1.2 mmol) was added to the suspension. Under ice-cooling, $NaBH(OAc)_3$ (250 mg, 1.2 mmol) was added to the reaction mixture. After stirring the reaction mixture for 5 min at 0 °C, stirring was continued for 72 h at ambient temperature. Then, a saturated aqueous solution of NaHCO3 was added and the mixture was extracted with ethyl acetate $(3\times)$. The combined organic layers were dried (Na_2SO_4) , filtered, and the solvent was removed in vacuo. The residue was taken up in methanol (10 mL) and concentrated sulfuric acid (0.03 mL, 55 mg, 0.56 mmol) was added. After stirring the reaction mixture for 72 h at 80 °C, the solvent was concentrated in vacuo, water was added, and the mixture was extracted with ethyl acetate $(3\times)$. The combined organic layers were dried (Na_2SO_4) , filtered, and the solvent was removed in vacuo. The residue was purified by flash column chromatography ($\emptyset = 3 \text{ cm}, h = 16 \text{ cm}, V =$ 20 mL, petroleum ether/ethyl acetate = $10:1 \rightarrow 4:1$) to give (S)-35 as a colorless oil (140 mg, 0.28 mmol, 50%). $R_f = 0.44$ (petroleum ether/ ethyl acetate = 10:1); $[\alpha]_{D}^{20} = -7.9$ (1.6, methanol); HPLC (method 1): $t_{\rm R} = 23.6$ min, purity 98.9%.

Methyl (R)-3-(dibenzylamino)-2-[(4-iodobenzyl)oxy]propanoate ((R)-35). Under a N₂ atmosphere, polymer-bound triphenylphosphine (\sim 3 mmol/g triphenylphosphine loading, 430 mg, 1.3 mmol) was added to a solution of (R)-28 (170 mg, 0.46 mmol) in a mixture of tetrahydrofuran (12 mL) and water (8 mL). After stirring the reaction mixture for 24 h at 40 °C, the mixture was filtered and the solvent was

removed in high vacuum. The residue was taken up in 1,2dichloroethane (10 mL) and freshly distilled benzaldehyde (0.10 mL, 100 mg, 0.98 mmol) was added to the suspension. Under icecooling, NaBH(OAc)₃ (210 mg, 0.97 mmol) was added to the reaction mixture. After stirring the reaction mixture for 5 min at 0 °C, stirring was continued for 72 h at ambient temperature. Then, a saturated aqueous solution of NaHCO3 was added and the mixture was extracted with ethyl acetate $(3\times)$. The combined organic layers were dried (Na_2SO_4) , filtered, and the solvent was removed in vacuo. The residue was taken up in methanol (10 mL) and concentrated sulfuric acid (0.03 mL, 55 mg, 0.56 mmol) was added. After stirring the reaction mixture for 72 h at 80 °C, the solvent was concentrated in vacuo, water was added, and the mixture was extracted with ethyl acetate (3×). The combined organic layers were dried (Na_2SO_4) , filtered, and the solvent was removed in vacuo. The residue was purified by flash column chromatography ($\emptyset = 3 \text{ cm}, h = 24 \text{ cm}, V =$ 20 mL, petroleum ether/ethyl acetate = $10:1 \rightarrow 4:1$) to give (*R*)-35 as a colorless oil (100 mg, 0.20 mmol, 43%). $R_f = 0.44$ (petroleum ether/ ethyl acetate = 10:1); $[\alpha]_{D}^{20}$ = +9.4 (1.2, methanol); HPLC (method 1): $t_{\rm R} = 23.6$ min, purity 100%.

Spectroscopic Data of (S)-**35** and (R)-**35**. ¹H NMR (DMSO-*d*₆): δ [ppm] = 2.72 (dd, *J* = 13.4/4.8 Hz, 1H, OCHCH₂N), 2.77 (dd, *J* = 13.4/6.2 Hz, 1H, OCHCH₂N), 3.48 (d, *J* = 13.7 Hz, 2H, N(CH₂Ph)₂), 3.60 (s, 3H, CO₂CH₃), 3.65 (d, *J* = 13.7 Hz, 2H, N(CH₂Ph)₂), 4.27 (dd, *J* = 6.2/4.8 Hz, 1H, OCHCH₂N), 4.38 (d, *J* = 12.0 Hz, 1H, OCH₂Ar), 4.50 (d, *J* = 12.0 Hz, 1H, OCH₂Ar), 7.11– 7.15 (m, 2H, 2'-H_{4-iodophenyl}, 6'-H_{4-iodophenyl}), 7.20–7.25 (m, 2H, 4"-H_{phenyl}), 7.25–7.29 (m, 4H, 2"-H_{phenyl}, 6"-H_{phenyl}), 7.29–7.33 (m, 4H, 3"-H_{phenyl}); ⁵C-M₂H_{4-iodophenyl}, and 7.68–7.73 (m, 2H, 3'-H_{4-iodophenyl}); ⁵C-H_{4-iodophenyl}); δ [ppm] = 51.5 (1C, CO₂CH₃), 54.7 (1C, OCHCH₂N), 93.6 (1C, C-4'_{4-iodophenyl}), 126.9 (2C, C-4"_{phenyl}), 128.1 (4C, C-3"_{phenyl}, C-5"_{phenyl}), 128.6 (4C, C-2"_{phenyl}), C-6'_{4-iodophenyl}), 137.0 (2C, C-3'_{4-iodophenyl}), 139.0 (2C, C-1"_{phenyl}), and 171.3 (1C, CO₂CH₃); IR (neat): $\tilde{\nu}$ [cm⁻¹] = 3026, 2799, 1747, 1589, 1484, 1452, 1201, 1096, 1006, 798, 745, 697, and 472; HRMS (*m*/z): [M + H]⁺ calcd for C₂₅H₂₇INO₃, 516.1030; found, 516.1031.

Methyl (S)-3-(*dibenzylamino*)-2-[(4-{[4-(morpholinomethyl)phenyl]ethynyl}benzyl)oxy]propanoate ((S)-36). Under a N₂ atmosphere, copper(I) iodide (5 mg, 0.026 mmol), bis-(triphenylphosphine)palladium(II) chloride (18 mg, 0.026 mmol), and diisopropylamine (3 mL) were added to a solution of (S)-35 (120 mg, 0.23 mmol) in dry THF (12 mL) at ambient temperature, and the mixture was stirred for 20 min. Then, 4-(4-ethynylbenzyl)morpholine (130 mg, 0.66 mmol) was added in two portions at an interval of 30 min. After stirring the reaction mixture for 24 h at ambient temperature, the solvent was removed in vacuo. The residue was dissolved in a mixture of petroleum ether and ethyl acetate (1:1) and filtered through a short silica gel column. The solvent was removed in vacuo and the residue was purified by flash column chromatography (\emptyset = 3 cm, h = 22 cm, V = 20 mL, petroleum ether/ ethyl acetate = 1:1, R_f = 0.44) to give (*S*)-**36** as a yellow oil (120 mg, 0.21 mmol, 87%). [α]_D²⁰ = -12.1 (1.1, methanol); HPLC (method 1): $t_{\rm R} = 19.4$ min, purity 98.1%.

Methyl (\hat{N})-3-(\hat{d} ibenzylamino)-2-[(4-{[4-(morpholinomethyl]phenyl]ethynyl]benzyl)oxy]propanoate ((\hat{R})-36). Under a N₂ atmosphere, copper(I) iodide (4 mg, 0.021 mmol), bis-(triphenylphosphine)palladium(II) chloride (14 mg, 0.020 mmol), and diisopropylamine (2.5 mL) were added to a solution of (\hat{R})-35 (100 mg, 0.19 mmol) in dry THF (12 mL) at ambient temperature, and the mixture was stirred for 20 min. Then, 4-(4-ethynylbenzyl)morpholine (120 mg, 0.61 mmol) was added in two portions at an interval of 30 min. After stirring the reaction mixture for 24 h at ambient temperature, the solvent was removed in vacuo. The residue was dissolved in a mixture of petroleum ether and ethyl acetate (1:1) and filtered through a short silica gel column. The solvent was removed in vacuo and the residue was purified by flash column chromatography (\emptyset = 2 cm, h = 19 cm, V = 10 mL, petroleum ether/ ethyl acetate = 1:1, $R_f = 0.44$) to give (*R*)-**36** as a yellow oil (92 mg, 0.16 mmol, 81%). $[\alpha]_D^{20} = +10.0$ (1.1, methanol); HPLC (method 1): $t_R = 19.4$ min, purity 100%.

Spectroscopic Data of (S)-**36** and (R)-**36**. ¹H NMR (DMSO-d₆): δ [ppm] = 2.31–2.39 (m, 4H, N(CH₂CH₂)₂O), 2.75 (dd, J = 13.4/ 4.8 Hz, 1H, OCHCH₂N), 2.80 (dd, J = 13.4/6.2 Hz, 1H, OCHCH₂N), 3.49 (s, 2H, NCH₂Ar), 3.51 (d, J = 13.7 Hz, 2H, N(CH₂Ph)₂), 3.55-3.60 (m, 4H, N(CH₂CH₂)₂O), 3.61 (s, 3H, CO_2CH_3), 3.67 (d, J = 13.7 Hz, 2H, N(CH_2Ph)₂), 4.29–4.32 (m, 1H, OCHCH₂N), 4.47 (d, J = 12.2 Hz, 1H, OCH₂Ar), 4.58 (d, J = 12.2Hz, 1H, OCH₂Ar), 7.21–7.26 (m, 2H, 4^m-H_{phenyl}), 7.27–7.34 (m, 8H, 2^m-H_{phenyl}, 3^m-H_{phenyl}, 5^m-H_{phenyl}, 6^m-H_{phenyl}), 7.34-7.39 (m, 4H, $2' - H_{4-\{[4-(morpholinomethyl)phenyl]ethynyl\}phenyl]ethynyl} benyl]ethynyl} benyl]ethynyl} benyl] benyl]ethynyl} 5'' - H_{4-\{[4-(morpholinomethyl)phenyl]ethynyl]phenyl} 5'' - H_{4-(morpholinomethyl)phenyl} 5'' - H_{4-[[4-(morpholinomethyl)phenyl]ethynyl]phenyl} 5'' - H_{4-[[4-(morpholinomethyl)phenyl]ethynyl]phenyl 5'' - H_{4-[[4-(morpholinomethyl)phenyl]ethynyl]phenyl 5'' - H_{4-[[4-(morpholinomethyl)phenyl]ethynyl 5'' - H_{4-[[4-(morpholinomethyl)phenyl]ethynyl 5'' - H_{4-[[4-(morpholinomethyl)phenyl]ethynyl 5'' - H_{4-[[4-(morpholinomethyl)phenyl]ethyn 5'' - H_{4-[[4-(morpholinomethyl)phenyl]ethyn 5'' - H_{4-[[4-(morpholinomethyl)phenyl]ethyn 5'' - H_{4-[[4-(morpholinomethyl]phenyl]ethyn 5'' - H_{4-[[4-(morpholinomethyl]phenyl]ethyl 5'' -$ H_{4-(morpholinomethyl)phenyl}), and 7.48-7.55 (m, 4H, 3'- $\begin{array}{l} H_{4-\{[4-(morpholinomethyl)phenyl)} & H_{1} \\ H_{4-\{[4-(morpholinomethyl)phenyl]ethynylphenyl]ethynylphenyl} & 2''-H_{4-(morpholinomethyl)phenyl)} & 6''-H_{4-(morpholinomethyl)phenyl)} & 1''-H_{4-(morpholinomethyl)phenyl)} & 1''-H_{4-(morpholinomethyl)phenyl) & 1''-H_{4-(morpholinomethyl)phenyl)} & 1''-H_{4-(morpholinomethyl)phenyl)} & 1''-H_{4-(morpholinomethyl)phenyl) & 1''-H_{4-(morpholinomethyl)phenyl)} & 1''-H_{4-(morpholinomethyl)phenyl) & 1''-H_{4-(morpholinomethyl)phenyl)} & 1''-H_{4-(morpholinomethyl)phenyl) & 1''-H_{4-(morpholinomethyl)phenyl) & 1''-H_{4-(morpholinomethyl)phenyl) & 1''-H_{4-(morpholinomethyl)phenyl) & 1''-H_{4-(morpholinomethyl)phenyl)} & 1''-H_{4-(morpholinomethyl)phenyl)$ $(1C, CO_2CH_3), 53.2$ (2C, N(CH₂CH₂)₂O), 54.7 (1C, OCHCH₂N), 58.0 (2C, $N(CH_2Ph)_2$), 62.0 (1C, $ArCH_2N$), 66.2 (2C, N-(CH₂CH₂)₂O), 71.1 (1C, OCH₂Ar), 77.7 (1C, OCHCH₂N), 89.0 (1C, C≡C), 89.3 (1C, C≡C), 120.8 (1C, C-1"_{4-(morpholinomethyl)phenyl}), $\begin{array}{c} 3' 4 - \{ [4 - (morpholinomethyl)phenyl]ethynyl \} p h e n y l \} p h e n y l \} p h e n y l \} C - 5' 4 - \{ [4 - (morpholinomethyl)phenyl]ethynyl \} phenyl \\ \end{array} \\ \begin{array}{c} C - 2'' 4 - (morpholinomethyl)phenyl \\ \end{array} \\ \end{array}$ 6 " $(4 - (morpholinomethyl)phenyl ethynyl)phenyb C 2 + (morpholinomethyl)phenyb C 3 + ((morpholinomethyl)phenyb C 3 + ((morpholinomethyl)phenyb 3 + (1 C, C - 1' 4 - (4 - (morpholinomethyl)phenyl)phenyl), 138.9 (1 C, C - 4'' 4 - ((morpholinomethyl)phenyl), 139.0 (2 C, C - 1''' phenyl), and 171.4 (1 C, CO₂CH₃); IR (neat): <math>\tilde{\nu}$ [cm⁻¹] = 3027, 2950, 2852, 2803, 1748, 1517, 1494, 1453, 1349, 1290, 1261, 1203, 1114, 1098, 1007, 978, 913, 866, 819, 746, 698, 540, and 515; HRMS (m/z): $[M + H]^+$ calcd for C38H41N2O4, 589.3061; found, 589.3081.

(5)-3-(Dibenzylamino)-N-hydroxy-2-[(4-{[4-(morpholinomethyl)phenyl]ethynyl]benzyl)oxy]propanamide ((S)-37). Under ice-cooling, an aqueous solution of hydroxylamine (50 wt %, 2 mL) was added to a solution of (S)-36 (78 mg, 0.13 mmol) in a mixture of THF (4 mL) and isopropanol (4 mL). After stirring the reaction mixture for 5 min at 0 °C, stirring was continued for 48 h at ambient temperature. Then, the solvent was removed in vacuo and the residue was purified by automatic flash column chromatography using a Biotage Isolera One system (10% \rightarrow 90% ACN in H₂O, Biotage SNAP Ultra C18 30 g). Fractions containing the desired product were combined and subjected to lyophilization to give (S)-37 as a colorless solid (37 mg, 0.063 mmol, 47%). mp 79 °C; $[\alpha]_D^{20} = -22.0$ (1.3, methanol); HPLC (method 2): $t_R = 13.5$ min, purity 100%.

(*R*)-3-(*Dibenzylamino*)-*N*-hydroxy-2-[(4-[[4-(morpholinomethyl)phenyl]ethynyl]benzyl)oxy]propanamide ((*R*)-37). Under ice-cooling, an aqueous solution of hydroxylamine (50 wt %, 2 mL) was added to a solution of (*R*)-36 (70 mg, 0.12 mmol) in a mixture of THF (4 mL) and isopropanol (4 mL). After stirring the reaction mixture for 5 min at 0 °C, stirring was continued for 48 h at ambient temperature. Then, the solvent was removed in vacuo and the residue was purified by automatic flash column chromatography using a Biotage Isolera One system (20% \rightarrow 100% ACN in H₂O, Biotage SNAP Ultra C18 30 g). Fractions containing the desired product were combined and subjected to lyophilization to give (*R*)-37 as a colorless solid (35 mg, 0.059 mmol, 50%). mp 79 °C; $[\alpha]_D^{20} = +28.5$ (1.1, methanol); HPLC (method 2): $t_R = 13.6$ min, purity 100%.

Spectroscopic Data of (S)-**37** and (R)-**37**. ¹H NMR (DMSO-*d*₆): δ [ppm] = 2.31–2.40 (m, 4H, N(CH₂CH₂)₂O), 2.67 (dd, *J* = 13.6/ 4.4 Hz, 1H, OCHCH₂N), 2.74 (dd, *J* = 13.6/7.3 Hz, 1H, OCHCH₂N), 3.49 (s, 2H, NCH₂Ar), 3.53 (d, *J* = 14.0 Hz, 2H, N(CH₂Ph)₂), 3.55–3.61 (m, 4H, N(CH₂CH₂)₂O), 3.64 (d, *J* = 14.0 Hz, 2H, N(CH₂Ph)₂), 4.04 (dd, *J* = 7.3/4.4 Hz, 1H, OCHCH₂N), 4.41 (d, *J* = 12.4 Hz, 1H, OCH₂Ar), 4.55 (d, *J* = 12.4 Hz, 1H, OCH₂Ar), 7.20–7.25 (m, 2H, 4^m-H_{phenvl}), 7.27–7.34 (m, 8H, 2^m-

H_{phenyb} 3^m-H_{phenyb} 5^m-H_{phenyb} 6^m-H_{phenyl}), 7.35-7.38 (m, 2H, 3ⁿ-H_{4-(morpholinomethyl)phenyl}, 5"-H_{4-(morpholinomethyl)phenyl}), 7.38–7.42 (m, 2H, $\begin{array}{l} 2'-H_{4-\{[4-(morpholinomethyl)phenyl]ethynyl\}phenyl\}} & 6'-\\ H_{4-\{[4-(morpholinomethyl)phenyl]ethynyl\}phenyl\}}, & 7.49-7.55 & (m, 4H, 3'-\\ \end{array}$ $\begin{array}{l} H_{4-\{[4-(morpholinomethyl)phenyl]ethynyl]ethynyl]ethynyl]phenyl, 5'-H_{4-\{[4-(morpholinomethyl)phenyl, 2''-H_{4-(morpholinomethyl)phenyl, 6''-H_{4-(morpholinomethyl)phenyl, 6''-H_{4 H_{4-(morpholinomethyl)phenyl}$, 8.93 (s br, 1H, CONHOH), and 10.78 (s br, 1H, CONHOH); ¹³C NMR (DMSO-d₆): δ [ppm] = 53.1 (2C, N(CH₂CH₂)₂O), 55.6 (1C, OCHCH₂N), 57.5 (2C, N(CH₂Ph)₂), 62.0 (1C, ArCH₂N), 66.2 (2C, N(CH₂CH₂)₂O), 70.5 (1C, OCH₂Ar), 76.8 (1C, OCHCH₂N), 89.1 (1C, C≡C), 89.3 (1C, C≡C), 120.8 (1C, C-1"_{4-(morpholinomethyl)phenyl}), 121.4 (1C, C-4'4-{[4-(morpholinomethyl)phenyl]ethynyl}phenyl), 126.8 (2C, C-4^{'''}phenyl), 127.8 $\begin{array}{c} (2C, C-2'_{4-{[4-(morpholinomethyl]phenyl]phenyl], 120:0 (2C, C-4-phenyl), 127:0 (2C, C-2'_{4-{[4-(morpholinomethyl]phenyl], 128:1 (4C, C-3'''phenyl), C-6''_{4-{[4-(morpholinomethyl]phenyl], 128:1 (4C, C-3'''phenyl), 128:6 (4C, C-2'''phenyl), 128:1 (4C, C-3'''phenyl), 128:6 (4C, C-2'''phenyl), 129:2 (2C, C-3''_{4-(morpholinomethyl)phenyl), 128:6 (4C, C-3'''_{4-(morpholinomethyl)phenyl), 131:17 (2C, C_{arom.}), 131:23 (2C, C_{arom.}), 138:7 (1C, C-3) \\ \end{array}$ $1'_{4-\{[4-(morpholinomethyl)phenyl]ethynyl\}phenyl})$, 138.8 (1C, C- $4''_{4-(morpholinomethyl)phenyl}$, 139.0 (2C, C- $1'''_{phenyl}$), and 166.8 (1C, C-2C) [2021] CONHOH); IR (neat): $\tilde{\nu}$ [cm⁻¹] = 3181, 3027, 2854, 2806, 1666, 1517, 1494, 1453, 1349, 1112, 1006, 914, 864, 819, 792, 746, 698, 539, and 517; HRMS (m/z): $[M + H]^+$ calcd for $C_{37}H_{40}N_3O_4$, 590.3013; found, 590.3019.

Methyl (S)-3-benzamido-2-[(4-iodobenzyl)oxy]propanoate ((S)-38a). Under a N₂ atmosphere, a 1.0 M solution of trimethylphosphane in toluene (1.5 mL, 1.5 mmol) was added to an ice-cooled mixture of benzoic acid (82 mg, 0.67 mmol), (S)-28 (220 mg, 0.6 mmol), and 2,2'-dithiodipyridine (75 mg, 0.34 mmol). After stirring the reaction mixture for 20 min at 0 °C, the ice-bath was removed, and the mixture was stirred for 24 h at ambient temperature. Then, water (1.5 mL) was added, and the mixture was stirred for 20 min. Afterward, the mixture was diluted with dichloromethane and washed with a saturated aqueous solution of NaHCO₃, an ice-cold solution of 1.0 M HCl, and water. The combined organic layers were dried (Na₂SO₄), filtered, and the solvent was removed in vacuo. The residue was purified by flash column chromatography ($\emptyset = 3 \text{ cm}, h =$ 21 cm, V = 20 mL, petroleum ether/ethyl acetate = 2:1, $R_f = 0.37$) to give (S)-38a as a colorless solid (220 mg, 0.51 mmol, 83%). mp 117 °C; $[\alpha]_{D}^{20} = -39.6$ (2.7, methanol); HPLC (method 1): $t_{R} = 22.7$ min, purity 99.5%.

Methyl (R)-3-benzamido-2-[(4-iodobenzyl)oxy]propanoate ((R)-38a). Under a N₂ atmosphere, a 1.0 M solution of trimethylphosphane in toluene (1.7 mL, 1.7 mmol) was added to an ice-cooled mixture of benzoic acid (82 mg, 0.67 mmol), (R)-28 (240 mg, 0.66 mmol), and 2,2'-dithiodipyridine (74 mg, 0.34 mmol). After stirring the reaction mixture for 20 min at 0 °C, the ice-bath was removed, and the mixture was stirred for 24 h at ambient temperature. Then, water (1.0 mL) was added, and the mixture was stirred for 20 min. Afterward, the mixture was diluted with dichloromethane and washed with a saturated aqueous solution of NaHCO₃, an ice-cold solution of 1.0 M HCl, and water. The combined organic layers were dried (Na₂SO₄), filtered, and the solvent was removed in vacuo. The residue was purified by flash column chromatography ($\emptyset = 3 \text{ cm}, h =$ 26 cm, V = 20 mL, petroleum ether/ethyl acetate = 2:1, $R_f = 0.37$) to give (R)-38a as a colorless solid (190 mg, 0.43 mmol, 65%). mp 117 °C; $[\alpha]_{D}^{20} = +38.3$ (6.8, methanol); HPLC (method 1): $t_{R} = 22.8$ min, purity 99.4%.

Spectroscopic Data of (S)-**38a** and (R)-**38a**. ¹H NMR (DMSOd₆): δ [ppm] = 3.45–3.54 (m, 1H, OCHCH₂NH), 3.60–3.71 (m, 4H, OCHCH₂NH (1H), CO₂CH₃), 4.19 (dd, *J* = 7.3/5.0 Hz, 1H, OCHCH₂NH), 4.41 (d, *J* = 12.3 Hz, 1H, OCH₂Ar), 4.58 (d, *J* = 12.3 Hz, 1H, OCH₂Ar), 7.10–7.17 (m, 2H, 2'-H_{4-iodophenyl}, 6'-H_{4-iodophenyl}), 7.43–7.50 (m, 2H, 3"-H_{benzoyl}, S"-H_{benzoyl}), 7.50–7.57 (m, 1H, 4"-H_{benzoyl}), 7.58–7.65 (m, 2H, 3'-H_{4-iodophenyl}, 5'-H_{4-iodophenyl}), 7.77–7.84 (m, 2H, 2"-H_{benzoyl}, 6"-H_{benzoyl}), and 8.67 (t, *J* = 5.9 Hz, 1H, CONH); ¹³C NMR (DMSO-*d*₆): δ [ppm] = 41.4 (1C, OCHCH₂NH), 51.9 (1C, CO₂CH₃), 70.7 (1C, OCH₂Ar), 76.5 (1C, OCHCH₂NH), 93.6 (1C, C-4'_{4-iodophenyl}), 127.2 (2C, C-2"_{benzoyl}), C-6"_{benzoyl}), 128.3 (2C, C-3"_{benzoyl}, C-5"_{benzoyl}), 129.9 (2C, C-2'_{4-iodophenyl}), C-6'_{4-iodophenyl}), 131.3 (1C, C-4"_{benzoyl}), 134.2 (1C, C-1"_{benzoyl}), 136.9 (2C, C-3'_{4-iodophenyl}), C-5'_{4-iodophenyl}), 137.5 (1C, C-1'_{4-iodophenyl}), 166.5 (1C, CONH), and 171.0 (1C, CO₂CH₃); IR (neat): $\tilde{\nu}$ [cm⁻¹] = 3279, 2948, 1741, 1635, 1532, 1485, 1257, 1205, 1139, 1121, 1007, 796, 694, and 433; HRMS (*m*/*z*): [M + Na]⁺ calcd for C₁₈H₁₈INNaO₄, 462.0173; found, 462.0162.

Methyl (S)-3-benzamido-2-[(4-{[4-(morpholinomethyl)phenyl]ethynyl]benzyl)oxy]propanoate ((S)-39a). Under a N2 atmosphere, copper(I) iodide (6.8 mg, 0.036 mmol), bis(triphenylphosphine)palladium(II) chloride (26 mg, 0.037 mmol), and diisopropylamine (3 mL) were added to a solution of (S)-38a (160 mg, 0.36 mmol) in dry THF (15 mL) at ambient temperature and the mixture was stirred for 20 min. Then, 4-(4-ethynylbenzyl)morpholine (170 mg, 0.82 mmol) was added in two portions at an interval of 30 min. After stirring the reaction mixture for 24 h at ambient temperature, the solvent was removed in vacuo. The residue was dissolved in a mixture of petroleum ether and ethyl acetate (1:4) and filtered through a short silica gel column. The solvent was removed in vacuo and the residue was purified by flash column chromatography ($\emptyset = 4 \text{ cm}, h = 13 \text{ cm},$ V = 30 mL, petroleum ether/ethyl acetate = 1:4, $R_f = 0.24$) to give (S)-39a as a yellow oil (170 mg, 0.34 mmol, 94%). $[\alpha]_{\rm D}^{20} = -31.5$ (3.4, methanol); HPLC (method 2): $t_{\rm R}$ = 14.1 min, purity 95.9%.

Methyl (R)-3-benzamido-2-[(4-{[4-(morpholinomethyl)phenyl]ethynyl]benzyl)oxy]propanoate ((R)-39a). Under a N₂ atmosphere, copper(I) iodide (27 mg, 0.14 mmol), bis(triphenylphosphine)palladium(II) chloride (65 mg, 0.093 mmol), and diisopropylamine (10 mL) were added to a solution of (R)-38a (190 mg, 0.44 mmol) in dry THF (5 mL) at ambient temperature and the mixture was stirred for 20 min. Then, 4-(4-ethynylbenzyl)morpholine (130 mg, 0.65 mmol) was added in two portions at an interval of 30 min. After stirring the reaction mixture for 24 h at ambient temperature, the solvent was removed in vacuo. The residue was dissolved in a mixture of petroleum ether and ethyl acetate (1:4) and filtered through a short silica gel column. The solvent was removed in vacuo and the residue was purified by flash column chromatography ($\emptyset = 3 \text{ cm}, h = 21 \text{ cm},$ V = 20 mL, petroleum ether/ethyl acetate = $1:2 \rightarrow 1:4$) to give (R)-**39a** as a yellow oil (150 mg, 0.29 mmol, 66%). $R_f = 0.24$ (petroleum ether/ethyl acetate = 1:4); $[\alpha]_{D}^{20}$ = +29.1 (3.5, methanol); HPLC (method 2): $t_{\rm R} = 14.1$ min, purity 96.1%.

Spectroscopic Data of (S)-39a and (R)-39a. ¹H NMR (DMSO d_6): δ [ppm] = 2.30-2.41 (m, 4H, N(CH₂CH₂)₂O), 3.57-3.61 (m, 7H, NCH₂Ar, N(CH₂CH₂)₂O, OCHCH₂NH (1H)), 3.61-3.72 (m, 4H, OCHCH₂NH (1H), CO₂CH₃), 4.24 (dd, J = 7.2/5.1 Hz, 1H, OCHCH₂NH), 4.50 (d, J = 12.5 Hz, 1H, OCH₂Ar), 4.66 (d, J = 12.5 Hz, 1H, OCH_2Ar), 7.32-7.42 (m, 4H, 2'-6′- $\begin{array}{l} H_{4-\{[4-(morpholinomethyl)phenyl]ethynyl]ethynyl}phenyl]ethynyl}phenyl, & 6'-\\ H_{4-\{[4-(morpholinomethyl)phenyl]ethynyl}phenyl, & 3''-H_{4-(morpholinomethyl)phenyl, } & 5''-\\ H_{4-(morpholinomethyl)phenyl), & 7.42-7.55 & (m, 7H, 3'-)\\ \end{array}$ $\begin{array}{l} H_{4-\{[4-(morpholinomethyl)phenyl]ethynyl]ethynyl]phenyl}, 5' - \\ H_{4+\{[4-(morpholinomethyl)phenyl]ethynyl]phenyl}, 2'' - H_{4-(morpholinomethyl)phenyl}, 6'' - \\ \end{array}$ H4-(morpholinomethyl)phenyb 3^{'''}-Hbenzoyb 4^{'''}-Hbenzoyb 5^{'''}-Hbenzoyb), 7.79-7.86 (m, 2H, 2^{'''}-H_{benzoyl}, 6^{'''}-H_{benzoyl}), and 8.68 (t, J = 5.9 Hz, 1H, CONH); ¹³C NMR (DMSO- d_6): δ [ppm] = 41.4 (1C, OCHCH₂NH), 51.9 (1C, CO_2CH_3), 53.1 (2C, $N(CH_2CH_2)_2O$), 62.0 (1C, ArCH₂N), 66.2 (2C, N(CH₂CH₂)₂O), 70.9 (1C, OCH₂Ar), 76.6 (1C, OCHCH₂NH), 89.0 (1C, C≡C), 89.3 (1C, C≡C), 120.8 (1C, C-1"_{4-(morpholinomethyl)phenyl}), 121.5 (1C, C-4'4-{[4-(morpholinomethyl)phenyl]ethynyl]phenyl], 127.2 (2C, C-2^mbenzoyl, C-6^{""}benzoyl), 127.8 (2C, C-2[']_{4-{[4-(morpholinomethyl)phenyl]ethynyl}phenyl} C- $\begin{array}{l} & & benzoy() \quad 12' \text{ is } (2c) \quad c \geq 4 + [1 + (\text{morpholinomethyl)phenyl]ethynyl]phenyl \\ & & 6' + \{[4 - (\text{morpholinomethyl)phenyl]ethynyl]phenyl), \quad 128.3 \quad (2C, C-3''' benzoyl), \\ & & 5''' benzoyl), \quad 129.2 \quad (2C, C-3'' + - (\text{morpholinomethyl})phenyl), \quad C-5''' + (\text{morpholinomethyl})phenyl), \quad 131.15 \quad (2C, Carom.), \quad 131.21 \quad (3C, C-4''' benzoyl), \\ & & Carom.), \quad 134.2 \quad (1C, C-1''' benzoyl), \quad 138.4 \quad (1C, C-1'' benzoyl), \quad 138.9 \quad (1C, C-1'' + [[4 - (\text{morpholinomethyl})phenyl]]ethynyl]phenyl]), \quad 138.9 \quad (1C, C-1''' benzoyl), \quad 138.9 \quad (1C, C-1''' benzoyl), \quad 128.9 \quad (1C, C-1''' benzoyl), \quad (1C, C-1'' benzoyl), \quad (1C, C-1'' benzoyl), \quad (1C, C-1''' benzoyl), \quad (1C, C-1'' benzoyl), \quad (1C, C-1''' benzoyl), \quad ($ $4''_{4-(morpholinomethyl)phenyl}$, 166.5 (1C, CONH), and 171.0 (1C, CO₂CH₃); IR (neat): $\tilde{\nu}$ [cm⁻¹] = 3336, 2951, 2855, 2808, 1745, 1645, 1518, 1290, 1205, 1113, 1006, 914, 864, 819, 693, and 540; HRMS (m/z): $[M + H]^+$ calcd for C₃₁H₃₃N₂O₅, 513.2384; found, 513.2351.

(S)-*N*-{3-(*Hydroxyamino*)-2-[(4-{[4-(morpholinomethyl)phenyl]ethynyl}benzyl)oxy]-3-oxopropyl}benzamide ((S)-13a). Under icecooling, an aqueous solution of hydroxylamine (50 wt %, 3 mL) was added to a solution of (S)-39a (140 mg, 0.27 mmol) in a mixture of THF (5 mL) and isopropanol (5 mL). After stirring the reaction mixture for 5 min at 0 °C, stirring was continued for 24 h at ambient temperature. Then, the solvent was removed in vacuo and the residue was purified by automatic flash column chromatography using a Biotage Isolera One system (10% \rightarrow 80% ACN in H₂O, Biotage SNAP Ultra C18 12 g). Fractions containing the desired product were combined and subjected to lyophilization to give (S)-13a as a colorless solid (56 mg, 0.11 mmol, 41%). mp 86 °C; $[a]_D^{20} = -47.0$ (1.0, methanol); HPLC (method 2): $t_R = 12.9$ min, purity 100%.

(*R*)-*N*-{3-(*Hydroxyamino*)-2-[(4-{[(4-(morpholinomethyl))phenyl]ethynyl]benzyl)oxy]-3-oxopropyl]benzamide ((*R*)-13a). Under icecooling, an aqueous solution of hydroxylamine (50 wt %, 2.5 mL) was added to a solution of (*R*)-39a (100 mg, 0.20 mmol) in a mixture of THF (4 mL) and isopropanol (4 mL). After stirring the reaction mixture for 5 min at 0 °C, stirring was continued for 36 h at ambient temperature. Then, the solvent was removed in vacuo and the residue was purified by automatic flash column chromatography using a Biotage Isolera One system (10% \rightarrow 75% ACN in H₂O, Biotage SNAP Ultra C18 12 g). Fractions containing the desired product were combined and subjected to lyophilization to give (*R*)-13a as a colorless solid (70 mg, 0.14 mmol, 69%). mp 86 °C; $[\alpha]_D^{20} = +44.7$ (1.7, methanol); HPLC (method 2): $t_R = 13.0$ min, purity 99.6%.

Spectroscopic Data of (S)-13a and (R)-13a. ¹H NMR (DMSO d_6 : δ [ppm] = 2.30–2.40 (m, 4H, N(CH₂CH₂)₂O), 3.45–3.51 (m, 3H, NCH₂Ar, OCHCH₂NH (1H)), 3.54-3.60 (m, 5H, N- $(CH_2CH_2)_2O_1$ OCHCH₂NH (1H)), 4.01 (dd, J = 7.9/4.8 Hz, 1H, OCHCH₂NH), 4.43 (d, J = 12.6 Hz, 1H, OCH₂Ar), 4.61 (d, J = 12.6 Hz, 1H, OCH_2Ar), 7.34-7.40 (m, 4H, 2'- $\begin{array}{l} H_{4-\{[4-(m \mbox{ or } p \mbox{ ol } i \mbox{ om } e \mbox{ th } y]) \mbox{ phenyl}] \mbox{ eth } y1] \mbox{ phenyl}, & 6'-\\ H_{4-\{[4-(m \mbox{ or } p \mbox{ ol } i \mbox{ om } e \mbox{ th } y]) \mbox{ phenyl}, & 3''-H_{4-(m \mbox{ or } p \mbox{ ol } i \mbox{ om } e \mbox{ th } y]) \mbox{ phenyl}, & 5''-\\ H_{4-(m \mbox{ or } p \mbox{ ol } i \mbox{ om } e \mbox{ th } y]) \mbox{ phenyl}, & 7.40-7.44 \mbox{ (m, 2H, 3'-)}, & 3'-\\ \end{array}$ 5"'-H_{benzoyl}), 7.49-7.51 (m, 2H, 2"-H_{4-(morpholinomethyl)phenyl, 6"-} H_{4-(morpholinomethyl)phenyl}), 7.51-7.55 (m, 1H, 4^m-H_{benzoyl}), 7.81-7.85 $(m, 2H, 2'''-H_{benzovl}, 6'''-H_{benzovl}), 8.60$ (t, J = 5.7 Hz, 1H, CONHCH₂), 8.98 (s br, 1H, CONHOH), and 10.89 (s br, 1H, CONHOH); ¹³C NMR (DMSO- d_6): δ [ppm] = 41.7 (1C, OCHCH₂NH), 53.2 (2C, N(CH₂CH₂)₂O), 62.0 (1C, ArCH₂N), 66.2 (2C, N(CH₂CH₂)₂O), 70.6 (1C, OCH₂Ar), 76.7 (1C, OCHCH₂NH), 89.0 (1C, C≡C), 89.3 (1C, C≡C), 120.8 (1C, C-1"4-(morpholinomethyl)phenyl), 121.4 (1C, C-4'4+{[4-(morpholinomethyl)phenyl]ethynyl}phenyl), 127.2 (2C, C-2^{'''}benzoyl) C- $\begin{array}{l} & (1+(morpholinomethyl)phenyl (2C, C-2'_4-[[4-(morpholinomethyl)phenyl]ethynyl]phenyl C-6'_4-[[4-(morpholinomethyl)phenyl], 128.2 (2C, C-3'''_{benzoyl}), 129.2 (2C, C-3''_4-(morpholinomethyl)phenyl), C-5'''_{benzoyl}, 129.2 (2C, C-3''_4-(morpholinomethyl)phenyl), C-5'''_4-(morpholinomethyl)phenyl), 131.10 (2C, C-2''_4-(morpholinomethyl)phenyl), C-5'''_4-(morpholinomethyl)phenyl), C-5'''_4-(morpholinomethyl)phenyl), C-5''_4-(morpholinomethyl)phenyl), C-5''_4-($ 3 ′ $\begin{array}{l} 3'_{4-\{[4-(morpholinomethyl)phenyl]ethynyl]ethynyl}phenyl, C-5'_{4-\{[4-(morpholinomethyl)phenyl]ethynyl}phenyl), 131.18 (1C, C-4'''_{benzoyl}), \end{array}$ C -131.22 (2C, C-2"₄(morpholinomethyl)phenyl), C-6"₄(morpholinomethyl)phenyl), 134.3 (1C, C-1"" b e n z o y 1), 138.7 (1C, C- $1'_{4-[[4-(morpholinomethyl)phenyl]ethynyl]phenyl)}, 138.8 (1C, C-4''_{4-(morpholinomethyl)phenyl}), 166.2 (1C, CONHOH), and 166.5 (1C, CONHCH₂); IR (neat): <math>\tilde{\nu}$ [cm⁻¹] = 3245, 2860, 2807, 1644, 1529, 1292, 1113, 1006, 864, 692, and 541; HRMS (m/z): $[M + H]^+$ calcd for C₃₀H₃₂N₃O₅, 514.2336; found, 514.2333.

NMR Experiments. Standard 1D and STD NMR spectra were acquired at 20 °C with a Bruker 600 MHz NMR spectrometer equipped with a 5 mm cryoprobe. Parameters for the STD experiments (saturation frequency and saturation time) were identical for all samples. Selective saturation of the protein NMR spectrum was achieved with the decoupler offset at 0.5 ppm, and nonsaturation control was performed at 15,000 Hz downfield. STD saturation time was 2 s. WaterLOGSY mixing time was 1.5 s. Two NOESY experiments were recorded with mixing times of 0.6 and 0.3 s. STD and WaterLOGSY spectra were recorded with the same NMR tubes

containing 3 μ M LpxC, compound 9 at 500 μ M, and fragments at 500 μ M. For NOESY experiments, NMR tubes contained 6 μ M LpxC.

Temperature was set to 293 K for all NMR experiments. Water suppression was achieved with the excitation sculpting sequence in all experiments.

STD signals were measured for protons in the aromatic region only. The STD effects were measured as the ratio between the intensities of the STD signal and the 1D signal ($I_{\text{STD}}/I_{\text{1D}}$). STD effects were then normalized by setting the largest STD effect to 100%.

Biological Evaluation. *Disk Diffusion assay.* The disc diffusion assays against *E. coli* BL21(DE3) and the defective strain *E. coli* D22 were performed as follows: liquid cultures of the bacteria were grown overnight in lysogeny broth $(LB)^{73}$ at 37 °C and 200 rpm. 150 μ L of an overnight cell suspension was spread evenly onto LB agar plates. 0.15 μ mol of each compound (dissolved in 10 or 15 μ L DMSO) were applied onto circular filter paper ($\emptyset = 6$ mm, Cytiva). Pure DMSO, serving as a negative control, and CHIR-090, serving as a positive control, were also spotted. The agar plates were incubated overnight at 37 °C, and the diameter of the zone of growth inhibition was measured for each compound. Each assay was performed at least three times on separate days.

The disc diffusion assays against *E. coli* TOP10 (Thermo Fisher, Waltham, USA), *E. coli* ATCC 35218, *K. pneumoniae* ATCC 700603, and *P. aeruginosa* ATCC 27853 were performed as follows: the bacteria were grown overnight on a Columbia blood agar plate (Oxoid, Basingstoke, UK), and one colony was suspended in sterile saline to yield a suspension of 0.5 McFarland standard. Using a sterile swab, the suspension was spread evenly onto Mueller–Hinton agar (Oxoid, Basingstoke, UK). 0.15 μ mol of each compound (dissolved in 10 μ L DMSO) was applied onto circular 6 mm diameter filter paper disks, which were then placed on the agar. After incubating the agar plates for 20 h at 37 °C, the diameter of the zone of growth inhibition was measured for each compound.

Minimum Inhibitory Concentration (MIC). The MIC values of the compounds were determined by means of the microdilution method using 96-well plates.

To determine the MIC values against *E. coli* BL21(DE3) and *E. coli* D22, the bacteria were grown overnight in LB at 37 °C and 200 rpm. The overnight suspension was diluted 1:1000 in fresh LB. 10 μ L of a 2-fold dilution series of the compounds in DMSO and 90 μ L of LB were dispensed to each well of a 96-well plate. Then, 100 μ L of the inoculated medium was added, resulting in 5 × 10⁵ cfu mL⁻¹, 5% DMSO, and a final concentration range of the test compounds between 64 and 0.016 μ g mL⁻¹. The plates were incubated for 20 h at 37 °C. The MIC was defined as the lowest concentration of the compounds that prevented visible growth after incubation. Each assay was performed at least three times on separate days.

LpxC Enzyme Assays. Protein Expression. E. coli LpxC C63A. The expression of *E. coli* LpxC C63A was performed essentially as previously described.⁷⁴ The C63A mutation lowers the undesired influence of the Zn^{2+} -concentration on the enzymatic activity.⁵⁰

The plasmid pET11EcLpxCC63A, which was kindly provided by Carol Fierke, ⁵⁰ was transformed into *E. coli* BL21(DE3) cells. The overnight culture was prepared by growing a single colony in 50 mL of LB supplemented with carbenicillin (0.1 mM) and glucose (0.5%) at 37 °C and 200 rpm. The next day, 2 mL of this culture was used to inoculate 400 mL of fresh LB containing carbenicillin (0.1 mM) and glucose (0.5%). After reaching an OD₆₀₀ of 0.6–0.8, the culture was cooled to 30 °C and induced with isopropyl β -D-1-thiogalactopyranoside (IPTG, 1 mM) and ZnCl₂ (100 μ M). After being grown for an additional 4 h at 30 °C, the cells were cooled on ice for 20 min and then harvested by centrifugation (4 °C, 5000g, 15 min) and stored at –20 °C.

P. aeruginosa LpxC. The expression of *P. aeruginosa* LpxC was based on the protocol for the expression of *E. coli* LpxC C63A. The plasmid pWY427, which was kindly provided by Ning Gao,²⁰ was transformed into *E. coli* BL21(DE3) cells. The overnight culture was prepared by growing a single colony in 50 mL of LB supplemented with kanamycin (0.1 mM) and glucose (0.5%) at 37 °C and 200 rpm. The next day, 2 mL of this culture was used to inoculate 400 mL of

fresh LB containing kanamycin (0.1 mM) and glucose (0.5%). After reaching an OD₆₀₀ of 0.3–0.4, the culture was induced with IPTG (500 μ M) and ZnCl₂ (100 μ M). After being grown for an additional 2 h at 37 °C, the cells were cooled on ice for 20 min and then harvested by centrifugation (4 °C, 5000g, 15 min) and stored at –20 °C.

Protein Purification. Unless otherwise specified, all steps were carried out at 4 $^{\circ}$ C.

E. coli LpxC C63A. The harvested cells were thawed on ice and resuspended in 50 mL of anion exchange (AEX) buffer [25 mM HEPES (pH = 7.0), 2 mM dithiothreitol (DTT)], containing benzamidine (15 μ g mL⁻¹) and phenylmethylsulfonyl fluoride (PMSF, 1 mM) as protease inhibitors. Afterward, the cells were disrupted by sonication (5 × 40 s). Then, the cellular debris were removed by centrifugation (4 °C, 5000g, 90 min), and the supernatant was filtered (0.2 μ m).

The cleared lysate was loaded onto a 20 mL AEX column (HiPrep Q HP 16/10, GE Healthcare) and eluted at a flow rate of 0.5 mL min⁻¹ using a linear potassium chloride gradient (0 M \rightarrow 0.5 M) in AEX-buffer. The fractions containing LpxC were concentrated using molecular weight cut off (MWCO) spin columns (10 kDa), loaded onto a 120 mL size exclusion (SEC) column (HiLoad 16/600 Superdex 200, GE Healthcare) and eluted at a flow rate of 0.5 mL min⁻¹ in SEC buffer [50 mM Bis/Tris (pH = 6.0), 150 mM NaCl].

P. aeruginosa LpxC. The harvested cells were thawed on ice and resuspended in 50 mL of AEX buffer [25 mM Tris–HCl (pH = 8.0), 2 mM DTT, 5% glycerol] containing benzamidine (15 μ g mL⁻¹) and PMSF (1 mM) as protease inhibitors. Afterward, the cells were disrupted by sonication (5× 40 s). Then, cellular debris were removed by centrifugation (4 °C, 5000g, 90 min), and the supernatant was filtered (0.2 μ m).

The cleared lysate was loaded onto a 20 mL AEX column (HiPrep Q HP 16/10, GE Healthcare) and eluted at a flow rate of 0.5 mL min⁻¹ using a linear sodium chloride gradient (0 M \rightarrow 0.5 M) in AEX buffer. The fractions containing LpxC were concentrated using MWCO spin columns (10 kDa), loaded onto a 120 mL SEC column (HiLoad 16/600 Superdex 200, GE Healthcare) and eluted at a flow rate of 0.5 mL min⁻¹ in SEC buffer [25 mM HEPES (pH = 8.0), 2 mM DTT, 5% glycerol].

The presence of the enzyme during the purification process was confirmed by sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) with Coomassie brilliant blue staining. The purified enzyme was quantified by use of a Nanodrop 2000C, diluted with SEC buffer to 0.5 mg mL⁻¹ and stored at -80 °C.

Enzyme Inhibition Assays. E. coli LpxC C63A. A fluorescencebased microplate assay for LpxC activity was performed as described by Clements et al.48 The wells in a black, nonbinding, 96-well fluorescence microplate (Greiner Bio One, Frickenhausen) were filled with 93 μ L of 26.9 μ M UDP-3-O-[(R)-3-hydroxymyristoyl]-Nacetylglucosamine in assay buffer [40 mM sodium morpholinoethanesulfonic acid (pH 6.0), 80 µM dithiothreitol, 0.02% Brij 35]. In order to assay the inhibitors at final concentrations from 20 nM up to 20 μ M, 2 μ L of a respective dilution of the compounds in DMSO were added. The addition of 5 μ L of a solution of purified LpxC (10 $\mu g \text{ mL}^{-1}$) in assay buffer led to final concentrations of 25 μM UDP-3-O-[(R)-3-hydroxymyristoyl]-N-acetylglucosamine, 15 nM E. coli LpxC C63A, 2% DMSO, and from 20 nM up to 20 μ M inhibitor. The microplate was incubated for 30 min at 37 °C in a plate shaker. Then, the biochemical reaction was stopped by adding 40 μ L of 0.625 M sodium hydroxide. The reaction mixture was further incubated for 10 min and neutralized by adding 40 μL of 0.625 M acetic acid. The deacetylated product UDP-3-O-[(R)-3-hydroxymyristoyl]glucosamine was converted into a fluorescing isoindole by adding 120 μ L of an *o*-phthaldialdehyde-2-mercaptoethanol solution, which was prepared by dissolving 10 mg of o-phthaldialdehyde in 1 mL of methanol, diluting the mixture with 24 mL of a sodium borate buffer (0.1 M), and finally adding 2.5 μ L of 2-mercaptoethanol.⁴⁹ Fluorescence was measured with a TriStar² S LB 942 plate reader (Berthold, Bad Wildbad) at 340 nm excitation and 460 nm emission wavelengths. Each assay was performed at least three times on separate days. The IC₅₀ values were calculated via Probit-log

concentration graphs with the aid of the software Origin and were subsequently converted into K_i values using the Cheng–Prusoff equation.^{75,76} The K_M value of *E. coli* LpxC C63A was determined experimentally using the LC–MS/MS-based LpxC assay (Supporting Information) and was found to be 3.6 μ M (Figure S8).

P. aeruginosa LpxC. The protocol of the LC-MS/MS-based *P. aeruginosa* LpxC assay was based on the *E. coli* LpxC C63A enzyme assay. Compared to the fluorescence-based enzyme assay, the substrate was diluted 1:10 and the inhibitors were diluted 1:4.

The wells in a black, nonbinding, 96-well fluorescence microplate (Greiner Bio One, Frickenhausen) were filled with 93 μ L of 2.69 μ M UDP-3-O-[(*R*)-3-hydroxymyristoyl]-*N*-acetylglucosamine in assay buffer [50 mM KH₂PO₄/K₂HPO₄ (pH = 7.5), 80 μ M dithiothreitol, 0.02% Brij 35]. In order to assay the inhibitors at final concentrations from 5 nM up to 5 μ M, 2 μ L of a respective dilution of the compounds in DMSO were added. The addition of 5 μ L of a solution of purified *P. aeruginosa* LpxC (5 μ g mL⁻¹) in assay buffer led to final concentrations of 2.5 μ M UDP-3-O-[(*R*)-3-hydroxymyristoyl]-*N*-acetylglucosamine, 7.5 nM *P. aeruginosa* LpxC, 2% DMSO, and from 5 nM up to 5 μ M inhibitor. The microplate was incubated for 30 min at 37 °C in a plate shaker. Then, the biochemical reaction was stopped by adding 40 μ L of 0.625 M hydrochloric acid. The reaction mixtures were further incubated for 10 min, sealed, and stored at -80 °C until analysis.

LC–MS/MS-Analysis. The reaction mixtures were separated by ultrahigh performance liquid-chromatography (1290 II Infinity UHPLC, Agilent Technologies), and the eluted compounds were analyzed by mass spectrometry using electrospray ionization in negative ion mode with a triple quadrupole linear ion trap mass spectrometer (QTRAP 5500, AB Sciex LLC).

UHPLC method: column: Nucleodur C18 Gravity-SB ($\emptyset = 3 \text{ mm}$, h = 100 mm, Macherey-Nagel), coupled to a Universal RP-guard column ($\emptyset = 2 \text{ mm}$, h = 4 mm, Macherey-Nagel); flow rate: 0.3 mL · min⁻¹; injection volume: 3.0 μ L; solvents: (A) 20 mM ammonium formate in water; (B) 1 mM ammonium formate in acetonitrile/ isopropanol/water (47.5:42.75:9.75); gradient elution: (B %): 0–1 min: 30%, 1–16 min: gradient from 30 to 90%, 16–17 min: 90%, 17–17.5 min: gradient from 90 to 30%, 17.5–21.5 min: 30%; detection: 12–19 min; $t_{\rm R}$ (1) = 12.2 min, $t_{\rm R}$ (2) = 13.0 min.

To analyze the eluted compounds by mass spectrometry, a MRM method was applied. The specific parameters of this method are given in Table S2 (Supporting Information). After detection and selection of the precursor ions (1: m/z 832; 2: m/z 790), both analytes were fragmented, leading to three identical product ions [m/z (product 1) 385, collision energy = -60 V; m/z (product 2) 159, collision energy = -80 V; m/z (product 3) 79, collision energy = -140 V]. The mass transitions 832 \rightarrow 79 (substrate 1) and 790 \rightarrow 79 (product 2) were used as quantifiers; the other mass transitions were used as qualifiers. The ratio between substrate 1 and product 2 was quantified by comparing the peak areas of the quantifiers. The percentual inhibition caused by each inhibitor concentration was determined with respect to the amount of the product formed in the noninhibited reaction after 30 min.

Each assay was performed at least two times on separate days. The IC₅₀ values were calculated via Probit-log concentration graphs with the aid of the software Origin and were subsequently converted into K_i values using the Cheng–Prusoff equation.^{75,76} The K_M value for the *P. aeruginosa* LpxC-catalyzed deacetylation of 1 was determined experimentally (Supporting Information) and found to be 4.7 μ M (Figure S9).

Assays to Determine the In Vitro Inhibition of LasB, MMPs, and TACE. Purification of LasB from *P. aeruginosa* PA14 supernatant and the subsequent performance of the FRET-based in vitro inhibition assay was performed as described previously.⁶⁶ The TACE (ADAM-17) inhibitor screening kit was purchased from Sigma-Aldrich (Saint Louis, MO). MMPs 1–3 along with the SensoLyte 520 Generic MMP Activity Kit Fluorimetric were purchased from AnaSpec (Fremont, CA, USA). The assays were performed according to the guidelines of the respective manufacturer. Fluorescence signals were measured using a CLARIOstar plate reader (BMG Labtech, Ortenberg, Germany).

Cytotoxicity Assay. An MTT-based assay was employed to evaluate the viability of HepG2 cells after challenge with selected inhibitors and performed as described previously.⁷⁷

Kinetic Turbidimetric Solubility. The desired compounds were sequentially diluted in DMSO in a 96-well plate. 1.5 μ L from each well was transferred into another 96-well plate and mixed with 148.5 μ L of PBS. Plates were shaken for 5 min at 600 rpm at room temperature, and the absorbance at 620 nm was measured. Absorbance values were normalized by blank subtraction and plotted using GraphPad Prism 8.4.2 (GraphPad Software, San Diego, CA, USA). Solubility (*S*) was determined based on the First X value of AUC function using a threshold of 0.005.

Log $D_{7,4}$. Log $D_{7,4}$ was analyzed using an HPLC-based method. The UV retention time of reference compounds with known log $D_{7,4}$ was determined and plotted toward their log $D_{7,4}$. Linear regression was used to determine the log $D_{7,4}$ of unknown compounds. Analysis was performed using a Vanquish Flex HPLC system with a variable wavelength detector (Thermo Fisher, Dreieich, Germany) with the following conditions: EC150/2 NUCLEODUR C18 Pyramid column, 5 μ M (Macherey Nagel, Düren, Germany); eluent A: 50 mM NH₄OAc pH 7.4, eluent B: acetonitrile, and flow: 0.6 mL/min; gradient elution: (B %): 0–2.5 min: gradient from 0 to 100%, 2.5–3.0 min: 100%, 3.0–3.2 min: gradient from 100 to 0%, 3.2–5.0 min: 0%.

ADME In Vitro Studies. The microsomal metabolic stability assay as well as the plasma protein binding assay were conducted as described previously.⁷⁸

HPLC–MS/MS Analysis. Samples were analyzed using an Agilent 1290 Infinity II HPLC system coupled to an AB Sciex QTrap 6500plus mass spectrometer. LC conditions were as follows: column: Agilent Zorbax Eclipse Plus C18, 50×2.1 mm, 1.8μ m; temperature: 30 °C; injection volume: 5μ L per sample; flow rate: 700 μ L min⁻¹. Samples were run under acidic conditions. Solvents: (A) water + 0.1% formic acid; (B) 95% acetonitrile/5% H₂O + 0.1% formic acid. Gradient elution: (A %): 0–0.1 min: 99%, 0.1–3.5 min: gradient from 99 to 50%, 3.5–3.8 min: gradient from 50 to 0%, 3.8–4.7 min: gradient from 0 to 99%. Mass transitions for controls and compounds are depicted in Table S3.

Metabolic Stability in Liver S9 Fractions. For the evaluation of combined phase I and phase II metabolic stability, the compound (1 μ M) was incubated with 1 mg/mL pooled mouse liver S9 fraction (Xenotech, Kansas City, USA), 2 mM nicotinamide adenine dinucleotide phosphate hydrogen (NADPH), 1 mM UDPGA, 10 mM MgCl₂, 5 mM glutathione (GSH), and 0.1 mM 3'phosphoadenosine 5'-phosphosulfate (PAPS) at 37 °C for 120 min. The metabolic stability of testosterone, verapamil, and ketoconazole were determined in parallel to confirm the enzymatic activity of mouse S9 fractions. The incubation was stopped after defined time points by precipitation of aliquots of S9 enzymes with 2 volumes of cold acetonitrile containing internal standard (150 nM diphenhydramine). Samples were stored on ice until the end of the incubation and the precipitated protein was removed by centrifugation (15 min, 4 °C, 4000g). Concentration of the remaining test compound at the different time points was analyzed by HPLC-MS/MS (TSQ Quantum Access Max, Thermo Fisher, Dreieich, Germany) and used to determine half-life $(t_{1/2})$.

Stability in Mouse Plasma. To determine stability in mouse plasma, the compound $(1 \ \mu M)$ was incubated with pooled CD-1 mouse plasma (Neo Biotech, Nanterre, France). Samples were taken at defined time points by mixing aliquots with 4 volumes of acetonitrile containing internal standard (125 nM diphenhydramine). Samples were stored on ice until the end of the incubation, and the precipitated protein was removed by centrifugation (15 min, 4 °C, 4000g, 2 centrifugation steps). Concentration of the remaining test compound at the different time points was analyzed by HPLC–MS/MS (TSQ Quantum Access MAX, Thermo Fisher, Dreieich, Germany). The plasma stability of procain, propantheline, and diltiazem were determined in parallel to confirm enzymatic activity.

Computational Methods. Molecular docking was performed using a recently developed and evaluated protocol as reported in our previous studies.^{32,79} This docking protocol was successful in redocking the cocrystallized inhibitors. The crystal structure of E. coli LpxC (PDB ID: 4MQY) in complex with LPC-138 (Figure S1) was retrieved from the Protein Data Bank (https://www.rcsb.org). The LpxC protein structure was chosen due to the similarity of LPC-138 to the compounds developed in the current work. The LpxC protein was prepared using Protein Preparation Wizard by adding the hydrogen atoms and missing side chains in the Schrödinger suite.⁸¹ Solvent molecules were removed. Tautomeric states and protonation states of the amino acids were adjusted with the PROPKA tool at pH 7.0. OPLS3e force field was applied to minimize the complex to remove the steric clashes, bad contacts, and unsuitable torsional angles. The inhibitor structures were prepared using the Ligprep tool by applying the OPLS3e force field. Subsequently, 64 conformers per ligand were generated using the Confgen tool with force field minimization on output conformers. Molecular docking studies were performed in Glide from the Schrödinger suite. Grid files were prepared with default settings by applying box-size as 15 Å \times 15 Å \times 15 Å. Standard Precision mode with flexible ligand sampling and enhanced planarity of conjugated π groups were used for docking studies. The validation of the docking protocol was done by redocking. The root-mean-square deviation (RMSD) value of the redocked ligand from 4MQY compared to its observed binding mode in the crystal structure was 0.78 Å. The docking results were visually analyzed in MOE.82

MD Simulation. The selected docking poses in complex with LpxC as well as the original crystal structure cocrystallized with the inhibitor LPC-138 (PDB ID: 4MQY) were subjected to MD simulation using Desmond (Schrödinger Suite 2019).^{82,83} LpxC-inhibitor complexes were simulated for two independent runs each of 50 ns. The System Builder panel was used to build the complexes. The complexes were solvated using a SPC water model and an orthorhombic box with 10 Å distance between the solute structures and the simulation box boundary. The box volume was then minimized. The whole system was neutralized by adding sodium ions that were placed 10 Å away from the inhibitor structure. The MD panel in Desmond was used to set the simulation parameters. The prepared system was relaxed using the default Desmond relaxation protocol for NPT ensemble followed by a production run utilizing the NPT ensemble at the temperature of 300 K using a Nosé-Hoover chain thermostat and a pressure of 1.01325 bar using a Martyna-Tobias-Klein barostat. The progress of the simulation was recorded every 100 ps. The Simulation Interaction Diagram panel was used for analyzing the RMSD, RMSF, and the interaction persistence of the ligands. The Simulation Event Analysis panel was used for calculating protein-inhibitor distances and hydrogen bond occupancies.

ASSOCIATED CONTENT

Supporting Information

The Supporting Information is available free of charge at https://pubs.acs.org/doi/10.1021/acs.jmedchem.4c01262.

Molecular formula strings (CSV)

Determination of enzymatic parameters, synthetic procedures and analytical data of compounds (S)-38b-n, (R)-38b-h,j,m,n, (S)-39b-n, (R)-39-h,j,m,n, (S)-13b-n, and (R)-13b-h,j,m,n, ¹H and ¹³C NMR spectra of representative compounds, and HPLC chromatograms of the test compounds (PDF)

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Notes

The authors declare no competing financial interest.

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ABBREVIATIONS

ILOE, interligand nuclear Overhauser effect; LB, lysogeny broth; MRM, multiple reaction monitoring; RMSF, root-meansquare fluctuation; STD, saturation-transfer difference; TACE, tumor necrosis factor- α converting enzyme; UDPGA, uridine diphosphate glucuronic acid; WaterLOGSY, Water-Ligand Observed via Gradient SpectroscopY

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