

Lepadins D–F: Antiplasmodial and Antitrypanosomal Decahydroquinoline Derivatives from the Tropical Marine Tunicate *Didemnum* sp.

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From a new tunicate species, belonging to the genus *Didemnum*, three alkaloids possessing an unusual and extremely rare decahydroquinoline skeleton and showing significant and selective antiplasmodial and antitrypanosomal activity were obtained as follows: (2*R**,3*S**,4*aR**,5*R**,8*aS**)-decahydro-3-hydroxy-5-(5'-hydroxyoctyl)-2-methylquinoline (lepadin D, **1**), its quaternary nitrogen derivative (**2**), (2*R**,2''*E*,3*S**,4*aR**,5*R**,8*aS**)-decahydro-3-hydroxy-5-(5'-hydroxyoctyl)-2-methyl-3-quinolinyl ester 2''-octenoic acid (lepadin E, **3**), and (2*S**,2''*E*,3*S**,4*aR**,5*R**,8*aS**)-decahydro-3-hydroxy-5-(5'-hydroxyoctyl)-2-methyl-3-quinolinyl ester 2''-octenoic acid (lepadin F, **4**). These isolates may well serve as lead structures for the development of new antimalarial drugs.

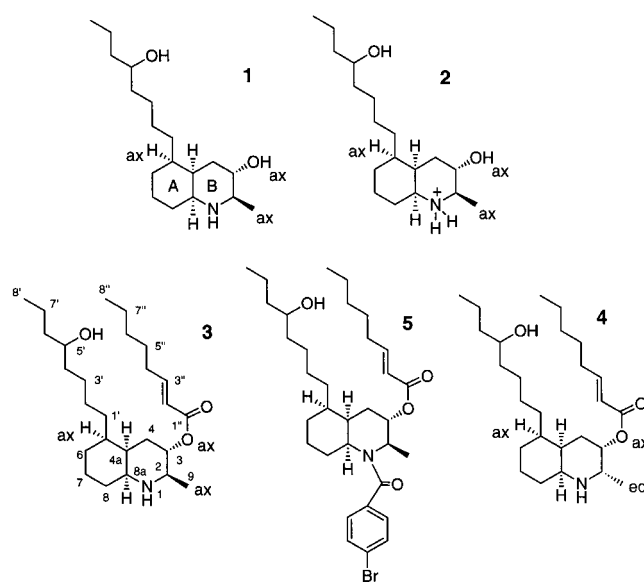
Introduction

The marine environment is still an unexploited resource in terms of antiparasitic agents or lead structures. Some sponge, tunicate, and marine fungal metabolites have, however, clearly demonstrated the potential of marine natural products in this area.¹ Tunicates of the genus *Didemnum* have yielded a variety of secondary metabolites including peptides,^{2–4} unusual nine-membered ring lactones,⁵ β -carboline,^{6,7} and indole-derived alkaloids,⁸ lamellarines,⁹ didemnenones,¹⁰ enterocin derivatives,¹¹ pyrroloacridine alkaloids such as plakinidine D,¹² small iodated aromatic compounds,^{12,13} and long chain amino alcohols¹⁴ and acids,¹⁵ many of which are halogenated. Several of these compounds were found to have interesting biological activities, e.g., mollamide,⁴ cyclodidemnoamide,² the didemnenones,¹⁰ and plakinidine D¹² are all cytotoxic against various cancer cell lines, while enterocin¹¹ and the didemnenones¹⁰ are antimicrobial. In the current project, the antiparasitic activity of *Didemnum*-derived secondary metabolites was investigated. These efforts resulted in the isolation of three new unusual alkaloids (lepadins D–F, **1**, **3**, and **4**) (Chart 1), with antiplasmodial and antitrypanosomal activity, from a new *Didemnum* species collected from Stanley Reef, the Great Barrier Reef. The new compounds are structurally related to the lepadins found in *Clavelina lepadiformis* collected in the North Sea^{16,17} and, together with these three compounds,^{16,17} are the only decahydroquinoline derivatives known from marine sources.

Results and Discussion

Structural Chemistry. Mass spectrometric (MS) analysis of **1** indicated it to have the molecular formula

Chart 1



$C_{18}H_{35}NO_2$. Because there were no multiple bonds present within the molecule, as deduced from the nuclear magnetic resonance (NMR) data, it was evident that the two elements of unsaturation indicated by the molecular formula of **1** must be due to the presence of two rings. From the ¹H and ¹³C NMR data of **1**, it was possible to account for the presence of $C_{18}H_{32}$ ($2 \times CH_3$, $10 \times CH_2$, and $6 \times CH$) within the molecule. According to the molecular formula of **1**, this left one nitrogen, two oxygen, and three hydrogen atoms to be accounted for. IR data of **1** showed it to contain OH and/or NH functionality (ν 3290 cm^{-1}). As the ¹³C NMR data evidenced the presence of two carbons with oxygen (δ 71.2 d, 72.2 d) and two with nitrogen attached (δ 54.0 d, 56.2 d), it was clear that the unaccounted for H_3NO_2 part of **1** was present in the form of two secondary hydroxyl groups and a secondary amine. After all protons had been assigned to their directly bound carbon partner, from the results of a two dimensional (2D) ¹H–

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Table 1. ¹H NMR Data for Compounds **1** (300 MHz, CD₃OD), **2** (400 MHz, CD₃OD), **3** (300 MHz, CDCl₃), and **4** (300 MHz, CDCl₃)^a

carbon	1	2	3	4
2	2.99 (dq, <i>J</i> = 4.1, 7.0)	3.46 (dq, <i>J</i> = 2.5, 7.0 Hz)	3.08 m	3.09 (dq, <i>J</i> = 1.9, 6.4 Hz)
3	3.70 (ddd, <i>J</i> = 4.1, 4.3, 4.4 Hz)	3.93 (ddd, <i>J</i> = 2.5, 3.1, 3.1 Hz)	4.78 m	4.92 brs
4	1.50 m, 1.87 (ddd, <i>J</i> = 4.4, 14.0, 14.0 Hz)	1.63 (brd, <i>J</i> = 14.2 Hz), 1.92 (ddd, <i>J</i> = 3.1, 14.2, 14.2 Hz)	1.52 m, 1.89 (ddd, <i>J</i> = 4.5, 11.7, 14.3 Hz)	1.62 (ddd, <i>J</i> = 2.1, 12.3, 12.3 Hz), 1.72 (brd, <i>J</i> = 12.3 Hz)
4a	2.38 m	2.60 m	2.17 m	2.05 m
5	1.53 m	1.59 m	1.47 m	1.42 m
6	1.12 (dddd, <i>J</i> = 3.3, 12.5, 12.5, 12.5 Hz), 1.38 m	1.13 (dddd, <i>J</i> = 3.1, 12.6, 12.7, 12.7 Hz), 1.32 m	1.09 m, 1.38 m	0.93 m, 1.36 m
7	1.79 m	1.89 m	1.13 m, 1.66 m	1.22 m, 1.31 m
8	1.66 (dddd, <i>J</i> = 3.8, 12.2, 12.5, 12.5 Hz), 1.72 m	1.84 m	1.55 m, 1.68 m	1.45 m, 1.78 m
8a	3.03 (ddd, <i>J</i> = 4.9, 6.0, 12.2 Hz)	3.48 m	2.95 m	2.89 (ddd, <i>J</i> = 4.1, 4.1, 12.2 Hz)
9	1.28 (d, <i>J</i> = 7.0 Hz)	1.48 (d, <i>J</i> = 7.1 Hz)	1.19 (d, <i>J</i> = 6.8 Hz)	1.01 (d, <i>J</i> = 6.4 Hz)
1'	1.34 m	1.34 m	1.26 m	1.15 m
2'	1.50 m	1.32 m	1.26 m	1.48 m
3'	1.47 m	1.34 m	1.46 m	1.19 m
4'	1.49 m	1.43 m	1.40 m	1.37 m
5'	3.56 m	3.57 m	3.56 m	3.54 m
6'	1.44 m	1.41 m	1.40 m	1.39 m
7'	1.51 m, 1.40 m	1.46 m	1.39 m	1.38 m
8'	0.97 (t, <i>J</i> = 7.0 Hz)	0.94 (t, <i>J</i> = 6.8 Hz)	0.90 (t, <i>J</i> = 7.0 Hz)	0.9 (t, <i>J</i> = 7.1 Hz)
2''			5.83 (d, <i>J</i> = 15.4 Hz)	5.89 (d, <i>J</i> = 15.4 Hz)
3''			6.99 (ddd, <i>J</i> = 6.8, 6.8, 15.4 Hz)	7.01 (ddd, <i>J</i> = 6.8, 6.8, 15.4 Hz)
4''			2.20 m	2.20 m
5''			1.26 m, 1.40 m	1.20 m
6''			1.29 m	1.31 m
7''			1.31 m	1.30 m
8''			0.90 (t, <i>J</i> = 6.8 Hz)	0.90 (t, <i>J</i> = 7.0 Hz)

^a All assignments are based on extensive 1D and 2D NMR measurements including COSY, HMQC, and HMBC.

¹³C heteronuclear multiple quantum coherence (HMQC) experiment, most of the planar structure of the molecule could be deduced from analysis of its ¹H–¹H correlation spectroscopy (COSY) data. Thus, starting with H₃–9, a continuous chain of ¹H–¹H coupling, via H–4a to H–5, H–5 to H₂–6, H₂–6 to H₂–7, H₂–7 to H₂–8, and then H₂–8 to H–8a, which further coupled to H–4a, was deduced and enabled ring A within the molecule to be completed. The ¹H–¹H spin system could be further extended by following the chain of proton couplings from H–5 through to H₃–8'. On the basis of the ¹H and ¹³C NMR chemical shifts of the resonances associated with CH–3 (δ 3.70 [ddd, *J* = 4.1, 4.3, 4.4 Hz] and 71.2 d) and CH–5' (δ 3.56 m and 72.2 d), it was evident that the two hydroxyl functions must reside at these positions, leaving the NH function to be placed between C–2 and C–8a, and in so doing complete the second ring (ring B) within **1** and its planar structure. The relative stereochemistry of the decahydroquinoline part of the molecule was determined mainly through analysis of ¹H–¹H coupling constants, and the results of nuclear Overhauser effect (NOE) difference experiments. As all of the ¹H–¹H coupling constants associated with H–3 were less than 5 Hz, it had to have an equatorial, and the 3-OH an axial, orientation. The coupling constants associated with the axially oriented protons at C–6 (*J* = 3.3, 12.5, 12.5, and 12.5 Hz) and C–8 (*J* = 3.8, 12.2, 12.5, and 12.5 Hz) clearly showed H–5 and H–8a (when considered part of the A ring) also to be axially oriented. NOE measurements made with **1** showed interactions to occur between H–4a and H–8a and H–5 and between H–8a and H–5 and H–7_{ax} and revealed H–4a, H–5, and H–8a to be on the same side of the molecule, the A ring to have a chair conformation, and rings A and B to be cis-fused, a deduction also supported by the absence of Bohlman

bands in the IR spectrum of **1**.¹⁸ With the conformation of the A ring fixed, the conformation of ring B was also deduced to be a chair on the basis of the fact that the 3-OH group had an axial orientation; H–4 demonstrated a trans-diaxial coupling with H–4a (14.0 Hz) and an axial–equatorial coupling with H–3. The orientation of the methyl group at C–2 was deduced as being axial on the basis of the relatively small ¹H–¹H coupling constant between H–2 and H–3 (4.1 Hz) and the clear NOE interactions between CH₃–9 and H–4_{ax} and H₂–8. Attempts to resolve the configuration at C–5' were unsuccessful. For **1**, (2*R**,3*S**,4*aR**,5*R**,8*aS**)-decahydro-3-hydroxy-5-(5'-hydroxyoctyl)-2-methylquinoline, which is similar in structure to lepadins A–C,^{16,17} the trivial name of lepadin D, is proposed.

The electron impact (EI) MS data of **2** were almost the same as those for **1**, which indicated that the two compounds were quite similar. Close examination of the ¹H and ¹³C NMR data sets (see Tables 1 and 2) for both compounds supported this deduction and showed the major differences between the two to arise from a change in the B ring. The only alteration in this part of the molecule that was consistent with all of the spectroscopic data was the presence of a quaternary nitrogen between C–2 and C–8a. All of the remaining spectroscopic data for **2**, particularly NOE difference and ROESY measurements, that showed NOE interactions to occur between H–8a and H–4a and H–5 and between CH₃–9 and H–4_{ax} and H₂–8 revealed it to have the same relative configuration as **1** and thus be best described as (2*R**,3*S**,4*aR**,5*R**,8*aS**)-decahydro-3-hydroxy-5-(5'-hydroxyoctyl)-2-methylquinolinium. Clearly, **2** can be generated by the action of dilute acid on **1** (as was treated with 0.1 M HCl, and **2** was indeed produced). As the work up of all of the material associated with this

Table 2. ^{13}C NMR Data for Compounds **1** (75.5 MHz, CD_3OD), **2** (100 MHz, CD_3OD), **3** (75.5 MHz, CDCl_3), and **4** (75.5 MHz, CDCl_3)^a

carbon	1	2	3	4
2	54.0 d	55.5 d	50.8 d	47.1 d
3	71.2 d	68.0 d	73.4 d	71.3 d
4	24.7 t	21.4 t	22.9 t	23.7 t
4a	33.3 d	31.1 d	33.9 d	33.0 d
5	40.4 d	40.2 d	39.0 d	39.5 d
6	28.3 t	27.3 t	27.5 t	25.8 t
7	25.7 t	25.9 t	23.8 t	26.6 t
8	30.3 t	26.9 t	31.1 t	25.4 t
8a	56.2 d	57.1 d	54.7 d	55.4 d
9	19.6 q	17.4 q	20.7 q	18.4 q
1'	34.4 t	34.0 t	32.9 t	33.1 t
2'	27.0 t	27.6 t	27.3 t	27.7 t
3'	28.3 t	28.0 t	27.7 t	27.0 t
4'	38.4 t	38.3 t	37.5 t	37.5 t
5'	72.2 d ^b	72.1 d	71.6 d	71.6 d
6'	40.8 t	40.8 t	39.8 t	39.8 t
7'	19.9 t	19.9 t	18.9 t	18.9 t
8'	14.5 q	14.5 q	14.0 q	14.2 q
1''			166.0 s	166.6 s
2''			121.4 d	121.2 d
3''			150.0 d	149.9 d
4''			32.2 t	32.2 t
5''			25.8 t	25.2 t
6''			31.4 t	31.4 t
7''			22.5 t	22.5 t
8''			14.2 q	14.0 q

^a All assignments are based on extensive 1D and 2D NMR measurements including COSY and HMQC. ^b Implied multiplicity by DEPT (C = s, CH = d, CH_2 = t, and CH_3 = q).

tunicate was strictly neutral, however, the likelihood that **2** was an artifact of isolation can be ruled out. It should also be noted that **2** was the only quaternary nitrogen compound detected in this study. As **2** was generated from the action of dilute HCl on **1**, it is not unlikely that the counter ion present in **2** is Cl^- , a deduction that is supported, to some extent, by the chemical ionization (CI) MS of **2**.

Compound **3** was analyzed for $\text{C}_{26}\text{H}_{47}\text{NO}_3$ by HREIMS. Of the four degrees of unsaturation implied by the molecular formula of **3**, two could be attributed to double bonds; a carbon-carbon double bond (δ [5.83 {d, $J = 15.4$ Hz}], 6.99 {ddd, $J = 6.8, 6.8, 15.4$ Hz}], 121.4 d, 150.0 d) was detected, which is part of an α,β unsaturated ester carbonyl group (δ 166.0 s; 264 nm [ϵ 2730]; 1715 cm^{-1}). As there were no other multiple bonds within the molecule, the two remaining elements of unsaturation had to be present in the form of two rings. From the ^1H and ^{13}C NMR data (see Tables 1 and 2) of **3**, it was possible to deduce that the only difference between it and **1** was the substitution of the secondary hydroxyl function at C-3 in **1** with a *2E*-octenoic acid ester function. This deduction was supported by all of the spectroscopic data for **3**. The relative configuration of **3** was deduced to be the same as that of **1** based on ^1H - ^1H coupling constant analysis and NOE difference measurements, which showed it to be (*2R**, *2''E*, *3S**, *4aR**, *5R**, *8aS**)-decahydro-3-hydroxy-5-(5'-hydroxyoctyl)-2-methyl-3-quinolinyl ester *2''*-octenoic acid, for which the trivial name lepadin E is proposed.

MS analysis of **4** showed it to have the identical molecular formula as **3**. All of its other spectroscopic data were also extremely similar to those found for lepadin E (**3**). Interpretation of the HMBC, HMQC, and ^1H - ^1H COSY spectra of **4** confirmed it to be a decahydroquinoline with the same substitution pattern as

that found in **3**; compounds **3** and **4** must thus be stereoisomers. The ^1H and ^{13}C NMR chemical shifts associated with CH-2, CH-3, CH_2 -8, and CH_3 -9 for **4** differ significantly from those of **3**. When the locations of these atoms within the molecule are considered, it is evident that the only change within the molecule that can be made to accommodate all of these differences, without significantly affecting ^1H - ^1H coupling constants, is inversion of the configuration at C-2. The results of NOE difference measurements made with **4**, in particular interactions between H-2 and H-4_{ax} and H-8_{ax}, between H-8a and H-4a and H-5, and between CH_3 -9 and H-2'' and H-3'', were also consistent with this deduction and showed the molecule to be best described as (*2S**, *2''E*, *3S**, *4aR**, *5R**, *8aS**)-decahydro-3-hydroxy-5-(5'-hydroxyoctyl)-2-methyl-3-quinolinyl ester *2''*-octenoic acid (lepadin F).

Attempts were made to prepare some para-bromobenzoate derivatives of the new lepadins in order to determine the absolute configuration for at least one of the new compounds. In each case, unfortunately, these reactions yielded only amides of the type shown by **5**, which were not crystalline. Esterification at C-5' was never observed under various reaction conditions.

Compounds **1**-**4** are extremely rare decahydroquinoline derivatives. Among marine natural products, this structural class is only known from the flat worm *Prosthecergeus villatus* and its prey, the tunicate *Clavelina lepadiformis*.^{16,17} In the terrestrial environment, poisonous frogs of the genus *Dendrobates* produce simple decahydroquinolines.¹⁹ Among plants, *Lycopodium* species are known to produce decahydroquinolines but as parts of more complex alkaloidal molecules.²⁰

Biological Activity. The isolated compounds were tested in several bioassays for their biological activity (Table 3). All compounds, except **2**, had weak antifungal activity against *Ustilago violacea* and/or *Eurotium repens* and several other microorganisms. Compounds **3** and **4** showed moderate inhibition of the enzyme tyrosine kinase p56^{lck} (TK). Antiplasmodial (*Plasmodium falciparum*) and antitrypanosomal (*Trypanosoma brucei rhodesiense*, *Trypanosoma cruzi*) activities were assessed for all compounds, and found to be significant and structure-related. Antiplasmodial and antitrypanosomal effects increase significantly from **1** to **4** (e.g., against *P. falciparum* clone K1, **1** has an IC_{50} of 6169 ng/mL [20 μmol] as compared to 208 ng/mL [0.4 μmol] for **4**). This 50-fold activity increase clearly results from the presence of the *2E*-octenoic acid ester function at C-3 in **3**-**5** in place of the secondary hydroxyl function found in **1** and **2**. Antiplasmodial activity also increases in going from **3** to **4**. As the only difference between these two molecules is the configuration at C-2, it must be concluded that the configuration at this center plays a significant role in determining the activity of the tested molecules. The mechanism of action of these compounds is unknown and needs to be investigated to see if it is similar to that of the fully aromatic quinoline type compounds, e.g., chloroquine.²¹ The conformationally mobile side chain at C-3 probably serves to stabilize any nonbonding interactions between the test compounds and a "receptor" molecule, which,

Table 3. Biological Activities of Compounds 1–5

substance tested	<i>Ustilago violacea</i> ^a	<i>Eurotium repens</i> ^a	tyrosine kinase ^b	<i>Plasmodium falciparum</i> , IC ₅₀ μg/mL		<i>Trypanosoma cruzi</i> , IC ₅₀ μg/mL	<i>Trypanosoma rhodesiense</i> , IC ₅₀ μg/mL	cytotoxicity ^c , IC ₅₀ μg/mL
				strain K1	strain NF54			
chloroquine				0.091	0.005			
qinghaosu				0.001	0.003			
melarsoprol						0.6	0.0026	
miconazol	27	20						
piceatannol			100					
1	2	1	NA	6.1(20)	10.0(33)	37.2	5.60	>30
2	NA ^d	NA	NA	4.0(12) ^e	10.0(33) ^e	37.3	1.97	>30
3	4	4	65	0.4(0.9)	0.9(2)	2.2	0.38	16.2
4	NA	2	86	0.2(0.4)	0.3(0.7)	2.6	0.23	18.3
5				0.5(0.8)	0.4(0.6)		36.85	

^a Number in this column indicates zone of inhibition, in millimeters, measured from the edge of a circular filter disk impregnated with test substance; sample concentration 0.02 mg/disk. ^b Tyrosine kinase p56^{lck} percent inhibition using 200 μg/mL of sample. ^c Rat skeletal muscle myoblast cells (L6). ^d NA, indicates a substance was inactive in the applied test. ^e On the basis of the counter ion being Cl.

in the case of the quinolines, is heme.²² Compounds **3** and **4** also have antitrypanosomal effects and are weakly cytotoxic.

Conclusions

Four (**1–4**) unusual decahydroquinolines were isolated from a new species of *Didemnum*. The most pronounced biological activities were seen for compounds **3** and **4** in antiplasmodial and antitrypanosomal tests. This biological activity seems to be dependent on the configuration at C-2 and the nature of the functionality at C-3 in the decahydroquinoline. Clearly, the low cytotoxicity of the molecules isolated and tested in this study makes them suitable as models for the development of potential therapeutic agents.

Experimental Section

General Experimental Procedures. These were performed as previously reported.²³

Animal Material. Animal material was obtained in May 1983, from Stanley Reef, the Great Barrier Reef, Queensland, Australia. Animals growing at 4–7 m depth were collected, deep frozen, and on return to the laboratory, freeze-dried. A voucher specimen is deposited at the Department for Pharmaceutical Biology, University of Bonn (voucher no. LL48).

Extraction and Isolation. Tunicate tissue (53.0 g) was exhaustively extracted with (CH₃)₂CO (2 L) followed by aqueous MeOH (80%) (2.5 L) to yield 1.76 g of a (CH₃)₂CO soluble material. Normal phase vacuum-liquid chromatography (NP VLC) of the (CH₃)₂CO solubles, using hexane with increasing proportions of EtOAc as eluent, followed by MeOH, afforded 14 fractions each of 110 mL. Thin-layer chromatography (TLC) and ¹H NMR examination of these fractions indicated VLC fraction 12 to be of further interest. VLC separation of this fraction employing normal phase silica and gradient elution from EtOAc to MeOH yielded eight fractions (12.1–12.8) each of 80 mL. VLC separation of fraction 12.2 employing normal phase silica and elution with a 9:1 mixture of CH₂Cl₂ and MeOH afforded a further six fractions (12.2.1–12.2.6). Normal phase column chromatography, employing EtOAc:MeOH:NH₃ 8:2:0.1 as eluent, of VLC fraction 12.2.3 yielded compound **2**. Normal phase column chromatography, employing EtOAc:MeOH:NH₃ 8:2:0.1 as eluent, of VLC fraction 12.2.4 yielded compound **1**. Normal phase column chromatography, employing EtOAc:(CH₃)₂CO:MeOH:H₂O:NH₃ 12:43:43:1:1 as eluent, of VLC fraction 12.3 gave a further 11 fractions (12.3.1–12.3.11). Normal phase high-performance liquid chromatography (HPLC) separation of fraction 12.3.3, using EtOAc:MeOH:NH₃ 9:1:0.1 as eluent, yielded compound **3**. Normal phase HPLC separation of fraction 12.3.6, using EtOAc:MeOH:NH₃ 9:1:0.1 as eluent, yielded compound **4**.

(2R*,3S*,4aR*,5R*,8aS*)-Decahydro-3-hydroxy-5-(5'-hydroxyoctyl)-2-methylquinoline (Lepadin D, 1). A clear oil (4 mg, 0.008%); [α]_D²² +3.0° (c 0.2, MeOH). IR ν_{max} (film): 3290, 3250, 2915, 1460 cm⁻¹. ¹H and ¹³C NMR data, see Tables 1 and 2. EIMS: m/z [M + H]⁺ 298 (2), [M]⁺ 297 (6), 282 (4), 264 (10), 254 (54), 236 (28), 168 (100), 119 (36). HREIMS: m/z 297.266 (calcd for C₁₈H₃₅NO₂ [M]⁺, 297.2668).

(2R*,3S*,4aR*,5R*,8aS*)-Decahydro-3-hydroxy-5-(5'-hydroxyoctyl)-2-methylquinolinium (2). A clear oil (9 mg, 0.017%); [α]_D²² -14.4° (c 0.27, MeOH). IR ν_{max} (film): 3340, 2930, 1455 cm⁻¹. ¹H and ¹³C NMR data, see Tables 1 and 2. EIMS: m/z [M]⁺ 298 (2), 297 (8), 282 (4), 264 (10), 254 (54), 236 (28), 168 (100). HREIMS: m/z 297.266 (calcd for C₁₈H₃₅NO₂ [M - H]⁺, 297.2668).

(2R*,2'E,3S*,4aR*,5R*,8aS*)-Decahydro-3-hydroxy-5-(5'-hydroxyoctyl)-2-methyl-3-quinolinyl Ester 2''-Octenoic Acid (Lepadin E, 3). A clear oil (15 mg, 0.028%); [α]_D²² -2.0° (c 0.1, MeOH). IR ν_{max} (film): 3360, 2925, 1715 cm⁻¹. UV λ_{max} (MeOH): 263 nm (ε 2110). ¹H and ¹³C NMR data, see Tables 1 and 2. EIMS: m/z [M]⁺ 421 (<1), 420 (<1), 406 (1), 378 (2), 292 (5), 279 (100), 236 (60), 206 (20), 178 (54), 150 (24). HREIMS: m/z 421.356 (calcd for C₂₆H₄₇NO₃, 421.3556).

(2S*,2'E,3S*,4aR*,5R*,8aS*)-Decahydro-3-hydroxy-5-(5'-hydroxyoctyl)-2-methyl-3-quinolinyl Ester 2''-Octenoic Acid (Lepadin F, 4). A clear oil (10 mg, 0.019%); [α]_D²² -1.5° (c 0.1, CHCl₃). IR ν_{max} (film): 3360, 2925, 1715 cm⁻¹. UV λ_{max} (MeOH): 264 nm (ε 2730). ¹H and ¹³C NMR data, see Tables 1 and 2. EIMS: m/z [M]⁺ 421 (<1), 420 (<1), 406 (1), 378 (2), 292 (5), 279 (100), 236 (60), 206 (20), 178 (54), 150 (24). HREIMS: m/z 421.356 (calcd for C₂₆H₄₇NO₃, 421.3556).

To a stirred CH₂Cl₂ solution of **3** (6 mg in 3 mL), 3 mol equiv of para-bromobenzoyl chloride and 2 mg of dimethyl aminopyridine were added. The resultant solution was stirred for 10 h at 24 °C, and the reaction was then quenched with H₂O (5 mL). The H₂O phase was then extracted with CH₂Cl₂ (3 × 5 mL). The combined CH₂Cl₂ solubles were purified by normal phase HPLC [(CH₃)₂CO:petroleum ether, 50:50 as eluent] to yield **5**.

Para-Bromobenzoyl Derivative of 3 (5). A clear oil (4 mg). ¹H and ¹³C NMR data, partially assigned data available from the authors by request. EIMS: m/z [M]⁺ 605, 603 (4, 4), 562, 560 (4, 5), 463, 461(100, 99). HREIMS: m/z 603.292 (calcd for C₃₃H₅₀⁷⁹BrNO₄, 603.2923).

Biological Testing. The antimicrobial,²⁴ TK inhibition,²⁵ reverse transcriptase (RT) inhibition,²⁶ antimalarial,²⁷ antitrypanosomal,^{28,29} and cytotoxicity³⁰ assays were carried out as previously described.

Agar Diffusion Test. Test organisms for the agar diffusion assays were the bacteria *Bacillus megaterium* de Bary (Gram positive) and *Escherichia coli* (Migula) Castellani and Chambers (Gram negative), the fungi *Microbotryum violaceum* (Pers.) Roussel, *Eurotium repens* Corda, and the green micro-alga *Chlorella fusca* Shih Krauss. Samples were prepared by taking 100 μL (200 μg) of the solution of test sample and

pipetting them onto a sterile antibiotic filter disk, which was then placed onto the appropriate growth medium and sprayed with a suspension of the corresponding test organism and subsequently incubated. Zones of inhibition were measured from the edge of filter disks. Samples displaying a growth inhibition zone ≥ 3 mm and/or a complete inhibition zone ≥ 1 mm were considered as positive.²⁴

Inhibition of TK. The TK inhibition assay was carried out using T cell tyrosine kinase p56^{lck} and a commercial test kit (ELISA). Sample concentrations were 200 $\mu\text{g}/\text{mL}$.²⁵

Inhibition of HIV-1 RT. Activity of recombinant HIV-1 RT was measured according to the ELISA protocol established by Eberle and Seibl.²⁶ Sample concentrations were 66 $\mu\text{g}/\text{mL}$. A sample was considered as active when the RT activity was reduced by 20% or more. All compounds tested in this assay (1–5) were found to be inactive.

Antimalarial Activity. Cultures of *P. falciparum* (chloroquine-sensitive strain NF54 and chloroquine-resistant strain K1) were maintained in human erythrocytes [(A+, Albumax, (Invitrogen)] in vitro. The antimalarial activity of extracts was assessed using an in vitro radio isotope incorporation method. IC₅₀ values of test extracts, reference, and positive control were determined from logarithmic dose–response curves and were expressed as a percentage of inhibited parasite specific ³H-hypoxanthine incorporation.²⁷ Samples with IC₅₀ < 10 $\mu\text{g}/\text{mL}$ were considered as active.

Activity Against Trypanosomes of African Origin. Activity of compounds against *T. brucei rhodesiense*, the causative agent of African sleeping sickness, was determined according to R az et al.²⁸ Briefly, parasites were propagated axenically in culture medium supplemented with horse serum. Cultures in logarithmic growth phase were propagated for 3 days in the presence of various drug concentrations. Viability of parasites was quantified using the dye Alamar Blue according to R az et al.²⁸ Drug sensitivity was regarded as positive when MIC < 100 $\mu\text{g}/\text{mL}$.

Activity of compounds against *T. cruzi* (Tulahuen strain), the causative agent of Chagas disease, was determined according to Buckner et al.²⁹ Briefly, a *T. cruzi* strain that expresses the *E. coli* β -galactosidase gene was propagated intracellularly in mammalian fibroblast cells in 96 well microtiter plates. Cultures were propagated for 5 days in the presence of drug, before viability of *T. cruzi*, amastigote forms, was determined. Transfected parasites catalyze a colorimetric reaction with chlorophenol red and β -D-galacto-pyranoside as substrate. The reaction was quantified employing an enzyme-linked immunosorbent assay. Drug sensitivity was regarded as positive when MIC < 100 $\mu\text{g}/\text{mL}$.

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