Chinikomycins A and B: Isolation, Structure Elucidation, and Biological Activity of Novel Antibiotics from a Marine *Streptomyces* sp. Isolate M045^{#,1}

Fuchao Li,^{†,‡,⊥} Rajendra P. Maskey,[†] Song Qin,[‡] Isabel Sattler,[†] Heinz H. Fiebig,[§] Armin Maier,[§] Axel Zeeck,[†] and Hartmut Laatsch^{*,†}

Department of Organic and Biomolecular Chemistry, University of Göttingen, Tammannstrasse 2, D-37077 Göttingen, Germany, Institute of Oceanology, Chinese Academy of Sciences, Nanhai Road 7, 266071 Qingdao, People's Republic of China, Graduate School of the Chinese Academy of Sciences, Yuquan Road 19A, 100039 Beijing, People's Republic of China, and Oncotest GmbH, Am Flughafen 12-14, D-79108 Freiburg, Germany

Received November 26, 2003

In our screening of marine Streptomycetes for bioactive principles, two novel antitumor antibiotics designated as chinikomycins A (2a) and B (2b) were isolated together with manumycin A (1), and their structures were elucidated by a detailed interpretation of their spectra. Chinikomycins A (2a) and B (2b) are chlorine-containing aromatized manumycin derivatives of the type 64-pABA-2 with an unusual *para* orientation of the side chains. They exhibited antitumor activity against different human cancer cell lines, but were inactive in antiviral, antimicrobial, and phytotoxicity tests.

The manumycins constitute a class of antibiotics with antibiotic, cytotoxic, and other biological activities; manumycin A (1) especially inhibits the Ras farnesyl transferase.² Structurally they contain a central six-membered ring $(m-C_7N \text{ unit})$ of biosynthetic origin from 3-amino-5hydroxybenzoic acid, and two 1,3-arranged polyketide chains. A methyl-branched unsaturated fatty acid is linked to the central ring via an amide bond. The second side chain extends from the mC₇N unit as a structurally conserved all-trans triene (except in colabomycins A and D and U-62,162) and terminates with a carboxamide group linked to a 2-amino-3-hydroxycyclopent-2-enone moiety $(C_5N \text{ unit})$.³ The first member of the manumycin family, manumycin A (1), was isolated in 1963 by Zähner and coworkers from Streptomyces parvulus Tü 64.4 Its constitution was determined in 1973,⁵ and the stereochemistry was reported in 1987³ and revised in 1998.⁶



In our screening program for bioactive principles from marine streptomycetes, two novel antibiotics belonging to the manumycin group were isolated, which we named chinikomycins A (**2a**) and B (**2b**) (niko in Nepali = to be cured). In addition, the strain also produced manumycin A (**1**). In this paper we report the isolation and the structure elucidation of **2a** and **2b** as well as the biological activity of these compounds.

Results and Discussion

The ethyl acetate extract of *Streptomyces* sp. isolate M045 drew our attention due to its biological activity against various microorganisms and microalgae. In the chemical screening of the extract, in addition to fats, fatty acids, and prodigiosins, a striking middle polar yellow band was observed, which was found to be responsible for the antibacterial activity.

On variation of the growth medium, temperature, and starting pH, the production of the bioactive component was increased to about 15-25 mg/L in fish flour medium (FF medium) with a pH starting at 6.5 and a cultivation temperature of 35 °C. The crude extract, obtained after usual workup7 of 60 1 L Erlenmeyer flasks each containing 250 mL of FF medium, was subjected to flash chromatography on silica gel with a MeOH/CH₂Cl₂ gradient to yield three fractions. Repeated separation of fraction II delivered manumycin A (1) and chinikomycin A (2a) as yellow solids and the red chinikomycin B (2b). The latter was produced by oxidation of 2a during its isolation and is therefore not a primary natural product. Treatment of fraction III with silver oxide delivered a further crop of chinikomycin B (2b). Manumycin A (1) was easily identified by a substructure search in AntiBase⁸ and by comparison of the NMR data with the literature values.²

The molecular weights of 2a and 2b were determined by ESIMS to be 568 and 566, respectively. Isotopic peaks of the molecular ions indicated both to contain one chlorine atom each. ESI HR mass spectrometry of the $[M + Na]^+$ quasi-molecular signals led to the molecular formulas $C_{31}H_{37}ClN_2O_6$ and $C_{31}H_{35}ClN_2O_6$ for **2a** and **2b**, and thus the same number of carbon atoms is found as for manumaycin A (1). The proton NMR spectrum of compounds 2a and **2b** both showed nine signals for protons at sp² carbon atoms, four methyl signals, and two further multiplets of five and six protons at δ 2.6–2.4 and 1.4–1.2, respectively. In addition, the spectrum of 2a contained five signals of acidic protons, whereas that of 2b showed only three. This and the conversion of **2a** into **2b** with silver oxide and the reverse reaction with aqueous dithionite solution indicated the red **2b** to be the quinone of the hydroguinone **2a**. The ¹³C NMR and APT spectra of **2a** showed only 27 instead of

10.1021/np030518r CCC: \$30.25 © 2005 American Chemical Society and American Society of Pharmacognosy Published on Web 02/26/2005

 [#] Dedicated to Prof. Dr. H.-G. Floss on the occasion of his 70th birthday.
 * To whom correspondence should be addressed. Tel: +49-551-393211.
 Fax: +49-551-399660. E-mail: hlaatsc@gwdg.de.

[†] University of Göttingen.

^{*} Institute of Oceanology, Chinese Academy of Sciences.

 $^{^{\}perp}$ Graduate School of the Chinese Academy of Sciences.

[§] Oncotest.

Table 1. ¹³C NMR (125.7 MHz, δ Values) Data of 1, 2a, and 2b

C no.	$2\mathbf{a}^a$	$\mathbf{2b}^{b}$	1^{b}	C no.	$2\mathbf{a}^a$	$\mathbf{2b}^{b}$	1^{b}	C no.	$2\mathbf{a}^a$	$2\mathbf{b}^b$	1^b
1	136.8	176.3	189.0	12	121.0	123.7	121.7	5''	141.6	144.2	142.7
2	128.6	137.7	128.0	13	166.1	165.2	165.6	6''	32.1	32.9	32.9
3	107.6	114.7	126.6	1′	n.v.	197.5	197.6	$7^{\prime\prime}$	36.4	37.0	37.1
4	150.3	185.2	71.2	2'	114.9	115.1	115.5	8″	29.2	29.7	29.8
5	117.0	138.4	57.4	3′	n.v.	174.6	174.5	9″	22.2	22.8	22.8
6	122.2	135.0	52.8	4'	n.v.	32.1	32.2	10''	13.9	14.0	14.1
7	130.4	127.3	136.6	5'	n.v.	25.6	25.7	2"-Me	14.1	14.1	14.0
8	132.8	144.3	131.4	1″	168.3	168.3	168.8	5"-Me	16.3	16.4	16.5
9	143.0	141.5	139.6	$2^{\prime\prime}$	128.5	127.6	128.4	6''-Me	20.6	20.6	20.7
10	129.8	136.3	131.7	3″	138.7	141.8	140.2				
11	142.6	142.8	143.4	4‴	130.0	129.9	129.9				

^{*a*} DMSO- d_6 . ^{*b*} CDCl₃, n.v. = not visible

31 signals. The spectra of compound **2b**, however, exhibited two further carbonyl signals and two additional methylene carbon signals as demanded by the molecular formula.

Interpretation of the 2D NMR spectra (H,H COSY, HMQC, HMBC, and NOESY) led to 2,4,6-trimethyl-2E,4E-deca-2,4-dienoic acid amide as substructure **I** of both compounds **2a** and **2b** (Figure 1). This fragment is known



Figure 1. (2*E*,4*E*)-2,4,6-Trimethyldeca-2,4-dienoic acid amide fragment (**I**) of chinikomycins A (**2a**) and B (**2b**) and of manumycin A (**1**) derived from H,H COSY (not shown), HMQC (not shown), HMBC (\rightarrow), and NOESY experiments (\leftrightarrow).

as part of manumycin $(1)^2$ and was finally confirmed by comparison of the ¹³C NMR data with the literature values (Table 1) and by the presence of 1 in the extracts.

With the aid of the coupling constants of the olefinic protons and their H,H COSY correlations, a system of three conjugated *trans* double bonds was constructed. By careful interpretation of the H,H COSY, HMQC, and HMBC couplings, substructures **II** and **III** were developed for compounds **2a** and **2b**, respectively (Figure 2). The carbon



Figure 2. Correlation of the HMBC (\rightarrow , only selected) and H,H COSY (\leftrightarrow) experiments in fragments II and III of chinikomycins A (**2a**) and B (**2b**), respectively.

skeleton of substructure **II** is part of 64-p-ABA-2 (2c),⁹ which was obtained by feeding of *Streptomyces parvulus* with 4-aminobenzoic acid; however, in 2a two additional hydroxyl groups and the chlorine atom are present.

In the substructures I and II of 2a and I and III of 2b, all expected 26 carbon atoms can be assigned easily. Since in 2a not all carbon signals were visible, further interpre-

tation was done on **2b**. The resulting substructure IV (Figure 3) was confirmed by comparison of the ¹³C NMR



Figure 3. Cyclopentan-1,3-dione fragment (**IV**) of the chinikomycins derived from H,H COSY (\leftrightarrow), HMQC (not shown), and HMBC (\rightarrow) spectra of **2a/2b**.

data with the respective part of manumycin A (1). Also in the latter, the carbon signals of the cyclopentan-1,3-dione moiety are broad,¹⁰ which explains the missing signals in **2a**. The substructure **IV** is additionally proven by the similar fragmentation pattern on ESI MS-MS of manumycin A (1) and chinikomycin A (**2a**): The quasi-molecular signals at m/z 573 ([M + Na]⁺) and 591 ([M + Na]⁺) of **1** and **2a** delivered ions at m/z 460 and 478, respectively, by loss of the 2-aminocyclopentan-1,3-dione fragment.

As the amide carbonyl group C-1" in I is overlapping with the anilide structure in II and III, and as IV must be connected with the triene chain of II/III as in 1 (see Tab. 1), the chlorine atom must be attached to the free valence at C-5 of the fragments II/III to complete the structures 2a and 2b of chinikomycins A and B.



Whereas 64-pABA-2 (**2c**) and a few related compounds have been obtained in a precursor-directed biosynthesis by feeding of *p*-aminobenzoic acid,⁹ the chinikomycins are the first truly natural manumycin antibiotics with a *para* orientation of the side chains. We realized, however, that a compound with shift values identical to those of **2a** had been isolated by one of the authors previously from *Streptomyces parvulus* Tü 64.¹¹ The isolated amount had been insufficient to solve the structure with the technical capabilities at that time.

The chlorohydrin 3 is not an intermediate in the biosynthesis of 1;² however, it may be formed from the latter by ring opening with chloride and could easily lose water at C-5. The carbocation thus formed could, after protonation, undergo a sextet rearrangement to deliver 4, and depro-

Table 2. Antitumor Activity of **2a** and **2b** in Human TumorCell Lines

	IC	50^a
tumor cell line	2a	2b
LXFA 629L	>6.0	>6.0
LXFL 529L	>6.0	>6.0
MAXF 401NL	2.41	3.04
MEXF 462NL	4.15	>6.0
RXF 944L	4.02	>6.0
UXF 1138L	>6.0	>6.0

^{*a*} IC₅₀ values in μ g/mL.

tonation at OH-4 yielding a carbonyl group and enolization at C-1/C-4 would result in 2a. It is obvious that the assumption of the alternative chlorohydrin 5 would finally deliver the isomer 6 even without a rearrangement of the carbon skeleton. There is, however, little doubt that 2a is the preferable structure: According to the HMBC data (Figure 2), the OH group at δ 10.04 must be connected with the carbon atom at δ 150.3 and be placed between the aromatic CH (C-3) and the ring atom bearing the conjugated chain (C-5). In 6, these correlations are substituted by less favorable 4- and 5-bond couplings of OH-6. Furthermore, H-7 shows a cross signal with the signal at δ 150.3 (C-4) which is due to a ${}^{3}J$ coupling in **2a**, but would require a ${}^{4}J$ coupling in **6**. Similar correlations supporting structure 2b are found in the HMBC spectrum of the corresponding quinone. Diagnostic is the strong cross signal between H-7 (δ 6.83) and δ 185.2 as expected for CO-4 in 2b; in 6, the corresponding coupling should be much weaker $({}^{4}J)$. The cross signal between H-3 and C-7 is very weak, as expected for 2b, but should be strong in 6. An NOE signal between H-3 and H-7/8 is not visible (but would be expected for 6), and a strong interaction between the OH-4 group (signal at δ 10.04) and H-3 is a final proof for their ortho orientation as in 2a. Also the experimental values agree better with the calculated ¹³C shifts¹² on the basis of **2a** than for **6**. It should be mentioned additionally that the formation of chinikomycin B by air oxidation is explained better by 2b than by an ortho-quinone derived from 6 with its higher oxidation potential.



Biological Activity. Chinikomycins A (**2a**) and B (**2b**) both exhibited moderate antitumor activity, **2a** being the more potent one. The hydroquinone **2a** selectively inhibited proliferation in cell lines of mammary cancer (MAXF 401NL, $IC_{50} = 2.41 \, \mu g/mL$), melanoma (MEXF 462NL, $IC_{50} = 4.15 \, \mu g/mL$), and renal cancer (RXF 944L, $IC_{50} = 4.02 \, \mu g/mL$). The quinone **2b** showed selective antitumor activity against the mammary cancer cell line MAXF 401NL ($IC_{50} = 3.04 \, \mu g/mL$) (see Table 2). Cell lines of non small cell lung cancer (LXFA 629L, LXFL 529L) and uterus

cancer (UXF 1138L) were not sensitive to 2a and 2b. The chinikomycins exhibited no antimicrobial or antiviral activities.

Experimental Section

General Experimental Procedures. Material and methods and antimicrobial tests were used as described earlier.⁷ ESIHRMS was measured on a Micromass LCT mass spectrometer coupled with a HP 1100 HPLC with a diode array detector. Reserpin (MW = 608) and Leucin-Enkephalin (MW = 555) were used as standards in positive and negative mode. R_f -values were measured on Polygram SIL G/UV₂₅₄ (Macherey-Nagel & Co., Düren, Germany).

FF Medium. Glucose (21 g), fish powder (5 g), wheat flour (10 g), yeast extract (1 g), magnesium sulfate (0.5 g), sodium chloride (1 g), calcium chloride (0.5 g), and trace element solution (10 mL) were dissolved in artificial seawater (0.5 L) and tap water (0.5 L). Trace element solution: ferrous sulfate (10.2 g), cobalt chloride (0.04 g), calcium chloride (0.04 g), manganese chloride (0.04 g), zinc sulfate (0.08 g), sodium borate (0.08 g), and sodium molybdate (0.74 g) were dissolved in 500 mL of water.

 M_2^+ Medium. Malt extract (10 g), yeast extract (4 g), and glucose (4 g) were dissolved in artificial seawater (0.5 L) and tap water (0.5 L).

LB Medium. Tryptone (10 g), yeast extract (5 g), sodium chloride (10 g), and glucose (5 g) were dissolved in artificial seawater (0.5 L) and tap water (0.5 L).

SM Medium. Defatted soybean flour (20 g) and mannitol (20 g) were dissolved in artificial seawater (0.5 L) and tap water (0.5 L).

ME Medium. Meat extract (1 g), glucose (10 g), peptone (2 g), and yeast extract (1 g) were dissolved in artificial seawater (0.5 L) and tap water (0.5 L).

CL Medium. Calcium chloride (45 g), glucose (5 g), and yeast extract (40 g) were dissolved in artificial seawater (0.5 L) and tap water (0.5 L).

Gause's Starch Medium. Soluble starch (20 g), potassium nitrate (1 g), potassium hydrogen phosphate (0.5 g), magnesium sulfate (0.5 g), sodium chloride (0.5 g), ferrous sulfate (0.01 g), and agar (18 g) were dissolved in 1 L of water containing 50% natural seawater and 0.1 mg/mL K₂Cr₂O₇. The pH of the solution was set to 7.2 before sterilization.

Taxonomy of the Producing Strain. The strain M045 has been derived from sediment of Jiaozhou Bay in China. It was isolated on Gause's starch medium with incubation at 28 °C. The pure culture was maintained on Gause's starch agar medium with K₂Cr₂O₇ at 4 °C. The strain forms a red vegetative mycelium and a white aerial mycelium. The substrate mycelium does not have transverse septa and no fragments. The aerial mycelium has many branches. The strain forms spiral sporophores (Spirales). The spores are oval with smooth surface. Melanin pigment is not produced on tyrosine agar,¹⁴ and water-soluble pigment is not produced on other media. The temperature optimum is at ~ 28 °C. The strain does not grow at 45 °C or at 10 °C. Gelatin and starch are degraded. Hydrogen sulfide is not produced. The strain is catalase positive, and nitrate reductase is not formed. Cell walls were purified and amino acids analyzed using thin-layer chromatography as described by Lechevalier and Lechevalier.¹⁵ The peptidoglycan cell wall of the strain contains major amounts of L-diaminopimelic acid (L-DAP) and glycine but no diagnostic sugars in whole cell hydrolysates of the strain (cell wall chemotype I).¹⁶ Due to its chemical and morphological features as well as the 16S rRNA (GenBank accession number AY644669), the strain can be assigned to the genus Streptomyces. The strain is deposited in the culture collection of marine actinomycetes at the Institute of Oceanology, Chinese Academy of Sciences, Nanhai Road 7, 266071 Qingdao, China.

Optimization of Chinikomycin A (2a) Production by Streptomyces sp. Isolate M045. In M_2^+ medium using our standard conditions for the screening of streptomycetes, the strain Streptomyces sp. isolate M045 produced chinikomycin A (2a) only in a very small amount ($\ll 1$ mg/L). To increase

Table 3. Results of the Optimization of Chinikomycin A (**2a**) Production (mg/L) by *Streptomyces* sp. Isolate M045 (pH values were adjusted by adding dilute NaOH prior to sterilization)

	pН	[6.5	pH 8.0		
medium	28 °C	35 °C	28 °C	35 °C	
M_2^+ medium	0	0	≪1	0	
SM medium	0	0	0	0	
ME medium	0	0	≪1	0	
CL medium	0	0	0	0	
LB medium	0	0	0	0	
FF medium	0	15 - 25	0	≪1	

the production of 2a, the strain was grown in different media under four different conditions. The 2a concentration was determined by integration of the HPLC signal and by comparison with a pure sample. The results of the optimization are listed in Table 3.

Fermentation of Streptomyces sp. Isolate M045 and Workup. The marine strain Streptomyces sp. isolate M045 was inoculated from its slant agar culture on M_2^+ agar plates with 50% seawater. It formed red colonies with a white aerial mycelium and a red coloration of the agar on incubation at 28 °C while shaking for 72 h. With fresh agar cultures of the strain, 60 1 L Erlenmeyer flasks each containing 250 mL of FF medium (set to the pH 6.5 before sterilization) were inoculated and grown for 4 days at 35 °C while shaking at 120 rpm. A dark brown culture broth with a yellowish gray mycelium was obtained.

The entire culture broth was mixed with ca. 1 kg of diatomaceous earth and pressed through a pressure filter, and both filtrate and residue were extracted with ethyl acetate. The combined extracts were evaporated in vacuo at 30 °C to dryness to yield 14 g of the crude extract, which contained yellow 2a, prodigiosins, and a large amount of fat and fatty acids. The extract was triturated with cyclohexane (250 mL), and the insoluble solid was separated by filtration through a sintered glass funnel to get extract A (cyclohexane soluble part) and the precipitate B (insoluble material). The solid B was subjected to silica gel column chromatography $(3 \times 60 \text{ cm}, 270 \text{ cm})$ g of silica gel, dichloromethane/methanol gradient, 1.0 L of CH2Cl2, 2.0 L of CH2Cl2/0.5% MeOH, 1.0 L of CH2Cl2/2% MeOH) and separated under TLC control into fractions I (80 mg), II (325 mg), and III (414 mg). Fraction I contained only fats and prodigiosins and was not further analyzed.

Fraction II was washed with cyclohexane/25% dichloromethane $(2 \times 25 \text{ mL})$ to get a yellow solid (fraction IIa) containing mainly 2a. On TLC, the yellow zone of 2a ($R_f =$ 0.50) turned red with sodium hydroxide. On silica gel and Sephadex LH 20, 2a is partially oxidized, forming red 2b. The crude material was separated twice by PTLC (20×20 cm, CH₂Cl₂/5% MeOH) and quickly recovered while the plates were still wet to deliver 50 mg of chinikomycin A (2a). The red zone that appeared during the separation yielded 3.5 mg of red chinikomycin B (2b). The cyclohexane/25% dichloromethanesoluble part (fraction IIb) showed strong antibacterial activity and contained a light yellow band on TLC, which was more stable than 2a and showed no color reaction with dilute sodium hydroxide but turned blue-violet on spraying with anisaldehyde/sulfuric acid. Purification of fraction IIb by PTLC (20 imes20 cm, CH₂Cl₂/7% MeOH) followed by Sephadex LH-20 yielded 90 mg of manumycin A (1, $R_f = 0.50$).

Extract A was similarly purified on silica gel columns first with a dichloromethane/methanol gradient and then with cyclohexane/70% ethyl acetate. The product was finally washed twice with cyclohexane (2×50 mL) to yield another 100 mg of **2a** and 30 mg of **1**.

Chinikomycin A (2a): yellowish brown solid; $R_f = 0.34$ (CH₂Cl₂/5% MeOH); UV/vis (MeOH) λ_{max} (log ϵ) 260 (4.46), 335 (s, 4.46), 395 (4.62) nm; OR $[\alpha]_D^{20}$ -86° (c 0.02, CHCl3); no CD effect visible; IR (KBr) ν_{max} 3425, 2965, 2924, 2851, 1657, 1620, 1538, 1501, 1383, 1319, 1228, 1173, 1084, 1008, 873, 662, 546 cm⁻¹; ¹H NMR (DMSO- d_6 , 500 MHz) δ 13.90 (1 H, s br, 3'-OH), 10.04 (1 H, s, 4-OH), 9.86 (1 H, s, 13-NH), 9.12 (1 H,

s, 2-NH), 9.06 (1 H, s, 1-OH), 7.47 (1 H, s, H-3), 7.45 (1 H, dd, ${}^{3}J = 15.5$, 11.1 Hz, H-8), 7.26 (1 H, dd, ${}^{3}J = 15.1$, 11.2 Hz, H-11), 6.97 (1 H, d, ${}^{3}J = 15.5$ Hz, H-7), 6.86 (1 H, dd, ${}^{3}J = 15.1$, 11.1 Hz, H-9), 6.82 (1 H, s, H-3"), 6.51 (1 H, d, ${}^{3}J = 14.6$ Hz, H-12), 6.49 (1 H, dd, ${}^{3}J = 14.6$, 11.2 Hz, H-10), 5.36 (1 H, d, ${}^{3}J = 9.5$ Hz, H-5"), 2.53 (4 H, m, H₂-4",5"), 2.51 (1 H, m, H-6"), 2.03 (3 H, s, 2"-Me), 1.82 (3 H, s, 4"-Me), 1.4–1.2 (6 H, m, H₂-7",8",9"), 0.96 (3 H, d, ${}^{3}J = 6.8$ Hz, 6"-Me), 0.88 (3 H, t, ${}^{3}J = 6.8$ Hz, H₃-10"); ${}^{13}C$ NMR (DMSO-d₆, 500 MHz), see Table 1; (+)-ESI-MS m/z (%) 615 ([M¹ + 2Na - H]⁺, 5), 613 ([M² + 2Na - H]⁺, 25), 593 ([M¹ + Na]⁺, 32), 591 ([M² + Na]⁺, 100); (-)-ESI-MS m/z (%) 569 ([M¹ - H]⁻, 52), 567 ([M² - H]⁻, 100); (+)-ESI HRMS m/z 591.2237 (calcd for [M² + Na]⁺, C₃₁H₃₇ClN₂O₆Na, 591.2237712).

Chinikomycin B (2b): red solid; $R_f = 0.58$ (CH₂Cl₂/5%) MeOH); UV/vis (MeOH) $\lambda_{\rm max}\,(\log\epsilon)$ 261 (4.68), 344 (4.72), 500 (3.98) nm; IR (KBr) ν_{max} 3425, 2963, 2923, 2857, 1655, 1620, 1502, 1457, 1382, 1318, 1297, 1233, 1173, 1121, 1083, 1035, 1008, 874, 851, 653, 546 cm⁻¹; ¹H NMR (CDCl₃, 500 MHz) δ 13.64 (1H, s br, 3'-OH), 8.60 (1H, s, 2-NH), 8.13 (1 H, s, 13-NH), 7.80 (1 H, dd, ${}^{3}J = 15.6$, 10.0 Hz, H-8), 7.60 (1 H, s, H-3), 7.37 (1 H, dd, ${}^{3}J = 14.8$, 10.3 Hz, H-11), 6.86 (1 H, s, H-3"), $6.83 (1 \text{ H}, \text{d}, {}^{3}J = 15.6 \text{ Hz}, \text{H-7}), 6.73 (1 \text{ H}, \text{dd}, {}^{3}J = 14.8, 10.1$ Hz, H-9), 6.68 (1 H, dd, ${}^{3}J = 14.8$, 10.3 Hz, H-10), 6.28 (1 H, d, ${}^{3}J = 14.8$ Hz, H-12), 5.40 (1 H, d, ${}^{3}J = 9.7$ Hz, H-5"), 2.62 $(2 \text{ H}, \text{ s br}, \text{H}_2\text{-}5'), 2.56 (2 \text{ H}, \text{ s br}, \text{H}_2\text{-}4'), 2.45 (1 \text{ H}, \text{ m}, \text{H}\text{-}6''),$ 2.08 (3 H, s 2"-Me), 1.83 (3 H, s, 4"-Me), 1.4-1.2 (6 H, m, H₂-7'', 8'', 9''), 0.97 (3 H, d, ${}^{3}J = 6.5$ Hz, 6''-Me), 0.86 (3 H, t, 7.0 Hz, H₃-10"); ¹³C NMR (CDCl₃, 500 MHz), see Table 1; (+)-ESI-MS m/z (%) 611 ([M² + 2Na - H]⁺); (-)-ESI-MS m/z (%) 567 ([M¹ – H]⁻, 29), 565 ([M² – H]⁻, 100); (–)-ESI HRMS m/z565.2104 (calcd for $[M^2 - H]^-$, $C_{31}H_{34}ClN_2O_6$, 565.2105274).

Oxidation of Chinikomycin A (2a). Fraction III was dissolved in 50 mL of dichloromethane and stirred under TLC control for 15 min with \sim 1 g of silver oxide until all yellow **2a** was transformed into red **2b**. Purification by PTLC (20 × 20 cm, CH₂Cl₂/5% MeOH) afforded 7 mg of chinikomycin B (**2b**).

Reduction of Chinikomycin B (2b). A red solution of chinikomycin B (**2b**, <1 mg) in dichloromethane (2 mL) and 0.1 N aqueous sodium dithionite solution (1 mL) was shaken in a test tube until the color of the organic phase had changed from red to yellow. The organic phase was separated and evaporated, and the residue was identified by TLC as chinikomycin A (**2a**).

Biological Activities. Antitumor tests were performed as described in ref 13. The chinikomycins (**2a/2b**) were inactive against *Bacillus subtilis*, *Streptomyces viridochromogenes* (Tü 57), *Staphylococcus aureus*, and *Escherichia coli*, the microal-gae *Chlorella vulgaris*, *Chlorella sorokiniana*, and *Scenedesmus subspicatus*, and the fungi *Mucor miehei* and *Candida albicans* at a concentration of 40 μ g/test plate, while manumycin (1) gave inhibition zones of 16–28 mm at 10 μ g/test plate. The respective activities of the crude extract were due to the manumycin A (1) content. In an antiviral assay with the EMC virus, the chinicomycins had at 100 μ g/mL no protective activity on A549 cells against the cytopathic effects of a viral infection.¹⁷

Acknowledgment. This work was supported by a grant from the Bundesministerium für Bildung and Forschung (BMBF, grant 03F0346A) and by the National High Technology Research and Development Program of China (863 Program, 2001AA624020) and the Key Innovative Project of CAS (KZCX3-SW-215). F.L. thanks the BMBF and the International Bureau at DLR for a scholarship (CHN 01/326). We are grateful to H. Zhumei for providing the strain and to Y. Takahashi of The Microbial Chemistry Research Center in Tokyo for very valuable hints during the structure elucidation.

References and Notes

 Article No. XXIV on Marine Bacteria. For Part XXIII, see: Veluri, R.; Oka, I.; Wagner-Döbler, I.; Laatsch, H. J. Nat. Prod. 2003, 66, 1520-1523.

- (2) Sattler, I.; Thiericke, R.; Zeeck, A. Nat. Prod. Rep. **1998**, 15, 221–240.
- (3) Zeeck, A.; Schroeder, K.; Frobel, K.; Grote, R.; Thiericke, R. J. Antibiot. 1987, 40, 1530-1540.
- (4) Buzzetti, F.; Gäumann, E.; Hütter, R.; Keller-Schierlein, W.; Neipp, L.; Prelog, V.; Zähner, H. Pharm. Acta Helv. 1963, 38, 871–874.
- (5) Schroeder, K.; Zeeck, A. Tetrahedron Lett. 1973, 50, 4995-4998.
- (6) Alcaraz, L.; Macdonald, G.; Ragot, J. P.; Lewis, N.; Taylor, R. J. K. J. Org. Chem. 1998, 63, 3526–3527.
- (7) Biabani, M. A. F.; Baake, M.; Lovisetto, B.; Laatsch, H.; Helmke, E.; Weyland, H. J. Antibiot. **1998**, 51, 333–340.
- (8) Laatsch, H. AntiBase 2002, A Natural Products Database for Rapid Structure Determination. Chemical Concepts, Weinheim 2002; see Internet http://www.gwdg.de/~ucoc/laatsch/.
- (9) Zeeck, A.; Sattler, I.; Boddien, C. DECHEMA Monogr. 1993, 129 (Wege zu neuen Produkten und Verfahren der Biotechnologie), 85– 95.
- (10) Shizuri, Y.; Ojika, M.; Yamada, K. *Tetrahedron Lett.* **1981**, *22*, 4291–4294.

- (11) Sattler, I. Vorläufer-dirigierte Biosynthese und Fermentation unter erhöhtem Sauerstoffpartialdruck-Isolierung und Strukturaufklärung neuer Verbindungen aus der Manumycingruppe. Ph.D. Thesis, University of Goettingen, Germany, 1992.
 (12) <u>ACD NMR simulation programs</u> Vers. 3.00; Advanced Chemistry
- ACD NMR simulation programs Vers. 3.00; Advanced Chemistry Development Inc.: Toronto, 1998.
 Dengler, W. A.; Schulte, J.; Berger, P. B.; Mertelsmann, R.; Fiebig,
- (13) Dengier, W. A.; Schulte, J.; Berger, P. B.; Mertelsmann, K.; Flebig, H. H. Anti-Cancer Drugs 1995, 6, 522–532.
- (14) Shirling, E. B.; Gottlieb, D. Int. J. Syst. Bacteriol. **1966**, *16*, 313–340.
- (15) Lechevalier, M. P.; Lechevalier, H. A. The Chemotaxonomy of actinomycetes. In Actinomycete Taxonomy, Special Publication 6; Society for Industrial Microbiology: Arlington, VA, 1980; pp 227– 291.
- (16) Lechevalier, M. P.; Lechevalier, H. A. Int. J. Syst. Bacteriol. 1970, 20, 435–443.
- (17) Busche, A. Identifizierung und Charakterisierung von Modulatoren der Interferon-y-Antwort. Ph.D. Thesis, University of Hanover, 2003. We thank A. Busche for performing the tests.

NP030518R