# Stereostructure of Amphotericin A 

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(Received for publication June 10, 1996)

The stereostructure of amphotericin A was established on the basis of NMR studies which contained DQF COSY, ROESY, ID TOCSY, HSQC and HMBC experiments.

Amphotericin A, a tetraene macrolide antibiotic, is a minor component ( $10 \%$ ) accompanying amphotericin B in the antibiotic complex produced by Streptomyces nodosus ${ }^{1,2)}$. Both components of the amphotericin complex, isolated in the pure crystalline forms, were reported to exhibit antifungal activity ${ }^{1,2)}$. The gross structure of amphotericin A, elucidated by MS and ${ }^{1} \mathrm{H}$ NMR, has been published ${ }^{3,4)}$. In this paper we report our NMR studies of amphotericin A which resulted in the assignment of its stereostructure.

## Results and Discussion

Amphotericin A(1), Fig. 1, was transformed into its $3^{\prime}-N$-acetyl methoxycarbonylmethylamide derivative (2), Fig. 1, which facilitated the purification process. The NMR studies of 2 , consisting of DQF-COSY, ROESY, 1D TOCSY, HSQC and HMBC experiments, resulted in full proton and carbon assignments, which are listed in Tables 1 and 2, respectively.

The vicinal coupling constants were measured from the $1 \mathrm{D}-{ }^{1} \mathrm{H}$ NMR spectrum, 1D-TOSCY experiments and, in a few cases, from the fine structure of the DQF-COSY cross-peaks.

The analysis of the DQF-COSY spectrum of $\mathbf{2}$ revealed all the connectivities within three structural blocks: $\mathrm{C}-2 \sim \mathrm{C}-12, \mathrm{C}-14 \sim \mathrm{C}-37$ and the sugar moiety (Fig. 1). These blocks were connected based upon the HMBC
experiment which displayed long-range heteronuclear connectivities. Thus, the blocks C-2 $\sim \mathrm{C}-12$ and C-14~C37 were connected via $\mathrm{C}-13$ due to couplings $\mathrm{H}-12 / \mathrm{C}-13$ and $\mathrm{H}-14 / \mathrm{C}-13$. The chemical shift of $\mathrm{H}-37$ ( 5.63 ppm ) associated with the $\mathrm{H}-2 / \mathrm{C}-1$ and $\mathrm{H}-37 / \mathrm{C}-1$ correlation revealed that the $\mathrm{C}-2 \sim \mathrm{C}-12$ fragment was connected via the lactone bond to $\mathrm{C}-37$. The location of the N acetylmycosaminyl substituent at $\mathrm{C}-19$ was pointed out by the $\mathrm{H}-1^{\prime} / \mathrm{C}-19$ correlation.

Thus, the gross structure of amphotericin A was confirmed as 1 which is shown in Fig. 1.

The geometries of the tetraene and diene chromophores, assigned as $20 \mathrm{E}, 22 \mathrm{E}, 24 \mathrm{E}, 26 \mathrm{E}$, and $30 \mathrm{E}, 32 \mathrm{E}$, respectively, were pointed out by the following set of vicinal coupling constants: $J_{20,21}=15.0 \mathrm{~Hz}, J_{22,23}=14.6$ $\mathrm{Hz}, J_{24,25}=15.0 \mathrm{~Hz}, J_{26,27}=15.0 \mathrm{~Hz}, J_{30,31}=15.0 \mathrm{~Hz}$, $J_{32,33}=15.0 \mathrm{~Hz}$.

The data required for the assignment of the geometry of the C-27~C-30 fragment could not be extracted from NMR spectra measured in the solvent system pyridine-$d_{5}$-methanol- $d_{4}$ 9:1 due to strong overlapping resonances of olephinic protons $\mathrm{H}-27$ and $\mathrm{H}-30$ as well as four protons of the C-28 and C-29 methylene groups.

Fortunately, the ${ }^{1} \mathrm{H}$ NMR spectrum of 2 measured in methanol- $d_{4}$ showed in the region of interest by the very well resolved proton resonances of $\mathrm{H}-27$ ( $\delta=5.72 \mathrm{ppm}$ ) and H-30 ( $\delta=5.58 \mathrm{ppm})$. This subsequently allowed via 1D-TOCSY experiments for the assignment of chemical

Fig. 1. The structure of amphotericin A (1) and its $3^{\prime}-N$-acetylamphotericin A methoxycarbonylmethylamide (2).

$1 \mathrm{R}^{1}=\mathrm{H}, \mathrm{R}^{2}=\mathrm{OH}$
$2 \mathrm{R}^{1}=\mathrm{CH}_{3} \mathrm{CO}, \mathrm{R}^{2}=\mathrm{NHCH}_{2} \mathrm{COOCH}_{3}$

Table 1. ${ }^{1} \mathrm{H}$ NMR data of $3^{\prime}-\mathrm{N}$-acetylamphotericin A methoxycarbonylmethylamide (2).

| No. | $\begin{gathered} \delta \\ (\mathrm{ppm}) \end{gathered}$ | Coupling partner ( $J, \mathrm{~Hz}$ ) | ROE to proton | No. | $\begin{gathered} \delta \\ (\mathrm{ppm}) \end{gathered}$ | Coupling partner ( $J, \mathrm{~Hz}$ ) | ROE to proton |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 2 a | 2.62 | 2b (15.0), 3 (4.5) | 2b, 3, 4a | 19 | 4.98 | 18a (3.8), 18b ( $\sim 0$, | $18 \mathrm{a}, \mathrm{b}, 20,21,1^{\prime}$ |
| 2 b | 2.72 | 2a (15.0), 3 (8.8) | 2a, 4b |  |  | 20 (7.7) |  |
| 3 | 4.67 | $\begin{gathered} 2 \mathrm{a}(4.5), 2 \mathrm{~b}(8.8) \\ 4 \mathrm{a}(4.4), 4 \mathrm{~b}(9.0) \end{gathered}$ | 2a, 4a 5 | 20 | 6.24 | 19 (7.7), 21 (15.0) | 11, 17, 19, 22 |
|  |  |  |  | 21 | 6.50 | 20 (15.0), 22 (10.6) | 19, 23 |
| 4 a | 1.82 | $\begin{aligned} & 3(4.4), 4 \mathrm{~b}(14.0), \\ & 5(4.0) \end{aligned}$ | 2a, 3, 4b, 5 | 22 | 6.38 | 21 (10.6), 23 (14.6) | 9, 11, 20, 24 |
|  |  |  |  | 23 | 6.21 | 22 (14.6), 24 (10.4) | 21, 25 |
| 4 b | 1.92 | $\begin{aligned} & 3(9.0), 4 \mathrm{a}(14.0) \\ & 5(8.4) \end{aligned}$ | $2 \mathrm{~b}, 4 \mathrm{a}$ | 24 | 6.31 | 23 (10.4), 25 (15.0) | 9, 22, 26 |
|  |  |  |  | 25 | 6.09 | 24 (15.0), 26 (10.3) | 23, 27 |
| 5 | 4.13 | $\begin{gathered} 4 \mathrm{a}(4.0), 4 \mathrm{~b}(8.4) \\ 6 \mathrm{a}(3.8), 6 \mathrm{~b}(8.1) \end{gathered}$ | 3, 7a, 28a, b | 26 | 6.14 | 25 (10.3), 27 (15.0) | 7b, 24, 28a, b |
|  |  |  |  | 27 | 5.65 | 26 (15.0) | 25 |
| 6 a | 1.86 | $\begin{gathered} 5(3.8), 6 \mathrm{~b}(12.0), \\ 7 \mathrm{~b}(\sim 9.0) \end{gathered}$ | $6 \mathrm{~b}, 7 \mathrm{a}, 8$ | 28a | 2.18 |  | 5,26 |
|  |  |  |  | 28b |  |  |  |
| 6 b | 2.03 | 5 (8.1), 6a (12.0) | 6a, 8 | 29a | 2.10 |  | 31 |
| 7 a | 1.98 | 7b (14.2), 8 (3.0) | 5, 6a, 7b, 8 | 29 b |  |  |  |
| 7 b | 2.14 | $\begin{aligned} & 6 \mathrm{a}(\sim 9.0), 7 \mathrm{a}(14.2), \\ & 8(9.4) \end{aligned}$ | 7a, 9, 26 | 30 |  | 5.64 | 31 (15.0) | 32$29, a, b, 33$ |
|  |  |  |  | 31 | 6.13 | 30 (15.0), 32 (10.2) |  |  |
| 8 | 3.82 | $\begin{aligned} & 7 \mathrm{a}(3.0), 7 \mathrm{~b}(9.4), \\ & 9(3.5) \end{aligned}$ | 6a, 6b, 7a, 10a | 32 | 6.29 | 31 (10.2), 33 (15.0) | 30, 34, 36, 37 |  |
|  |  |  |  | 33 | 5.74 | 32 (15.0), 34 (8.5) | 31, 35, 36, Me34 |  |
| 9 | 4.15 | $\begin{aligned} & 8(3.5), 10 \mathrm{a}(4.5), \\ & 10 \mathrm{~b}(9.8) \end{aligned}$ | 7a, 11, 22, 24 | 34 | 2.61 | $\begin{gathered} 33(8.5), 35(7.7) \\ \operatorname{Me} 34(6.8) \end{gathered}$ | 32, 36, 37, Me34 |  |
| 10a | 1.87 | $\begin{aligned} & 9(4.5), 10 \mathrm{~b}(14.0), \\ & 11(5.6) \end{aligned}$ | $8,10 \mathrm{~b}, 12 \mathrm{a}$ | Me34 | 1.23 | $34(6.8)$ | 33, 34, 35 |  |
|  |  |  |  | 35 | 3.49 | 34 (7.7), 36 (4.0) | $\begin{gathered} 33,36, \mathrm{Me} 34, \\ \mathrm{Me} 36 \end{gathered}$ |  |
| 10 b | 2.05 | $\begin{aligned} & 9(9.8), 10 \mathrm{a}(14.0), \\ & 11(9.6) \end{aligned}$ | 10a, 12b |  |  |  |  |  |
|  |  |  |  | 36 | 2.19 | $\begin{gathered} 35(4.0), 37(3.0) \\ \text { Me36 (6.9) } \end{gathered}$ | $32,33,34,35,37$ |  |
| 11 | 4.97 | $\begin{aligned} & 10 \mathrm{a}(5.6), 10 \mathrm{~b}(9.6) \\ & 12 \mathrm{a}(5.6), 12 \mathrm{~b}(9.9) \end{aligned}$ | 9, 20, 22 |  |  |  |  |  |
|  |  |  |  | Me36 | 1.09 | 36 (6.9) | 35, 36, Me37 |  |
| 12a | 1.86 | 11 (5.6), 12b (13.8) | 10a, 12b, 14a | 37 | 5.63 | 36 (3.0), Me37 (6.4) | 32, 34, 36, Me37 |  |
| 12b | 2.03 | 11 (9.9), 12a (13.8) | 10b, 12a, 14b | Me37 | 1.36 | 37 (6.4) | 37, Me36 |  |
| 14a | 1.73 | 14 b (13.6), 15 (11.5) | 12a, 14b, 16 | $1^{\prime}$ | 5.08 | $2^{\prime}(\sim 0)$ | $18 \mathrm{~b}, 19,3^{\prime}, 5^{\prime}$ |  |
| 14b | 2.51 | 14a (13.6), 15 (4.2) | 12b, 14a, 15 | 2 | 4.43 | $1^{\prime}(\sim 0), 3^{\prime}(3.3)$ | 17 |  |
| 15 | 5.09 | $\begin{aligned} & 14 \mathrm{a}(11.5), 14 \mathrm{~b}(4.2) \\ & 16(10.1) \end{aligned}$ | 14b, 17 | $3 '$ | 4.53 | $2^{\prime}(3.3), 4^{\prime}(9.5)$ | $1^{\prime}, 5^{\prime}$ |  |
|  |  |  |  | $4^{\prime}$ | 3.92 | $3^{\prime}(9.5), 5^{\prime}(9.3)$ | Me5 ${ }^{\prime}$ |  |
| 16 | 2.79 | 15 (10.1), 17 (10.1) | 14a, 18a | $5 '$ | 3.70 | $4^{\prime}(9.3), \mathrm{Me}^{\prime}(6.0)$ | $1^{\prime}, 3^{\prime}, \mathrm{Me} 5^{\prime}$ |  |
| 17 | 5.01 | $\begin{aligned} & 16(10.1), 18 \mathrm{a}(9.0) \\ & 18 \mathrm{~b}(7.0) \end{aligned}$ | 15, 18b, 20, $2^{\prime}$ | Me5' | 1.48 | $5^{\prime}(6.0)$ | $4^{\prime}, 5^{\prime}$ |  |
|  |  |  |  | Gly Ha | 3.94 | Gly Hb (18) | Gly Hb |  |
| 18a | 2.26 | $\begin{aligned} & 17(9.0), 18 b(13.5) \\ & 19(3.8) \end{aligned}$ | 16,18b, 19 | Gly Hb | 4.48 | Gly Ha (18) | Gly Ha |  |
|  |  |  |  | OMe | 3.65 |  |  |  |
| 18b | 2.70 | $\begin{aligned} & 17(7.0), 18 \mathrm{a}(13.5) \\ & 19(\sim 0) \end{aligned}$ | 17, 18a, 19, 1' | N -Ac | 2.01 |  |  |  |

Table 2. ${ }^{13} \mathrm{C}$ chemical shifts of $3^{\prime}-N$-acetylamphotericin A methoxycarbonylmethylamide (2).

| Carbon No. | $\begin{gathered} \delta \\ {[\mathrm{ppm}]} \end{gathered}$ | Carbon No. | $\begin{gathered} \delta \\ {[\mathrm{ppm}]} \end{gathered}$ | Carbon No. | $\begin{gathered} \delta \\ {[\mathrm{ppm}]} \end{gathered}$ | Carbon <br> No. | $\begin{gathered} \delta \\ {[\mathrm{ppm}]} \end{gathered}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 1 | 71.4 | 15 | 66.1 | 29 | 32.5 | $3^{\prime}$ | 56.2 |
| 2 | 43.7 | 16 | 59.2 | 30 | 131.7 | $4^{\prime}$ | 72.1 |
| 3 | 68.0 | 17 | 67.1 | 31 | 131.5 | $5^{\prime}$ | 74.3 |
| 4 | 44.4 | 18 | 38.1 | 32 | 130.5 | Me5' | 18.4 |
| 5 | 70.7 | 19 | 77.0 | 33 | 135.6 | $\mathrm{NH}-\mathrm{Ac}$ |  |
| 6 | 47.2 | 20 | 135.0 | 34 | 41.8 | CO | 171.2 |
| 7 | 30.0 | 21 | 131.0 | 35 | 77.6 | Me | 22.7 |
| 8 | 74.4 | 22 | 132.5 | 36 | 40.6 | GlyOMe |  |
| 9 | 74.0 | 23 | 132.6 | 37 | 71.5 | CH2 | 41.2 |
| 10 | 41.1 | 24 | 132.8 | Me34 | 17.3 | CO | 160.8 |
| 11 | 68.0 | 25 | 131.5 | Me36 | 12.5 | OMe | 52.0 |
| 12 | 35.8 | 26 | 133.0 | Me37 | 17.0 | COOR | 174.0 |
| 13 | 98.1 | 27 | 134.0 | $1{ }^{\prime}$ | 98.8 |  |  |
| 14 | 45.5 | 28 | 32.5 | $2^{\prime}$ | 70.9 |  |  |

shifts of H-26 ( $\delta=6.16 \mathrm{ppm}), \mathrm{H}-28 \mathrm{a}(\delta=2.20 \mathrm{ppm})$, H-28b ( $\delta=2.27 \mathrm{ppm}), ~ H-29 \mathrm{a} \quad(\delta=2.12 \mathrm{ppm}), ~ \mathrm{H}-29 \mathrm{~b}$ ( $\delta=2.23 \mathrm{ppm})$ and $\mathrm{H}-31(\delta=6.01 \mathrm{ppm})$.

The 2D-NOESY spectrum of 2 measured in methanol$d_{4}$ revealed that the $\mathrm{C}-28$ methylene protons exhibited NOE's to H-26, H-27 and H-30 while C-29 methylene protons showed NOE's to H-27, H-30 and H-31. This assignment associated with vicinal coupling constants $J_{27,28 \mathrm{a}}=6.8 \mathrm{~Hz}, J_{27,28 \mathrm{~b}}=6.8 \mathrm{~Hz}, J_{29 \mathrm{a}, 30}=5.4 \mathrm{~Hz}$ and $J_{29 \mathrm{~b}, 30}=7.4 \mathrm{~Hz}$ pointed out the transoidal geometry of the $\mathrm{C}-27 \sim \mathrm{C}-30$ fragment. It should be mentioned that the difference between $J_{29 \mathrm{a}, 30}$ and $J_{29 \mathrm{~b}, 30}$ could imply that the planes of tetraene and diene chromophores were slightly deviated from ideal coplanarity.

The chair conformation of the glycosidically bound mycosamine (Fig. 2), belonging to the $D$-series established earlier ${ }^{3,4)}$, was pointed out by a set of vicinal coupling constants: $J_{1^{\prime}, 2^{\prime}} \sim 0 \mathrm{~Hz}, J_{2^{\prime}, 3^{\prime}}=3.3 \mathrm{~Hz}, J_{3^{\prime}, 4^{\prime}}=$ $9.5 \mathrm{~Hz}, J_{4^{\prime}, 5^{\prime}}=9.3 \mathrm{~Hz}$. Thus the $\mathrm{H}-3^{\prime}, \mathrm{H}-4^{\prime}$ and $\mathrm{H}-5^{\prime}$ were found to be in axial positions while the $\mathrm{H}-2^{\prime}$ was equatorial. The axial position of $1^{\prime}-\mathrm{H}$ resulted from strong $\mathrm{H}-\mathbf{1}^{\prime} / \mathrm{H}-3^{\prime}$ and $\mathrm{H}-\mathbf{1}^{\prime} / \mathrm{H}-5^{\prime}$ ROE's (Table 1). This pointed out the $\beta$-configuration of the glycosidic bond.

The conformation of the C-13~C-19 fragment, (Fig. 2), was derived as follows: The proposed chair conformation of the hemiketal ring ( $\mathrm{C}-13 \sim \mathrm{C}-17$ ) resulted from the vicinal coupling constants: $J_{14 \mathrm{a}, 15}=11.5 \mathrm{~Hz}, J_{15,16}=$ 10.1 Hz and $J_{16,17}=10.1 \mathrm{~Hz}$ indicating axial positions of the appropriate protons, while the $\mathrm{H}-12 \mathrm{a} / \mathrm{H}-14 \mathrm{a}$ and

Fig. 2. The stereostructure of $3^{\prime}-N$-acetylamphotericin A methoxycarbonylmethylamide with diagnostic ROE's depicted as biderectional arrows:


H-12b/H-14b ROE's defined equatorial position of the $\mathrm{C}-12$ substituent (Fig. 2). This assigment was further supported by the strong ROE's: H-14a/H-16 and H-16/ $\mathrm{H}-18 \mathrm{a}$ observed in the ROESY spectrum. The geometry of the $\mathrm{C}-17 \sim \mathrm{C}-19$ fragment was reflected by $J_{17,18 \mathrm{a}}=$ $9.0 \mathrm{~Hz}, J_{17,18 \mathrm{~b}}=7.0 \mathrm{~Hz}, J_{18 \mathrm{a}, 19}=3.8 \mathrm{~Hz}$ and $J_{18 \mathrm{~b}, 19} \sim 0$ Hz . The configuration of the $\mathrm{C}-13 \sim \mathrm{C}-19$ fragment was then correlated with that of the mycosaminyl substituent by the $\mathrm{H}-1^{\prime} / \mathrm{H}-19, \mathrm{H}-1^{\prime} / \mathrm{H}-18 \mathrm{~b}$ and $\mathrm{H}-2^{\prime} / \mathrm{H}-17$ ROE's. Thus the absolute configuration of the $\mathrm{C}-13 \sim \mathrm{C}-19$ fragment was established as $13 R, 15 S, 16 R, 17 S$ and $19 R$.
The conformation of the $\mathrm{C}-13 \sim \mathrm{C}-19$ fragment associated with $\mathrm{H}-17 / \mathrm{H}-20$ and $\mathrm{H}-19 / \mathrm{H}-21$ ROE's allocated the C-12 equatorial substituent of the hemiketal ring above the tetraene chromophore plane.

The conformation of the $\mathrm{C}-8 \sim \mathrm{C}-12$ fragment was established based upon the vicinal coupling constants: $J_{9,10 \mathrm{~b}}=9.8 \mathrm{~Hz}$ and $J_{11,12 \mathrm{~b}}=9.9 \mathrm{~Hz}$ and the ROE's $\mathrm{H}-9 / \mathrm{H}-11$ and $\mathrm{H}-8 / \mathrm{H}-10 \mathrm{a}$ which pointed out "full stretched" conformation of this fragment. Consequently, the fragment $\mathrm{C}-8 \sim \mathrm{C}-12$ was found to be allocated above the tetraene chromophore plane due to the ROE's: $\mathrm{H}-11 / \mathrm{H}-20, \mathrm{H}-11 / \mathrm{H}-22$ and $\mathrm{H}-9 / \mathrm{H}-24$.
The C-2~C-5 fragment was found to adopt also "full stretched" conformation reflected by vicinal coupling constants: $J_{2 \mathrm{~b}, 3}=8.8 \mathrm{~Hz}, J_{3,4 \mathrm{~b}}=9.0 \mathrm{~Hz}$ and $J_{4 \mathrm{~b}, 5}=8.4 \mathrm{~Hz}$ and ROE's: $\mathrm{H}-2 \mathrm{~b} / \mathrm{H}-4 \mathrm{~b}$ and H-3/H-5.

The vicinal coupling constants $J_{5,6 \mathrm{a}}=3.8 \mathrm{~Hz}, J_{5,6 \mathrm{~b}}=$ $8.1 \mathrm{~Hz}, J_{6 \mathrm{a}, 7 \mathrm{~b}} \sim 9.0 \mathrm{~Hz}, J_{7 \mathrm{a}, 8}=3.0 \mathrm{~Hz}$ and $J_{7 \mathrm{~b}, 8}=9.4 \mathrm{~Hz}$ pointed out the relative allocation of the fragments $\mathrm{C}-2 \sim \mathrm{C}-5$ and $\mathrm{C}-8 \sim \mathrm{C}-12$.
Thus the $\mathrm{C}-2 \sim \mathrm{C}-12$ fragment was found to adopt a "full stretched" conformation and was allocated above the planes of the tetranene and diene chromophores. This assignment was in full agreement with the observed $\mathrm{H}-11 / \mathrm{H}-20, \mathrm{H}-11 / \mathrm{H}-22, \mathrm{H}-9 / \mathrm{H}-24, \mathrm{H}-7 \mathrm{~b} / \mathrm{H}-26$ and $\mathrm{H}-5 /$ H-28ab ROE's which are depicted in Fig. 2.

The absolute configuration of $\mathrm{C}-2 \sim \mathrm{C}-12$ fragment was assigned as $3 R, 5 S, 8 R, 9 S$ and $11 R$ based upon correlation with $\mathrm{C}-13 \sim \mathrm{C}-17$ fragment by $\mathrm{H}-12 \mathrm{a} / \mathrm{H}-14 \mathrm{a}$ and H-12b/H-14b ROE's.
The geometry of the $\mathrm{C}-33 \sim \mathrm{C}-37$ fragment was derived from vicinal coupling constants: $J_{33,34}=8.5 \mathrm{~Hz}, J_{34,35}=$ $7.7 \mathrm{~Hz}, J_{35,36}=4.0 \mathrm{~Hz}, J_{36,37}=3.0 \mathrm{~Hz}$ associated with the ROE's H-33/H-35, H-33/H-36, H-35/Me36 and Me36/ Me37 observed in ROESY spectrum of 2. The absolute configuration of this fragment was deduced as follows. The C-2~C-12 fragment was found to be allocated above the planes of the tetrane and diene chromophores. This fact determined the condition for closing the macrolac-
tone ring which required that the $\mathrm{C}-35, \mathrm{C}-36$ and $\mathrm{C}-37$ had to be located also above the planes of the diene and tetraene chromophores. This requirement could be covered by only one enantiomer of the $\mathrm{C}-33 \sim \mathrm{C}-37$ fragment. Thus the absolute configuration of the $\mathrm{C}-33 \sim$ C-37 fragment was assigned as $34 S, 35 R, 36 R$ and $37 S$.

## Experimental

Amphotericin A (1)
Amphotericin A in pure, crystalline form with $\mathrm{E}_{1}^{1 \%} \mathrm{~cm}=$ 1.350 at 304 nm (methanol), was supplied by E.R. Squibb and Sons, Princeton, N.J., U.S.A.
$3^{\prime}-N$-Acetylamphotericin A Methoxycarbonylmethylamide (2)
This derivative was obtained by the procedures previously described ${ }^{5,6)}$. The products were purificated by flash chromatography on MN -Kieselgel with the solvent system chloroform-methanol-water, 5:0.5:0.05 (v/v) and $5: 1: 0.1(\mathrm{v} / \mathrm{v})$.

## NMR Spectra

Spectra were recorded with a Varian Unity 500 Plus spectrometer in the solvent system, pyridine- $d_{5}$ -methanol- $d_{4} 9: 1(\mathrm{v} / \mathrm{v})$, with a sample concentration of $10 \mathrm{mg} / \mathrm{ml}$. The spectra of ${ }^{13} \mathrm{C}$ and $1 \mathrm{D}-{ }^{1} \mathrm{H}$ were collected with standard parameters. $2 \mathrm{D}-{ }^{1} \mathrm{H}$ spectra were measured in phase sensitive mode with a spectral width of 4723.7 Hz .

Double-quantum filtered COSY spectra were acquired in $4094 \times 800$ matrix and were processed in $8 \mathrm{~K} \times 4 \mathrm{~K}$ matrix (digital resolution 1.0 Hz and 2.0 Hz ).

ROESY spectra were acquired with a mix time of 0.4 seconds in $2048 \times 386$ matrix with 16 accumulations per increment in $2 \mathrm{~K} \times 1 \mathrm{~K}$ matrix (digital resolution 2.0 Hz and 4.1 Hz ).
HSQC and HMBC spectra were performed with pulse field gradients. The HSQC spectrum was acquired in phase sensitive mode. The spectral windows for ${ }^{1} \mathrm{H}$ and ${ }^{13} \mathrm{C}$ axes were 3399.6 Hz and 12574.7 Hz , respectively. Data were collected in $1024 \times 200$ matrix and processed in $1 \mathrm{~K} \times 1 \mathrm{~K}$ matrix. The HMBC spectrum was acquired
in absolute value mode with broadband decoupling. The spectral windows for ${ }^{1} \mathrm{H}$ and ${ }^{13} \mathrm{C}$ axes were 3399.6 Hz and 12574.7 Hz , respectively. Data were collected in $832 \times 256$ matrix and processed in $1 \mathrm{~K} \times 1 \mathrm{~K}$ matrix.
${ }^{1} \mathrm{H}$ NMR spectra were also recorded with methanol- $d_{4}$ solvent.

1D-TOCSY spectra were recorded with tophat shaped pulse $(100 \sim 300 \mathrm{~ms})$ with a mix time of $0.005 \sim 0.090$ seconds.

NOESY spectrum was acquired with mix time 0.3 seconds, with spectral width of 4096.3 Hz , in $2048 \times 341$ matrix and was processed in $2 \mathrm{~K} \times 1 \mathrm{~K}$ matrix.

## Acknowledgements

This work was supported by the State Committee for Scientific Research, Warsaw, Poland (grant No. 406329101).

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