

Molecular and Crystal Structure of a Nonionic Detergent: Nonanoyl-*N*-methylglucamide

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In crystals of the nonionic detergent nonanoyl-*N*-methylglucamide, molecules are packed parallel in sheets with adjacent sheets being arranged in a head-to-tail fashion; an unusual conformation of the polar head moiety explains why this compound forms micelles whereas the related alkylglyconamides do not.

Crystallization of membrane proteins depends critically on the detergent used. Small nonionic detergents have been found useful for growing three-dimensional crystals of such proteins.^{1,2} In our attempts to improve the size and diffraction of crystals³ of the mitochondrial pore-protein from *Neurospora crassa*,⁴ we use, among others, nonanoyl-*N*-methylglucamide (**1**) (MEGA-9)⁵ as detergent. At concentrations above the critical micelle concentration (c.m.c.), this amphiphile readily forms micelles at room temperature whereas the related compound *n*-octyl-*D*-gluconamide (**2**) does so only when heated to 90 °C.⁶ The crystal structure analysis of the latter molecule, which differs from MEGA-9 mainly by a reversed amide group and by lacking the *N*-methyl group, revealed an all-*trans* conformation of both the alkyl chain and the glucose moiety.⁷ In order to shed light on the structural

foundations of the different properties of MEGA-9, we have now determined its molecular and crystal structure by *X*-ray

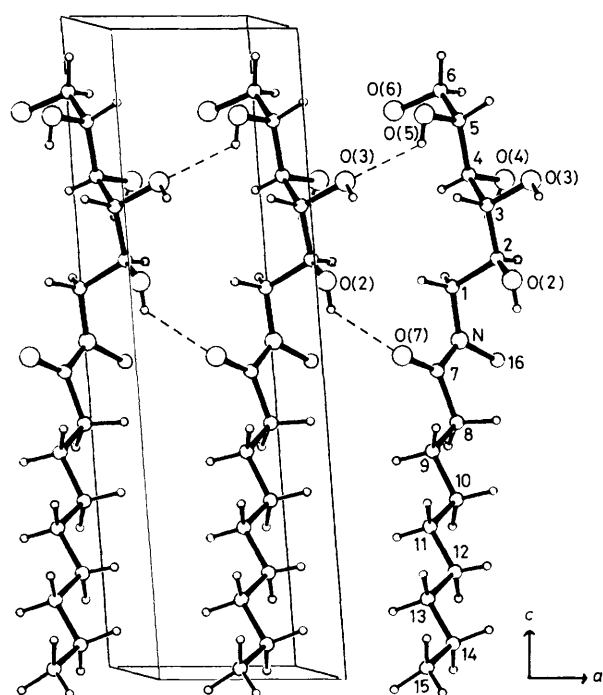
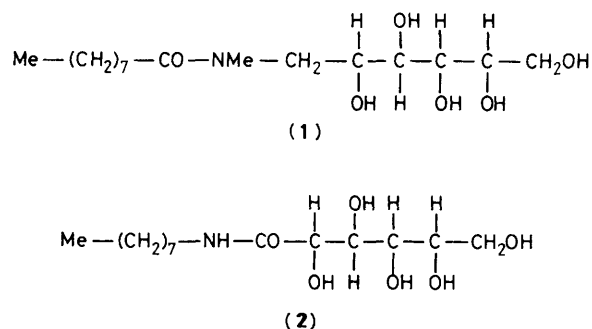


Figure 1. The crystal structure of MEGA-9 (**1**) with the atom numbering scheme, showing the parallel packing of molecules.

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Table 1. Hydrogen bond geometry in crystals of nonanoyl-*N*-methylglucamide (MEGA-9). Distances in Å, angles in °; standard deviations in parentheses; D, donor atom; A, acceptor atom.

D-H...A	Translation	D...A distance	H...A distance	D-H-A angle
O(2)-H(O2)...O(7)	1 + x, y, z	2.85(1)	2.25(2)	132.2(8)
O(3)-H(O3)...O(4)	x, 1 + y, z	2.75(2)	1.74(3)	154.4(9)
O(4)-H(O4)...O(2)	x, 1 - y, z	3.38(2)	2.40(3)	148.6(9)
O(5)-H(O5)...O(3)	1 - x, y, z	2.76(2)	2.44(3)	122.5(9)
O(6)-H(O6)...O(5)	x, 1 - y, z	2.76(1)	1.81(2)	166.4(8)

diffraction.‡ Owing to inherent difficulties in obtaining single crystals of nonionic detergents, the only other *X*-ray structure reported of such a compound is, to the best of our knowledge, that of 1-decyl- α -D-glucopyranoside.⁸

The molecular structure of MEGA-9 is shown in Figure 1 together with the atom numbering scheme. As in *n*-octyl-D-glucuronamide (**2**),⁷ the alkyl chain is in an all-*trans* conformation which includes the amide bond. However, there is an important difference in the polar head moiety, where a kink is introduced in MEGA-9 by the synclinal torsion angle C(1)-C(2)-C(3)-C(4) [$-57(1)^\circ$].

The molecules of MEGA-9 are packed parallel in sheets in the *ac*-plane, resulting in a head-to-tail packing of molecules (along *c* axis) in adjacent sheets. The same unusual feature has been observed with octyl-D-glucuronamide (**2**).⁷ However, the hydrogen bonding is quite different in the two structures owing to the methylation of the amide group and the different

conformation of the polar head moiety in MEGA-9. The latter is involved in five intermolecular hydrogen bonds three of which are rather weak (Table 1). In contrast, hydrogen bonding between the polar head groups in adjacent octyl-D-glucuronamide molecules (**2**) is much stronger⁷ favouring the aggregation to aqueous gels observed with this compound.⁶

From this observation it appears that not too strong hydrogen bonding of the polar head moieties is a pre-requisite for micelle formation.

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References

- 1 H. Michel, *Trends Biochem. Sci.*, 1983, **8**, 56.
- 2 R. M. Garavito and J. A. Jenkins, in 'Structure and Function of Membrane Proteins,' eds. E. Quagliariello and F. Palmieri, Elsevier, 1983, pp. 205-210.
- 3 A. Müller-Fahrnow, R. Hilgenfeld, W. Saenger, H. Freitag, and W. Neupert, unpublished work.
- 4 H. Freitag, W. Neupert, and R. Benz, *Eur. J. Biochem.*, 1982, **123**, 629.
- 5 J. E. K. Hildreth, *Biochem. J.*, 1982, **207**, 363.
- 6 B. Pfannemüller and W. Welte, *Chem. Phys. Lipids*, 1985, **37**, 227.
- 7 V. Zabel, A. Müller-Fahrnow, R. Hilgenfeld, W. Saenger, B. Pfannemüller, V. Enkelmann, and W. Welte, *Chem. Phys. Lipids*, 1986, **39**, 313.
- 8 P. C. Moews and J. R. Knox, *J. Am. Chem. Soc.*, 1976, **98**, 6628.
- 9 M. Steifa and W. Saenger, *Acta Crystallogr., Sect. A*, 1984, **40**, C409.

‡ *Crystal data*: crystals of MEGA-9 (Oxyl, Bobingen, FRG) grown from acetone-methanol (1:1), $C_{16}H_{33}NO_6$, $M = 335$, triclinic, space group *P*1, $a = 4.985(2)$, $b = 5.603(2)$, $c = 17.449(7)$ Å, $\alpha = 85.4(3)$, $\beta = 86.0(3)$, $\gamma = 76.2(3)^\circ$, $U = 468.5$ Å³, $Z = 1$, $D_c = 1.19$ g cm⁻³, $\mu(\text{Cu-K}\alpha) = 0.67$ cm⁻¹.

1445 unique intensity data ($2\theta_{\text{max}} = 120^\circ$) were recorded on a redesigned⁹ Stoe four-circle diffractometer using Ni-filtered Cu- $K\alpha$ -radiation and a $2\theta/\omega$ scan technique. The structure was solved by a combination of direct methods and difference Fourier techniques, and refined by full-matrix least-squares (all non-hydrogen atoms were anisotropic). All hydrogen atoms could be located from difference syntheses and isotropically refined. The final conventional *R*-factor was 0.055.

Atomic co-ordinates, bond lengths and angles, and thermal parameters have been deposited at the Cambridge Crystallographic Data Centre. See Notice to Authors, Issue No. 1, 1986.