

Figure 2
The unit cell in the title compound. Only water H atoms are shown.

1.5180 (11) Å. Most intrachain bond angles of the hydrocarbon chain fall within the narrow range 113.7 (3)–114.8 (3)°, with N1–C1–C3 slightly larger and C1–C2–C3 slightly smaller. The N–O distance compares with a value of 1.404 Å (corrected for libration) in trimethylamine oxide (Caron *et al.*, 1964) and a value of 1.399 Å (mean of three) in *N,N*-dimethylethanolamine *N*-oxide (Maia *et al.*, 1984).

The unit cell is shown in Fig. 2, which illustrates the hydrogen-bonded regions near $x = 0$, where the water molecules associate with the head-to-head hydrophilic ends of the fatty amine *N*-oxide molecules. In the interior of the cell, the parallel hydrocarbon chains interdigitate along the [010] direction. The hydrophilic region is shown in Fig. 3. Each water molecule donates two hydrogen bonds and accepts one, linking the *N*-oxide groups into six-oxygen centrosymmetric rings. These rings are further linked by $\text{H}_2\text{O} \cdots \text{H}_2\text{O}$ hydrogen bonds to form two-dimensional arrays (see Table 2 for hydrogen-bond details).

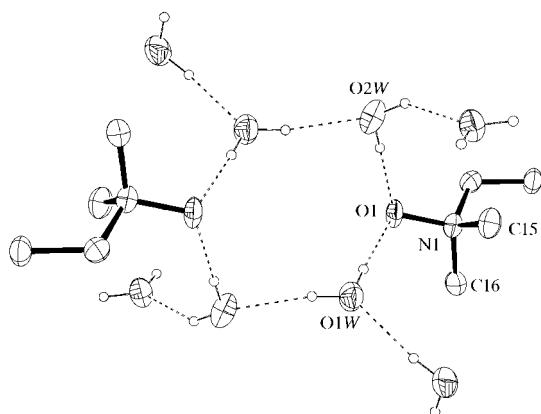


Figure 3
The hydrogen bonding in the title compound. Only the first two C atoms of the C_{14} chain are illustrated.

Experimental

The title compound was prepared by reacting liquid *N,N*-dimethyl-*n*-tetradecylamine with 70% hydrogen peroxide in the presence of CO_2 as catalyst. A blanket of CO_2 was maintained throughout the addition

of the peroxide to the liquid amine. The amine oxidation process is strongly exothermic and the reaction temperature was regulated by the rate of peroxide addition (Elnagar, 2000). The resulting soft solid (m.p. 317 K) was crystallized from ethyl methyl ketone.

Crystal data

$\text{C}_{16}\text{H}_{35}\text{NO} \cdot 2\text{H}_2\text{O}$
 $M_r = 293.48$
Monoclinic, $P2_1/c$
 $a = 22.782$ (8) Å
 $b = 8.110$ (5) Å
 $c = 9.995$ (5) Å
 $\beta = 91.19$ (2)°
 $V = 1846.3$ (16) Å³
 $Z = 4$

$D_x = 1.056$ Mg m⁻³
Mo $K\alpha$ radiation
Cell parameters from 3471 reflections
 $\theta = 2.5$ – 26.0 °
 $\mu = 0.07$ mm⁻¹
 $T = 100$ K
Plate, colorless
 $0.48 \times 0.17 \times 0.03$ mm

Data collection

Nonius KappaCCD diffractometer
(with an Oxford Cryosystems
Cryostream cooler)
 ω scans with κ offsets
15 919 measured reflections
3508 independent reflections

1746 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.076$
 $\theta_{\text{max}} = 26.0$ °
 $h = -28 \rightarrow 28$
 $k = -10 \rightarrow 10$
 $l = -11 \rightarrow 11$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.086$
 $wR(F^2) = 0.195$
 $S = 1.06$
3508 reflections
196 parameters
H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0530P)^2 + 1.8197P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.004$
 $\Delta\rho_{\text{max}} = 0.32$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.20$ e Å⁻³

Table 1

Selected geometric parameters (Å, °).

| | | | |
|-------------|-----------|-------------|-----------|
| O1–N1 | 1.411 (3) | N1–C15 | 1.489 (4) |
| N1–C16 | 1.477 (4) | N1–C1 | 1.494 (4) |
| O1–N1–C16 | 108.0 (2) | C16–N1–C1 | 111.8 (3) |
| O1–N1–C15 | 109.7 (2) | C15–N1–C1 | 111.6 (3) |
| C16–N1–C15 | 109.4 (3) | N1–C1–C2 | 117.0 (2) |
| O1–N1–C1 | 106.3 (2) | C3–C2–C1 | 109.5 (2) |
| O1–N1–C1–C2 | 173.4 (3) | N1–C1–C2–C3 | 172.1 (3) |

Table 2

Hydrogen-bonding geometry (Å, °).

| $D-H \cdots A$ | $D-H$ | $H \cdots A$ | $D \cdots A$ | $D-H \cdots A$ |
|---|-------|--------------|--------------|----------------|
| O1W–H1W ⁱ ···O1 | 0.88 | 1.85 | 2.727 (3) | 174 |
| O1W–H2W ⁱ ···O2W ⁱ | 0.87 | 1.97 | 2.832 (4) | 170 |
| O2W–H3W ⁱ ···O1 | 0.88 | 1.87 | 2.741 (4) | 177 |
| O2W–H4W ⁱ ···O1W ⁱⁱ | 0.88 | 2.07 | 2.829 (4) | 144 |

Symmetry codes: (i) $2 - x, 1 - y, -z$; (ii) $2 - x, y - \frac{1}{2}, \frac{1}{2} - z$.

H atoms on C atoms were placed in calculated positions, with C–H distances in the range 0.98–0.99 Å, and thereafter treated as riding. A torsional parameter was refined for each methyl group. Water H atoms were located in difference maps, and their O–H distances were restrained. For all H atoms, $U_{\text{iso}} = 1.5U_{\text{eq}}$ of the attached atom for methyl and water H atoms or $1.2U_{\text{eq}}$ for all other H atoms.

Data collection: *COLLECT* (Nonius, 2000); cell refinement: *HKL SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *HKL DENZO* (Otwinowski & Minor, 1997) and *SCALEPACK*; program(s) used to solve structure: *SIR97* (Altomare *et al.*, 1999); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997).

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: SQ1004). Services for accessing these data are described at the back of the journal.

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