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# 2-(2-Naphthyloxy)acetate derivatives. II. A new class of antiamnesic agents 

S. Thamotharan, ${ }^{\text {a }}$ V. Parthasarathi, ${ }^{\text {a }}{ }^{\text {P P. Gupta, }}{ }^{\text {b }}$ D. P. Jindal, ${ }^{\text {b }}+$ P. Piplani ${ }^{\text {b }}$ and Anthony Linden ${ }^{\text {c }}$

${ }^{\text {a }}$ Department of Physics, Bharathidasan University, Tiruchirappalli 620 024, India, ${ }^{\text {b }}$ University Institute of Pharmaceutical Sciences, Panjab University, Chandigarh 160014 , India, and ${ }^{\text {c Institute of Organic Chemistry, University of Zürich, }}$ Winterthurerstrasse 190, CH-8057 Zürich, Switzerland Correspondence e-mail: vpsarati@yahoo.com

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The title compounds, 4-(2-naphthyloxymethylcarbonyl)morpholine, $\mathrm{C}_{16} \mathrm{H}_{17} \mathrm{NO}_{3}$, (I), and 4-methyl-1-(2-naphthyloxymethylcarbonyl)piperazine, $\mathrm{C}_{17} \mathrm{H}_{20} \mathrm{~N}_{2} \mathrm{O}_{2}$, (II), are potential antiamnesics. The morpholine ring in (I) and the piperazine ring in (II) adopt chair conformations. In (I), the molecules are linked by weak intermolecular $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ interactions into chains that have a graph-set motif of $C(10)$, while in (II), the molecules are linked by weak intermolecular $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ interactions that generate two $C(7)$ graph-set motifs. The dihedral angle between the naphthalene moiety and the best plane through the morpholine ring is $20.62(4)^{\circ}$ in (I), while the naphthalene moiety is oriented nearly perpendicular to the mean plane of the piperazine ring in (II).

## Comment

The conformation of molecules with antiamnesic activity has attracted considerable interest (Amato et al., 1991), and the present structure determinations are part of our research program on biologically active 2-(2-naphthyloxy)acetate derivatives. We report here the preparation and the X-ray crystal structures of 4-(2-naphthyloxymethylcarbonyl)morpholine, (I), and 1-(2-naphthyloxymethylcarbonyl)-4-methylpiperazine, (II), both of which are new classes of antiamnesic agents (Thamotharan et al., 2003, and references therein).

(I)

(II)

Views of the molecules of (I) and (II), with the atomic numbering schemes, are shown in Figs. 1 and 2, respectively.

[^0]The corresponding bond lengths and angles in (I) and (II) are essentially equivalent and are comparable to those in the related structures 1-(2-naphthyloxymethylcarbonyl)piperidine and 3-methyl-1-(2-naphthyloxymethylcarbonyl)piperidine (Thamotharan et al., 2003). In the molecule of (I), the central C2/O11/C12/C13/N14 fragment is planar, with a maximum deviation of 0.023 (1) $\AA$ for atom C13. In contrast, only the O11/C12/C13/N14 segment is planar in (II), while atom C2 is twisted about the $\mathrm{O} 11-\mathrm{C} 12$ bond so that it lies almost perpendicular to the O11/C12/C13/N14 plane (Table 3). In (I), the central unit is almost coplanar with the naphthalene moiety, the angle between the planes being $3.67(7)^{\circ}$, while the dihedral angle between the central unit and the naphthalene moiety in (II) is $76.94(15)^{\circ}$. The central unit is twisted $27.79(14)^{\circ}$ out of the mean plane of the piperazine ring in (II), whereas it is twisted $21.73(6)^{\circ}$ out of the best plane passing through the morpholine ring in (I). The $\mathrm{C} 2-\mathrm{O} 11-\mathrm{C} 12-\mathrm{C} 13$ and $\mathrm{O} 11-\mathrm{C} 12-\mathrm{C} 13-\mathrm{N} 14$ torsion angles show that the central unit has an antiperiplanar conformation in (I), while the corresponding torsion angles in (II) represent synclinal and antiperiplanar conformations, respectively.

The morpholine ring in (I) has a chair conformation [the Cremer \& Pople (1975) puckering parameters are $Q=0.550(1) \AA, \quad q_{2}=0.045(1) \AA, \quad q_{3}=0.548(1) \AA, \quad \theta=$ $4.7(1)^{\circ}$ and $\varphi_{2}=345.8(19)^{\circ}$ for the atom sequence O17$\mathrm{C} 16-\mathrm{C} 15-\mathrm{N} 14-\mathrm{C} 19-\mathrm{C} 18]$. The dihedral angle between the naphthalene moiety and the morpholine ring is $20.62(4)^{\circ}$. In (II), the puckering parameters of the piperazine ring $\left[Q=0.5585(3) \AA, \quad q_{2}=0.016(3) \AA, \quad q_{3}=0.558(3) \AA, \quad \theta=\right.$ $0.7(3)^{\circ}$ and $\varphi_{2}=131(10)^{\circ}$ for the atom sequence $\mathrm{N} 14-\mathrm{C} 15-$ $\mathrm{C} 16-\mathrm{N} 17-\mathrm{C} 18-\mathrm{C} 19$ ] are indicative of a chair conformation, and the methyl substituent at atom N 17 is in an equatorial position. The sum of the angles around N 17 is $331^{\circ}$, revealing an essentially pyramidal geometry at atom N17. In contrast,


Figure 1
A view of the molecule of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the $50 \%$ probability level and H atoms are represented by circles of arbitrary radii.


Figure 2
A view of the molecule of (II), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the $50 \%$ probability level and H atoms are represented by circles of arbitrary radii.
atom N14 in (I) and (II) does not exhibit a pyramidal geometry because of the amide-type cojugation with carbonyl atom O13. The sum of the angles around atom N14 in (I) and (II) is $360^{\circ}$ (Tables 1 and 3). The naphthalene moiety is oriented nearly perpendicular to the mean plane of the piperazine ring.

The exocyclic $\mathrm{C} 1-\mathrm{C} 2-\mathrm{O} 11$ and $\mathrm{C} 13-\mathrm{N} 14-\mathrm{C} 15$ angles deviate significantly from the usual value of $120^{\circ}$ (Tables 1 and 3), perhaps as a result of steric repulsion $[\mathrm{H} 1 \cdots \mathrm{H} 121=2.21 \AA$ $(2.10 \AA), \mathrm{H} 121 \cdots \mathrm{H} 151=2.25 \AA(\mathrm{H} 121 \cdots \mathrm{H} 152=2.26 \AA)$ and $\mathrm{H} 122 \cdots \mathrm{H} 151=2.17 \AA(\mathrm{H} 122 \cdots \mathrm{H} 152=2.11 \AA)$; the values in parentheses correspond to (II)].

As can be seen from Table 2, in (I), atom C1 acts as a donor in a weak intermolecular $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ interaction with atom O 17 of an adjacent molecule. This interaction links the molecules into chains that run parallel to the $c$ axis and have a graph-set motif of $C(10)$ (Bernstein et al., 1995).

In (II), atom C1 acts as a donor for a weak intermolecular $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ interaction with carbonyl atom O 13 of an adjacent molecule (Table 4). This interaction produces a continuous chain that runs parallel to the $c$ axis and has a graph-set motif of $C(7)$. In addition, atom C 5 is involved in an intermolecular $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ interaction with atom O 11 of a different molecule. This interaction links the molecules into a continuous chain that runs parallel to the $a$ axis and has a graph-set motif of $C(7)$ (Bernstein et al., 1995).

## Experimental

For the preparation of (I), methyl 2-(2-naphthyloxy)acetate ( 0.5 g ) was reacted with morpholine. The oily product obtained was treated with water, and the resulting precipitate was filtered off, dried and crystallized from acetone to afford crystals of (I) (yield 0.532 g ,
$84.8 \%$; m.p. $416-418 \mathrm{~K}$ ). For the preparation of (II), methyl 2-(2naphthyloxy)acetate ( 0.5 g ) was reacted with $N$-methylpiperazine. The oily product obtained was treated with water, and the resulting precipitate was filtered off, dried and crystallized from ethanol to afford crystals of (II) (yield $0.514 \mathrm{~g}, 78.17 \%$; m.p. 383-385 K).

## Compound (I)

Crystal data
$\mathrm{C}_{16} \mathrm{H}_{17} \mathrm{NO}_{3}$
$M_{r}=271.31$
Monoclinic, $P 2_{1 / n} n$
$a=10.5594$ (2) $\AA$
$b=9.2956$ (2) $\AA$
$c=14.1012$ (3) $\AA$
$\beta=109.0917$ (9) ${ }^{\circ}$
$V=1307.98(5) \AA^{3}$
$Z=4$
$D_{x}=1.378 \mathrm{Mg} \mathrm{m}^{-3}$
Mo $K \alpha$ radiation
Cell parameters from 4035 reflections
$\theta=2.0-30.0^{\circ}$
$\mu=0.10 \mathrm{~mm}^{-1}$
$T=160$ (2) K
Prism, colourless
$0.28 \times 0.25 \times 0.23 \mathrm{~mm}$
Data collection
Nonius KappaCCD diffractometer $\varphi$ and $\omega$ scans with $\kappa$ offsets
37467 measured reflections 3815 independent reflections 2574 reflections with $I>2 \sigma(I)$

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.051$
$w R\left(F^{2}\right)=0.148$
$S=1.03$
3815 reflections
181 parameters
H -atom parameters constrained

Table 1
Selected bond and torsion angles ( ${ }^{\circ}$ ) for (I).

| C13-N14-C15 | $125.35(11)$ | C15-N14-C19 | $113.52(10)$ |
| :--- | :--- | :--- | :--- |
| C13-N14-C19 | $120.73(10)$ | C1-C2-O11 | $125.24(11)$ |
|  |  |  |  |
| C2-O11-C12-C13 | $177.19(10)$ | $\mathrm{O} 11-\mathrm{C} 12-\mathrm{C} 13-\mathrm{N} 14$ | $178.29(10)$ |

Table 2
Hydrogen-bonding geometry $\left({ }_{\mathrm{A}},^{\circ}\right)$ for (I).

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| $\mathrm{C} 1-\mathrm{H} 1 \cdots \mathrm{O}^{\mathrm{C}} 7^{\mathrm{i}}$ | 0.95 | 2.42 | $3.3359(15)$ | 161 |

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

## Compound (II)

## Crystal data

| $\mathrm{C}_{17} \mathrm{H}_{20} \mathrm{~N}_{2} \mathrm{O}_{2}$ | Mo $K \alpha$ radiation |
| :--- | :--- |
| $M_{r}=284.35$ | Cell parameters from 1547 |
| Orthorhombic, $P c a 2_{1}$ | reflections |
| $a=13.7560(3) \AA$ | $\theta=2.0-25.0^{\circ}$ |
| $b=12.7389(2) \AA$ | $\mu=0.08 \mathrm{~mm}^{-1}$ |
| $c=8.5031(2) \AA$ | $T=160(2) \mathrm{K}$ |
| $V=1490.05(5) \AA^{3}$ | Plate, colourless |
| $Z=4$ | $0.23 \times 0.13 \times 0.07 \mathrm{~mm}$ |
| $D_{x}=1.268 \mathrm{Mg} \mathrm{m}^{-3}$ |  |

## Data collection

Nonius KappaCCD diffractometer
$\omega$ scans with $\kappa$ offsets
17668 measured reflections
1414 independent reflections
1169 reflections with $I>2 \sigma(I)$

$$
\begin{aligned}
& R_{\text {int }}=0.079 \\
& \theta_{\max }=25.0^{\circ} \\
& h=-16 \rightarrow 16 \\
& k=-15 \rightarrow 15
\end{aligned}
$$

## Refinement

Refinement on $F^{2}$
$w=1 /\left[\sigma^{2}\left(F_{o}^{2}\right)+(0.0575 P)^{2}\right]$
where $P=\left(F_{o}^{2}+2 F_{c}^{2}\right) / 3$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.039$
$w R\left(F^{2}\right)=0.095$
$S=1.06$
1414 reflections
192 parameters
H -atom parameters constrained
$(\Delta / \sigma)_{\max }<0.001$
$\Delta \rho_{\text {max }}=0.18 \mathrm{e}^{-3}$
$\Delta \rho_{\min }=-0.16$ e $\AA^{-3}$
Extinction correction: SHELXL97
Extinction coefficient: 0.014 (3)

Table 3
Selected bond and torsion angles ( ${ }^{\circ}$ ) for (II).

| C13-N14-C15 | $126.0(2)$ | C18-N17-C20 | $109.9(2)$ |
| :--- | ---: | :--- | :--- |
| C13-N14-C19 | $121.2(2)$ | C16-N17-C20 | $110.7(2)$ |
| C15-N14-C19 | $112.4(2)$ | C1-C2-O11 | $125.1(2)$ |
| C18-N17-C16 | $110.0(2)$ |  |  |
| C2-O11-C12-C13 | $84.0(3)$ | O11-C12-C13-N14 | $175.6(2)$ |

Table 4
Hydrogen-bonding geometry $\left({ }^{\circ},{ }^{\circ}\right)$ for (II).

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| C1-H1 $\cdots \mathrm{O}_{1} 3^{\text {ii }}$ | 0.95 | 2.55 | $3.471(3)$ | 163 |
| C5-H5 $\mathrm{O}_{1} 1^{\text {iii }}$ | 0.95 | 2.49 | $3.236(3)$ | 135 |

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

For compound (I), all H atoms were placed in idealized positions ( $\mathrm{C}-\mathrm{H}=0.95-0.99 \AA$ ) and constrained to ride on their parent atoms, with $U_{\text {iso }}(\mathrm{H})$ values of $1.2 U_{\text {eq }}(\mathrm{C})$. For compound (II), methyl H atoms were constrained to an ideal geometry $(\mathrm{C}-\mathrm{H}=0.98 \AA)$, with $U_{\text {iso }}(\mathrm{H})$ values of $1.5 U_{\text {eq }}(\mathrm{C})$, but were allowed to rotate freely about the $\mathrm{C}-\mathrm{C}$ bond. All remaining H atoms were placed in idealized positions ( $\mathrm{C}-$ $\mathrm{H}=0.95-0.99 \mathrm{~A}$ ) and constrained to ride on their parent atoms, with $U_{\text {iso }}(\mathrm{H})$ values of $1.2 U_{\text {eq }}(\mathrm{C})$. Although the molecule is achiral, the structure possesses a polar axis. Because of the absence of any
significant anomalous scatterers in the compound, attempts to confirm the absolute structure by refinement of the Flack (1983) parameter in the presence of 1099 sets of Friedel equivalents led to an inconclusive value (Flack \& Bernardinelli, 2000) of -0.1 (15) for this parameter. Therefore, the absolute direction of the polar axis was assigned arbitrarily and the Friedel pairs were merged before the final refinement. Reflection 110 was partially obscured by the beam stop and was omitted.

For both compounds, data collection: COLLECT (Nonius, 2000); cell refinement: $D E N Z O-S M N$ (Otwinowski \& Minor, 1997); data reduction: $D E N Z O-S M N$ and SCALEPACK (Otwinowski \& Minor, 1997); program(s) used to solve structure: SIR92 (Altomare et al., 1994); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP-3 for Windows (Farrugia, 1997); software used to prepare material for publication: PARST97 (Nardelli, 1995) and PLATON (Spek, 2003).

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: SK1649). Services for accessing these data are described at the back of the journal.

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[^0]:    $\dagger$ Deceased.

