| NIA—CIA | 1.378 (7) | N3A—C11A | 1.382 (6) |
|---------------|-----------|--------------|-----------|
| N1A-C11A | 1.314 (6) | N3A—C19A | 1.451 (6) |
| N2A—C1A | 1.435 (6) | C2A—C3A | 1.489 (7) |
| N2A—C2A | 1.379 (6) | C3A—C4A | 1.368 (7) |
| N2A—C12A | 1.445 (6) | C3A—C11A | 1.429 (6) |
| N3A—C10A | 1.389 (6) | C4A—C5A | 1.418 (7) |
| C1A—N1A—C11A | 118.4 (4) | C2A—C3A—C11A | 117.6 (4) |
| C1A—N2A—C2A | 123.9 (4) | C3A—C4A—C5A | 120.6 (4) |
| C1A—N2A—C12A | 118.4 (4) | C4AC5AC10A | 117.7 (4) |
| C10A—N3A—C11A | 122.5 (4) | N3A-C10A-C5A | 120.4 (4) |
| C11A—N3A—C19A | 119.2 (4) | N1A—C11A—N3A | 118.1 (4) |
| N1A—C1A—N2A | 119.5 (4) | N1A-C11A-C3A | 125.2 (4) |
| N2A—C2A—C3A | 114.2 (4) | N3A—C11A—C3A | 116.7 (4) |
| C2A—C3A—C4A | 120.3 (4) | | |
| | | | |

Table 1. Selected geometric parameters (Å, °)

The space group was uniquely determined from the systematic absence: 0k0 when k = 2n + 1. Bijvoet pairs were not averaged. The non-H atoms were refined anisotropically and H atoms were placed in calculated positions but not refined.

Data collection: MSCIAFC Diffractometer Control Software (Molecular Structure Corporation, 1992). Cell refinement: MSCIAFC Diffractometer Control Software. Data reduction: TEXSAN PROCESS (Molecular Structure Corporation, 1995). Program(s) used to solve structure: direct methods SHELXS86 (Sheldrick, 1985). Program(s) used to refine structure: TEXSAN LS. Software used to prepare material for publication: TEXSAN FINISH.

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3-[(*Z*)-Piperidin-1-ylmethylidene]-2,3-dihydro-1,4-benzodioxan-2-one

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Abstract

The crystal structure of the title compound has been determined in order to obtain the geometry of one isomer of $C_{14}H_{15}NO_3$. The molecule has the unusual feature of a planar dioxane ring and several bond angles are enlarged due to steric hindrance.

Comment

In connection with our investigations into 1,4-benzodioxane chemistry, the reaction between 1,4-benzodioxane-2-carboxylic acid and some amines has been studied (Ruiz *et al.*, 1996), and the title compound, (I), was isolated. Although spectroscopic data show that (I) is clearly different from its isomer 2-piperidinylcarbonyl-1,4-benzodioxane, X-ray diffraction analysis has permitted the determination of the structure and geometry of this new unexpected compound.



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The 1,4-benzodioxo ($C_6H_4O_2$) fragment of the 1,4benzodioxane moiety has the same geometric values as those observed in a study of 29 structures containing this moiety, obtained from the Cambridge Structural Database (Allen & Kennard, 1993). The O1—C2— C7—O3 torsion angle has a value of $-2.2(3)^\circ$; this value varies between 0.0 and 4.6° in the previously mentioned study.

The dioxane ring is planar (Fig. 1), with the largest deviation from the mean plane being 0.018 (2) Å for the C1 atom. This ring is usually not planar, with atoms C1 and C8 being out of the plane defined by the remaining four atoms, but the C_{sp^2} character of C1 in the title structure produces the planarity of the ring.



Fig. 1. ORTEP (Brueggemann & Schmid, 1990) drawing of (I) showing the atomic numbering scheme. Displacement ellipsoids are drawn at the 50% probability level.

The largest and smallest bond angles [O3-C8-C1 122.71(15)] and $C8-C1-O1 116.92(15)^\circ$, respectively] have also been observed in compounds with the 1,4-benzodioxan-2-yl moiety [average values 122.2(5)] and 114.8(5)°, respectively; Leger, Colleter & Carpy, 1983; Dzvinchuk *et al.*, 1990].

The piperidine ring has the typical chair form, with atoms C11 and C14 0.655 (3) Å out of the plane defined by the remaining four atoms. Atoms O3, C8, C1, C9 and N are in a plane. The steric hindrance between the piperidine group and O3 produces an increase of the C8—C9—N bond angle to 132.4 (2)° and a shortening of the H9…H10A contact distance to 2.07 (4) Å.

Experimental

Single crystals of the title compound were grown from a methylene chloride solution at 269 K.

Crystal data

| $C_{14}H_{15}NO_{3}$ | Mo $K\alpha$ radiation |
|----------------------|-------------------------------|
| $M_r = 245.27$ | $\lambda = 0.71069 \text{ Å}$ |

Monoclinic $P2_1/a$ a = 8.994 (3) Å b = 16.534 (4) Å c = 9.428 (2) Å $\beta = 118.11$ (3)° V = 1236.6 (6) Å³ Z = 4 $D_x = 1.317$ Mg m⁻³ D_m not measured

Data collection

Enraf-Nonius CAD-4 diffractometer ω -2 θ scans Absorption correction: none 3741 measured reflections 3591 independent reflections 2230 reflections with $I > 2\sigma(I)$

Refinement

 $\Delta \rho_{\text{max}} = 0.312 \text{ e } \text{\AA}^{-3}$ $\Delta \rho_{\text{min}} = -0.375 \text{ e } \text{\AA}^{-3}$ Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.066$ $wR(F^2) = 0.312$ Extinction correction: S = 0.985SHELXL93 3591 reflections Extinction coefficient: 202 parameters 0.000(5)H atoms: see below Scattering factors from $w = 1/[\sigma^2(F_o^2) + (0.1659P)^2]$ International Tables for where $P = (F_o^2 + 2F_c^2)/3$ Crystallography (Vol. C) $(\Delta/\sigma)_{\rm max} = 0.001$

Table 1. Selected geometric parameters (Å, °)

Cell parameters from 25

reflections $\theta = 12 - 21^{\circ}$

 $\mu = 0.093 \text{ mm}^{-1}$

 $0.4 \times 0.2 \times 0.2$ mm

T = 293 (2) K

Prism

Colourless

 $R_{\rm int} = 0.040$

 $k = 0 \rightarrow 23$

 $l = 0 \rightarrow 13$

 $\theta_{\rm max} = 30.11^{\circ}$

 $h = -12 \rightarrow 11$

3 standard reflections

frequency: 120 min

intensity decay: none

| 02—C1 | 1.189 (2) | C7—O3 | 1.377 (2) |
|----------|-------------|----------|-------------|
| C101 | 1.391 (2) | O3—C8 | 1.384 (2) |
| C1—C8 | 1.440 (2) | C8—C9 | 1.372 (2) |
| 01—C2 | 1.375 (2) | C9N | 1.339 (2) |
| 02—C1—O1 | 116.3 (2) | C7—O3—C8 | 117.52 (13) |
| O2—C1—C8 | 126.8 (2) | C9—C8—O3 | 119.60 (14) |
| 01—C1—C8 | 116.92 (15) | C9-C8-C1 | 117.70 (15) |
| C2-01-C1 | 119.81 (13) | O3-C8-C1 | 122.71 (15) |
| 01—C2—C7 | 121.93 (15) | N-C9-C8 | 132.4 (2) |
| 01—C2—C3 | 118.5 (2) | C9-N-C10 | 120.4 (2) |
| C6—C7—O3 | 117.87 (15) | C9-N-C14 | 125.74 (15) |
| O3—C7—C2 | 121.02 (15) | | |

The positions of nine H atoms were determined from a difference map and the positions of the remaining six H atoms (H4, H6, H11, H11A, H14 and H14A) were computed. The isotropic displacement parameters and coordinates were refined for the first nine H atoms. An overall isotropic U and constrained atomic coordinates (riding model) were refined for the remaining six H atoms.

Data collection: CAD-4 Software (Enraf-Nonius, 1989). Cell refinement: CAD-4 Software. Data reduction: CFEO (Solans, 1978). Program(s) used to solve structure: SHELXS86 (Sheldrick, 1990). Program(s) used to refine structure: SHELXL93 (Sheldrick, 1993). Molecular graphics: ORTEP (Brueggemann & Schmid, 1990). Software used to prepare material for publication: CIFTAB (Sheldrick, 1994).

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Ismine[†]

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Abstract

The title compound, $C_{15}H_{15}NO_3$, is an alkaloid isolated from several Narcissus species and used to treat a variety of human medical problems. The structure has two molecules in the asymmetric unit which play different roles in the hydrogen-bonding scheme. This produces differences in the geometry of each molecule.

Comment

Ismine, (I), is an alkaloid isolated from whole plants of several Narcissus species (Viladomat *et al.*, 1990, 1992; Codina *et al.*, 1990) and from *Boophane flava* (Viladomat *et al.*, 1995). The Narcissus are Amaryllidacea species endemic to the north of the Iberian Peninsula. Plants of this genus have been used throughout history to treat a variety of human medical problems (Bastida, Viladomat & Codina, 1997).



The spectroscopic data of ismine are very close to those of 5,6-dihydrobicolorine, also isolated previously from several Narcissus species. In order to confirm its solid-state molecular structure, an X-ray crystallographic study was undertaken.

The X-ray study shows that only the assignment of the aromatic protons H7 and H10 should be interchanged in the ¹H NMR taking into account the HMBC (heteronuclear multiple bond correlation) and HMQC (heteronuclear multiple quantum correlation) correlations (Viladomat *et al.*, 1997).



Fig. 1. The molecular structure of (I) showing the two independent molecules with 50% probability displacement ellipsoids.

[†] Systematic name: {6-[2-(methylamino)phenyl]-2H-1,3-benzodioxol-5-yl}methanol.