

Table 4. Selected geometric parameters (\AA , $^\circ$) for (5) at 193 K

C1—O1	1.443 (4)	C7—C8	1.548 (7)
C1—C2	1.475 (5)	C8—C9	1.475 (5)
C1—C14	1.527 (5)	C9—C10	1.189 (5)
C1—C13	1.536 (5)	C10—C11	1.486 (5)
C2—C3	1.187 (5)	C11—O11	1.450 (4)
C3—C4	1.483 (5)	C11—C15	1.522 (5)
C4—C5	1.495 (6)	C11—C12	1.531 (5)
C5—C6	1.500 (6)	C12—C13	1.533 (5)
C6—C7	1.528 (7)		
C2—C1—C14	109.2 (3)	C9—C8—C7	110.8 (4)
C2—C1—C13	110.2 (3)	C10—C9—C8	175.2 (4)
C14—C1—C13	108.4 (3)	C9—C10—C11	175.9 (4)
C3—C2—C1	173.3 (4)	C10—C11—C15	112.3 (3)
C2—C3—C4	175.3 (4)	C10—C11—C12	109.1 (3)
C3—C4—C5	111.7 (3)	C15—C11—C12	112.1 (3)
C4—C5—C6	118.0 (4)	C11—C12—C13	111.5 (3)
C5—C6—C7	113.6 (4)	C12—C13—C1	117.1 (3)
C6—C7—C8	112.0 (4)		
C1—C2—C3—C4	4 (9)	C11—C12—C13—C1	—154.6 (3)
C8—C9—C10—C11	0 (10)		

Space group $I2/a$ was used, which is the third choice for space group $C2/c$ (No. 15; *International Tables for Crystallography*, 1983, Vol. A), in order to avoid large correlation coefficients between the $sc + z$ coordinates when β is much larger than 90° . At 193 K, there are fewer measurable reflections (*ca* 2%) as N is proportional to the volume of the unit cell, however, the number of observed reflections increased by *ca* 8%.

For both measurements, data collection: *STADI-4* (Stoe & Cie, 1996a); cell refinement: *STADI-4*; data reduction: *X-RED* (Stoe & Cie, 1996b); program(s) used to solve structures: *SHELXS86* (Sheldrick, 1990); program(s) used to refine structures: *SHELXL93* (Sheldrick, 1993); molecular graphics: *PLATON* (Spek, 1990); software used to prepare material for publication: *SHELXL93*.

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Lists of structure factors, anisotropic displacement parameters, H-atom coordinates and complete geometry have been deposited with the IUCr (Reference: NA1269). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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2-Amino-5-methyl-10b*H*-1,3,4-thiadiazolo[3,2-*c*]quinazolin-6-ium-10b-carboxylate Hydrate

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Abstract

The $C_{11}H_{10}N_4O_2S$ molecule in the title compound, $C_{11}H_{10}N_4O_2S \cdot H_2O$, is a new heterocycle. It comprises three fused rings and exists in a zwitterionic form. The two six-membered rings are approximately coplanar, but the presence of a saturated C_{sp^3} atom at the junction of the five-membered and central six-membered rings imposes a marked non-coplanarity on these, with a dihedral angle of $23.1(2)^\circ$. The molecule is thus chiral and the crystal structure is a racemate incorporating two molecules of water for each *R/S* pair of molecules. There is an extensive hydrogen-bonding network.

Comment

Many heterocyclic compounds exhibit bioactivity (Grayson, 1982), especially as antifungal, antitumour and antiviral agents (Iradyan *et al.*, 1990). Several heterocyclic compounds with a composition similar to that of the tricyclic CAM molecule in the title compound, $CAM \cdot H_2O$, are essentially planar and hence achiral. We have determined the structure of $CAM \cdot H_2O$ by X-ray diffraction methods (Gilmore, 1983) as part

2082 reflections
208 parameters
H atoms refined with fixed
 U_{iso}

Extinction correction: none
Scattering factors from *International Tables for X-ray Crystallography* (Vol. IV)

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Table 1. Selected geometric parameters (\AA , $^{\circ}$)

S—C(11)	1.768 (2)	N(2)—N(3)	1.398 (2)
S—C(2)	1.830 (2)	N(2)—C(2)	1.469 (2)
O(1)—C(1)	1.233 (2)	N(3)—C(11)	1.308 (3)
O(2)—C(1)	1.255 (2)	N(4)—C(11)	1.327 (3)
N(1)—C(9)	1.325 (3)	C(1)—C(2)	1.571 (3)
N(1)—C(8)	1.404 (3)	C(2)—C(3)	1.498 (3)
N(2)—C(9)	1.318 (2)	C(3)—C(4)	1.391 (3)
C(11)—S—C(2)	87.2 (1)	N(2)—C(2)—S	102.0 (1)
C(9)—N(1)—C(8)	122.5 (2)	C(3)—C(2)—C(1)	109.4 (1)
C(9)—N(2)—N(3)	121.4 (2)	C(3)—C(2)—S	115.8 (1)
C(9)—N(2)—C(2)	123.2 (2)	C(1)—C(2)—S	108.8 (1)
N(3)—N(2)—C(2)	115.1 (1)	C(8)—C(3)—C(2)	117.5 (2)
C(11)—N(3)—N(2)	108.2 (2)	C(3)—C(8)—N(1)	119.0 (2)
O(1)—C(1)—O(2)	128.7 (2)	N(2)—C(9)—N(1)	119.3 (2)
O(1)—C(1)—C(2)	116.8 (2)	N(3)—C(11)—N(4)	123.2 (2)
O(2)—C(1)—C(2)	114.3 (2)	N(3)—C(11)—S	117.5 (1)
N(2)—C(2)—C(3)	110.2 (1)	N(4)—C(11)—S	119.3 (2)
N(2)—C(2)—C(1)	110.3 (1)		

Table 2. Hydrogen-bonding geometry (\AA , $^{\circ}$)

$D\cdots H\cdots A$	$D\cdots A$	$D\cdots H\cdots A$
N(1)—H(3) \cdots O(3)	2.738 (3)	175
N(4)—H(1) \cdots O(2 ⁱⁱ)	2.782 (3)	173
N(4)—H(2) \cdots N(3 ⁱⁱ)	3.119 (3)	178
O(3)—H(11) \cdots O(1 ⁱⁱⁱ)	2.775 (3)	175
O(3)—H(12) \cdots O(2 ^{iv})	2.832 (2)	174

Symmetry codes: (i) $-x, 1 - y, -z$; (ii) $1 - x, 2 - y, -z$; (iii) $1 + x, y, z$; (iv) $1 - x, 1 - y, 1 - z$.

Data collection: *CAD-4 Software* (Enraf-Nonius, 1989). Cell refinement: *CAD-4 Software*. Data reduction: *TEXSAN PROCESS* and *HKL* (Molecular Structure Corporation, 1989). Program(s) used to refine structure: *TEXSAN LS* and *FOURIER*. Molecular graphics: *ORTEPII* (Johnson, 1976) and *PLUTO* (Motherwell & Clegg, 1978).

Lists of atomic coordinates, displacement parameters, structure factors and complete geometry have been deposited with the IUCr (Reference: CF1099). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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1-Acetyl-4-[(1*R*,2*S*,5*R*)-3-methoxy-carbonyl]piperazine-2,5-dione†

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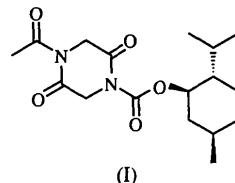
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Abstract

The title compound, $C_{17}H_{26}N_2O_5$, assumes a skewed-boat conformation. The degree of folding of the piperazine-dione ring given by the Hooker parameter, β , was found to be 32.06° .

Comment

X-ray crystallographic studies have shown that the presence of N and/or α -C substituents play an important role in determining the conformations of piperazine-2,5-diones (Karle, 1981). As part of our studies (Chai & King, 1995) to exploit such conformational effects to direct reactions stereoselectively, the title compound, (I), was synthesized.



The crystal structure shows that the methoxy-carbonyl substituent extends away from the piperazine-dione ring, whereas the latter adopts a skewed boat conformation. This is in contrast to the conformations adopted by glycine anhydride and its derivatives. For

† Alternative name: (1'*R*,2'*S*,5'*R*)-3-methyl 4-acetyl-2,5-dioxopiperazine-1-carboxylate.