Lists of structure factors, anisotropic displacement parameters, Hatom coordinates, complete geometry and torsion angles have been deposited with the IUCr (Reference: MU1138). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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# 2,2-Dimethyl-5-(2-methyl-4-phenyl-3,4diaza-2-butenylidene)-1,3-dioxane-4,6-dione

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#### Abstract

The molecular conformation of the title compound,  $C_{15}H_{16}N_2O_4$ , is described in terms of three planar sections which are mutually twisted. The main intermolecular contacts are N—H···O hydrogen bonds where the O···N separation is 3.063 (3) Å.

#### Comment

The study of the structure of the title compound, (1), was undertaken to determine whether there were conformational reasons for its ready cyclization to give a 3-oxopyridazinecarboxylic acid upon thermolysis in solution, while the corresponding unsubstituted compound (2) is recovered unchanged under similar conditions (McNab & Stobie, 1982). The structure of (1) is also related to that of the propenylidene Meldrum's acid derivative (3), which we have described recently (Blake, McNab & Monahan, 1991).



The conjugated system from C5 to N10 in (1) is exclusively *trans*, as found for (3), and so the cyclization reaction cannot be explained by a ground-state proximity of N10 and C4.

A feature of the structure of (3) is that electron delocalization occurs along the propenyl chain such that the lengths of all the C—C and C=C bonds are comparable [1.385 (8)–1.409 (8) Å]. In the case of (1), the system is much more localized [C=C 1.355 (3), C— C 1.439 (3) Å], partly due to poorer electron donation from the hydrazone N atom N10 compared with the terminal amino group in (3), and also partly due to the non-planarity of the system (see below). This reduced electron donation is also reflected in the lengths of the C4—C5 and C5—C6 bonds [both 1.469 (3) Å], which are substantially longer than the corresponding values for (3) [1.449 (8), 1.417 (8) Å].

The presence of the C8M methyl substituent in (1) has a dramatic effect on the angles subtended at C8 and C5. Non-bonded contacts between the C8M methyl group and O4 lead to widening of the C7-C8-C8M and C8M—C8—N9 angles to 125.7(2) and  $123.7(2)^{\circ}$ . respectively, with a concomitant reduction in the C7-C8—N9 angle to  $110.5 (2)^{\circ}$ . The C4—C5—C7 angle is widened to 126.0(2)°, compared with an angle of  $120.9(5)^{\circ}$  in (3). This distortion is almost entirely at the expense of the endocyclic C4-C5-C6 angle  $[116.5 (2)^{\circ}, \text{ compared with } 121.7 (5)^{\circ} \text{ in } (3)].$  A similar trend at C5 has been noted for the corresponding angles in the dimethylaminomethylene Meldrum's acid, which has analogous steric constraints (Blake, McNab & Monahan, 1991). In (1), the two exocyclic N-C-C angles involving the phenyl group differ significantly, possibly because of repulsion between the ortho H6P atom and the lone pair on N9.

The molecular conformation of (1) can be described in terms of three planar sections: the phenyl ring, the five-atom chain C5—C7—C8—N9—N10 with its C8M substituent, and four atoms (O1, O3, C4 and C6) of the Meldrum's acid ring. The angle between the normals to the phenyl ring and the chain is  $10.2 (2)^{\circ}$  and the angle between the normals to the chain and the plane of the Meldrum's acid ring is  $20.5 (2)^{\circ}$ . In contrast, there is a much greater degree of coplanarity between the chain and the Meldrum's acid ring in (3). The mean deviation from the plane of the chain is 0.058 Å, while the four atoms defining the Meldrum's acid plane 01 C2 C2A

C2B O3

C4 04

C5

C6 06 C7 **C**8 C8M

N9 N10

C1PC2P

C3P

C4P

C5P C6P

are essentially coplanar (mean deviation 0.0013 Å). Within the Meldrum's acid ring, C5 is displaced in one direction from the plane by 0.276(3)Å, while O4 and O6 are displaced in the opposite direction by 0.330(3) and 0.244(3)Å, respectively; the greater deviation of O4 may be explained by the proximity of C8M. These distortions may be sufficient to lower the activation energy for bond rotation relative to that in the unsubstituted compound (2) so that thermal cyclization may take place.

The principal intermolecular contacts are hydrogen bonds between N10—H10 and O6 at (x, 1+y, z), which link the molecules to form chains parallel to the b axis.



Fig. 1. A view of the molecule with the atom-numbering scheme. Displacement ellipsoids enclose 50% probability surfaces.

## **Experimental**

The title compound was obtained by the reaction of PhNHN=C(Me)CHO with 2,2-dimethyl-1,3-dioxane-4,6dione. Crystals were grown from ethanol (McNab & Stobie, 1982).

#### Crystal data

$C_{15}H_{16}N_2O_4$ $M_2 = 288.30$	Mo $K\alpha$ radiation	Table 2. Selected geometric parameters (Å, °)			
Triclinic $P\overline{1}$ a = 7.544 (3) Å b = 8.576 (3) Å	$\lambda = 0.71073 \text{ A}$ Cell parameters from 12 reflections $\theta = 15-16^{\circ}$ $\mu = 0.098 \text{ mm}^{-1}$	01—C6 01—C2 C2—03 C2—C2A C2—C2B	1.349 (3) 1.435 (3) 1.430 (2) 1.499 (3) 1.508 (3)	C5—C7 C5—C6 C6—O6 C7—C8 C8—N9	1.355 (3) 1.469 (3) 1.208 (3) 1.439 (3) 1.309 (3)
c = 12.014 (7)  Å $\alpha = 78.01 (3)^{\circ}$ $\alpha = 78.226 (2)^{\circ}$	$\mu = 0.098$ mm T = 295 (2) K Lath	O3—C4 C4—O4 C4—C5	1.354 (2) 1.206 (3) 1.469 (3)	C8—C8M N9—N10 N10—C1P	1.490 (3) 1.330 (2) 1.400 (2)
$p = 83.30 (3)^{\circ}$ $\gamma = 71.41 (3)^{\circ}$ $V = 719.6 (6) Å^{3}$ Z = 2 $D_{x} = 1.330 \text{ Mg m}^{-3}$ Data collection Stoe Stadi-4 diffractometer $\omega - 2\theta$ scans Absorption correction: none	0.75 × 0.33 × 0.075 mm Orange $R_{int} = 0.0116$ $\theta_{max} = 22.52^{\circ}$ $h = -7 \rightarrow 8$ $k = -8 \rightarrow 0$	$\begin{array}{c} C6 - 01 - C2 \\ O3 - C2 - 01 \\ O3 - C2 - C2A \\ O1 - C2 - C2A \\ O3 - C2 - C2B \\ O1 - C2 - C2B \\ C4 - C2 - C2B \\ C4 - O3 - C2 \\ O4 - C4 - O3 \\ O4 - C4 - O3 \\ O4 - C4 - C5 \\ O3 - C4 - C5 \\ C7 - C5 - C6 \\ \end{array}$	118.2 (2) 109.1 (2) 107.0 (2) 105.9 (2) 110.4 (2) 113.5 (2) 118.4 (2) 118.2 (2) 125.6 (2) 115.9 (2) 126.0 (2) 117.1 (2)	C4-C5-C6 O6-C6-O1 O6-C6-C5 O1-C6-C5 C5-C7-C8 N9-C8-C7 N9-C8-C8 M C7-C8-C8M C7-C8-C8M C7-C8-C8M C8-N9-N10 N9-N10-C1P C6P-C1P-N10 C2P-C1P-N10	116.5 (2) 118.2 (2) 125.5 (2) 116.2 (2) 132.9 (2) 110.5 (2) 123.7 (2) 125.7 (2) 119.2 (2) 119.06 (15) 121.38 (10) 118.60 (10)
1858 measured reflections1847 independent reflections1499 observed reflections $[I > 2\sigma(I)]$	$l = 0 \rightarrow 12$ 3 standard reflections frequency: 120 min intensity decay: 2.0%	C4C5C7C8 C6C5C7C8 C5C7C8N9 C5C7C8C8 <i>M</i> C7C8N9N10	16.6 (4) -170.8 (2) -169.2 (2) 14.2 (4) 177.4 (2)	C8 <i>M</i> —C8—N9—N10 C8—N9—N10—C1 <i>P</i> N9—N10—C1 <i>P</i> —C6 <i>P</i> N9—N10—C1 <i>P</i> —C2 <i>P</i>	-5.9 (3) 179.0 (2) -8.5 (2) 171.33 (13)

Refinement

Refinement on $F^2$	$\Delta \rho_{\rm max} = 0.216 \ {\rm e} \ {\rm \AA}^{-3}$
$R[F^2 > 2\sigma(F^2)] = 0.0397$	$\Delta \rho_{\rm min} = -0.200 \ {\rm e} \ {\rm \AA}^{-3}$
$wR(F^2) = 0.1394$	Extinction correction:
S = 1.124	SHELXL93 (Sheldrick,
1845 reflections	1993)
185 parameters	Extinction coefficient:
H10 was constrained to lie	0.046 (9)
1.00 Å from N10; other	Atomic scattering factors
H atoms were placed in	from International Tables
calculated positions.	for Crystallography (1992
$w = 1/[\sigma^2(F_o^2) + (0.1P)^2]$	Vol. C, Tables 4.2.6.8 and
where $P = (F_o^2 + 2F_c^2)/3$	6.1.1.4)
$(\Delta/\sigma)_{\rm max} = 0.367$	

## Table 1. Fractional atomic coordinates and equivalent isotropic displacement parameters (Å<sup>2</sup>)

$$U_{\text{eq}} = (1/3) \sum_i \sum_j U_{ij} a_i^* a_i^* \mathbf{a}_i \cdot \mathbf{a}_j.$$

x	y	Z	Uea
0.2374 (2)	0.2956 (2)	0.82567 (12)	0.0474 (5)
0.2004 (3)	0.4137 (2)	0.7209 (2)	0.0408 (6)
0.2316 (4)	0.3119 (3)	0.6290 (2)	0.0569 (7)
0.0055 (3)	0.5346 (3)	0.7267 (2)	0.0554 (7)
0.3367 (2)	0.5016 (2)	0.69950 (11)	0.0439 (4)
0.3673 (3)	0.5714 (2)	0.7839 (2)	0.0400 (6)
0.4654 (2)	0.6617 (2)	0.76169 (13)	0.0602 (5)
0.2926 (3)	0.5147 (2)	0.8982 (2)	0.0355 (5)
0.2617 (3)	0.3502 (3)	0.9186 (2)	0.0412 (6)
0.2589 (3)	0.2608 (2)	1.01041 (13)	0.0615 (5)
0.2684 (3)	0.5915 (2)	0.9891 (2)	0.0384 (5)
0.2622 (3)	0.7571 (2)	1.0000 (2)	0.0364 (5)
0.2414 (3)	0.9034 (3)	0.9051 (2)	0.0491 (6)
0.2675 (2)	0.7654 (2)	1.10718 (14)	0.0381 (5)
0.2704 (2)	0.9075 (2)	1.13387 (13)	0.0387 (5)
0.2730 (2)	0.9166 (2)	1.24865 (8)	0.0354 (5)
0.3019 (2)	1.05723 (14)	1.27502 (9)	0.0419 (6)
0.3058 (2)	1.06988 (15)	1.38855 (11)	0.0490 (6)
0.2809 (2)	0.9418 (2)	1.47570 (8)	0.0561 (7)
0.2520 (3)	0.8012 (2)	1.44930 (9)	0.0610 (7)
0.2481 (2)	0.78853 (15)	1.33577 (11)	0.0526 (6)

, °)

The phenyl ring (C1P to C6P) was constrained as a regular hexagon with C—C distances of 1.390 Å. Data collection and cell refinement: DIF4 (Stoe & Cie, 1988a). Data reduction: *REDU4* (Stoe & Cie, 1988b). Program(s) used to solve structure: *SHELXS86* (Sheldrick, 1990). Program(s) used to refine structure: *SHELXL93* (Sheldrick, 1993). Molecular graphics: *SHELXTL/PC* (Sheldrick, 1992).

We thank the SERC for funding towards the provision of a four-circle diffractometer.

Lists of structure factors, anisotropic displacement parameters, Hatom coordinates, bond distances and angles involving non-H atoms and torsion angles have been deposited with the IUCr (Reference: HA1123). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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## (2RS)-4,4-Dimethyl-2-[(1SR)-1-phenylethyl]-1-pyrrolidinol

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#### Abstract

In order to determine the stereospecificity of the thermal cyclization of [(E)-2,2-dimethyl-5-phenyl-4-hexenyl]hydroxylamine (1), the relative configuration of the title compound, C<sub>14</sub>H<sub>21</sub>NO, (2), has been established by single-crystal X-ray diffraction. The pyrrolidine ring adopts an envelope conformation with the N atom out

of the ring plane. The molecules form hydrogen-bonded pairs through a centre of inversion  $[O \cdots N \ 2.808 \ (5) \text{ Å} and O - H \cdots N \ 160 \ (4)^{\circ}]$ .

### Comment

The thermal cyclization of *N*-alkenylhydroxylamines, first reported by House and co-workers (House, Manning, Melillo, Lee, Haynes & Wilkes, 1976; House & Lee, 1976) and independently discovered by us (Oppolzer, Siles, Snowden, Bakker & Petrzilka, 1979), was initially proposed to occur *via* a radical chain mechanism. As an alternative to the thermal conversion of *N*-alkenyl-*N*-methylhydroxylamines to cyclic *N*-oxides, a retro-Cope elimination mechanism has also been postulated (Ciganek, 1990). However, compelling proof of either mechanism has not yet been presented.



In order to study the alkene faciality of this process, the (*E*)-5,5-disubstituted 4-alkenylhydroxylamine (1) was cyclized by heating it in degassed benzene under reflux (18 h) to provide the *N*-hydroxypyrrolidine (2) (m.p. 358–359 K) in 81% yield. Under analogous reaction conditions, the Z isomer of (1) gave an epimer of (2) (oil, yield 81%) without cross contamination (Oppolzer, Spivey & Bochet, 1994).



Fig. 1. View of the *N*-hydroxypyrrolidine (2) with the atomic labelling of non-H atoms shown. Ellipsoids are shown at the 30% probability level.