Received 22 March 2004

Accepted 23 March 2004

Online 27 March 2004

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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#### Key indicators

Single-crystal X-ray study T = 150 KMean  $\sigma(\text{C-C}) = 0.002 \text{ Å}$ Disorder in main residue R factor = 0.047 wR factor = 0.141 Data-to-parameter ratio = 26.4

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

# 2-(Adamantan-1-yl)-5-methylbenzo[*d*]-[1,3]oxazin-4-one

At 150 K, the benzo[d][1,3]oxazin-4-one heterocycle in the title compound, C<sub>19</sub>H<sub>21</sub>NO<sub>2</sub>, lies on a crystallographic mirror plane. This group is planar despite the resulting unfavourable steric interaction between the proximal 5-methyl and 4-carbonyl groups.

### Comment

The title compound, (I), has a planar benzo[d][1,3]oxazin-4one heterocycle that lies on a crystallographic mirror plane. The molecular structure of (I) is shown in Fig. 1 and selected bond distances and angles are given in Table 1. Within the oxazin-4-one group the C=O and C=N double bonds are clearly localized, but of the two formally single C-O bonds, O2-C2 is significantly shorter than O2-C1. The bicyclic heterocycle is planar despite the unfavourable steric interaction between the 5-methyl and 4-carbonyl groups, but the planarity allows  $\pi$ -stacking of these groups in the direction of the *b* axis (Fig. 2), with an inter-layer distance of 3.3662 (4) Å (Table 2). The widened bond angles of 128.39 (12), 121.75 (11) and 123.51 (12)° for O1-C1-C4, C1-C4-C5 and C4-C5-C9, respectively, still result in a short  $O1\cdots$ C9 distance of 2.838 (2) Å.



### Experimental

Suitable crystals of the title compound, (I), were prepared by the attempted recrystallization of 2-[(adamantane-1-carbonyl)-amino]-6-methylbenzoic acid from petroleum ether (80/100)-toluene.



### Figure 1

View of the molecular structure of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level and H atoms are represented by spheres of arbitrary size. Only one orientation of the disordered methyl group is shown.

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# organic papers

#### Crystal data

 $C_{19}H_{21}NO_2$   $M_r = 295.37$ Monoclinic,  $P2_1/m$  a = 8.3514 (12) Å b = 6.7324 (5) Å c = 13.3203 (19) Å  $\beta = 104.275$  (11)° V = 725.81 (16) Å<sup>3</sup> Z = 2

#### Data collection

Stoe IPDSII area-detector diffractometer  $\varphi$  and  $\omega$  scans Absorption correction: none 10561 measured reflections 3325 independent reflections

#### Refinement

Refinement on $F^2$ $R[F^2 > 2\sigma(F^2)] = 0.048$	$w = 1/[\sigma^2(F_o^2) + (0.081P)^2]$ where $P = (F^2 + 2F^2)/3$
$wR(F^2) = 0.141$	$(\Delta/\sigma)_{\rm max} < 0.001$
S = 1.00	$\Delta \rho_{\text{max}} = 0.49 \text{ e A}^{-3}$
126 parameters	Extinction correction: SHELXL97
H-atom parameters constrained	Extinction coefficient: 0.032 (7)

 $D_x = 1.352 \text{ Mg m}^{-3}$ 

Cell parameters from 8117

2132 reflections with  $I > 2\sigma(I)$ 

Mo  $K\alpha$  radiation

reflections

 $\theta = 2.5 - 34.7^{\circ}$  $\mu = 0.09 \text{ mm}^{-1}$ 

T = 150 (2) K

 $R_{\rm int} = 0.061$ 

 $\theta_{\rm max} = 34.7^\circ$ 

 $h = -13 \rightarrow 13$ 

 $k = -10 \rightarrow 9$ 

 $l = -21 \rightarrow 17$ 

Block, colourless  $0.60 \times 0.25 \times 0.20$  mm

### Table 1

Selected geometric parameters (Å, °).

O1-C1	1.2010 (16)	N1-C3	1.4019 (16)
O2-C2	1.3750 (15)	C1-C4	1.4625 (17)
O2-C1	1.3940 (15)	C3-C4	1.4082 (18)
N1-C2	1.2762 (15)		
C2-O2-C1	122.07 (9)	N1-C2-C10	123.61 (11)
C2-N1-C3	117.81 (11)	O2-C2-C10	111.75 (9)
O1-C1-O2	116.33 (11)	N1-C3-C4	122.43 (10)
O1-C1-C4	128.39 (12)	C3-C4-C1	117.77 (11)
O2-C1-C4	115.28 (10)	C5-C4-C1	121.75 (11)
N1-C2-O2	124.64 (11)	C4-C5-C9	123.51 (12)

#### Table 2

Contact distances (Å).

$\overline{C1 \cdots C7^{i}}$	3.3984 (4)	$C4 \cdot \cdot \cdot C8^i$	3.4884 (5)
$C3 \cdot \cdot \cdot C5^i$	3.4038 (4)	$N1 \cdots C6^i$	3.5083 (5

Symmetry code: (i) -x, -y, 1 - z.

### Figure 2

Packing diagram for (I), viewed down the b axis. Only one orientation for the disordered methyl group is shown.

All H atoms were initially located in a difference Fourier map. The methyl H atoms were constrained to an ideal geometry with a C–H distance of 0.98 Å, but the group was allowed to rotate freely about its X-C bond. In its final position, the methyl group is not bisected exactly by the mirror plane and hence is disordered 50:50 about the mirror plane. All other H atoms were placed in geometrically idealized positions, with C–H distances of 0.95–1.00 Å.  $U_{\rm iso}({\rm H})$  values were set at  $1.2U_{\rm eq}({\rm C})$  for all of the H atoms.

Data collection: X-AREA (Stoe & Cie, 2001); cell refinement: X-AREA; data reduction: X-RED32 (Stoe & Cie, 2001); program(s) used to solve structure: X-STEP32 (Stoe & Cie, 2001) and WinGX (Farrugia, 1999); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP-3 for Windows (Farrugia, 1997); software used to prepare material for publication: WinGX.

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