

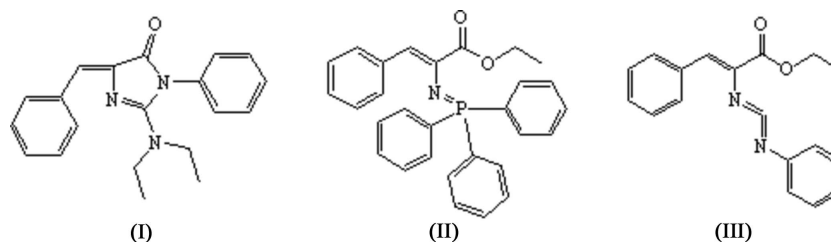
## Gui-Hua Li and Yang-Gen Hu\*

Department of Medicinal Chemistry, Yongyang  
Medical College, Shiyan 442000, People's  
Republic of ChinaCorrespondence e-mail:  
huyangg111@yahoo.com.cn

## Key indicators

Single-crystal X-ray study  
 $T = 292$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.003$  Å  
Disorder in main residue  
 $R$  factor = 0.056  
 $wR$  factor = 0.169  
Data-to-parameter ratio = 16.8For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.4-Benzylidene-2-diethylamino-1-phenyl-  
1*H*-imidazol-5(4*H*)-oneIn the crystal structure of the title compound,  $\text{C}_{20}\text{H}_{21}\text{N}_3\text{O}$ ,  
intramolecular  $\text{C}-\text{H}\cdots\text{N}$  hydrogen bonds stabilize the  
conformation of the molecule.Received 25 March 2006  
Accepted 29 March 2006

## Comment

Derivatives of 4*H*-imidazol-4-ones have shown biological  
activity (Lacroix *et al.*, 2000). Some 2-alkylaminoimidazolones  
exhibit good antibacterial and antifungal activities (Trivedi *et al.*,  
2002). Recently, in our work on the synthesis of biologi-  
cally active imidazolinones, we have developed a facile  
synthesis of 2-dialkylamino-4*H*-imidazolin-4-ones (Hu *et al.*,  
2004; Xin & Hu, 2006). The title compound, (I), may be used  
as a new precursor for obtaining bioactive molecules and its  
structure is reported here (Fig. 1 and Table 1).The five-membered imidazolone ring is planar, with a  
maximum deviation of 0.012 (2) Å for atoms N2. The C1–C6  
phenyl ring is only slightly twisted with respect to the imida-  
zolone ring [dihedral angle = 72.1 (1)°]. The diethylamino  
substituent (C18/C17/N3/C19/C20) is disordered over two  
sites, with refined occupancies of 0.829 (4) and 0.171 (4)  
(Fig. 1). Two intramolecular  $\text{C}-\text{H}\cdots\text{N}$  hydrogen-bonding  
interactions are present which stabilize the conformation of  
the molecule (Table 2). There are no hydrogen-bonding or  $\pi$ -  
 $\pi$  interactions (Fig. 2).

## Experimental

To a solution of *N*-(1-ethoxycarbonyl-2-phenylethen-1-yl)imino-  
triphenylphosphorane, (II) (3 mmol), in dry dichloromethane (15 ml)  
was added phenyl isocyanate (3 mmol) under nitrogen at room  
temperature. After being left to stand for 8 h, the solvent was  
removed under reduced pressure, and diethyl ether/petroleum ether  
(1:2, 20 ml) was added to precipitate triphenylphosphine oxide. After  
filtration, the solvent was removed to give the carbodiimide (III),  
which was used directly without further purification. Diethylamine  
(3 mmol) was added to a solution of the carbodiimide in acetonitrile  
(15 ml). The mixture was stirred for 6 h, concentrated under reduced  
pressure and the residue recrystallized from dichloromethane/  
petroleum ether (1:4) to give the title compound, (I) (yield 92%; m.p.

397 K). Suitable crystals were obtained by vapour diffusion of ethanol into dichloromethane at room temperature.

Crystal data

C<sub>20</sub>H<sub>21</sub>N<sub>3</sub>O  
*M<sub>r</sub>* = 319.40  
 Orthorhombic, *Pbca*  
*a* = 18.8275 (10) Å  
*b* = 8.9784 (5) Å  
*c* = 20.9951 (11) Å  
*V* = 3549.0 (3) Å<sup>3</sup>

*Z* = 8  
*D<sub>x</sub>* = 1.196 Mg m<sup>-3</sup>  
 Mo *K*α radiation  
 μ = 0.08 mm<sup>-1</sup>  
*T* = 292 (2) K  
 Block, yellow  
 0.30 × 0.30 × 0.20 mm

Data collection

Bruker SMART 4K CCD area-detector diffractometer  
 φ and ω scans  
 Absorption correction: multi-scan (*SADABS*; Sheldrick, 2003)  
*T<sub>min</sub>* = 0.978, *T<sub>max</sub>* = 0.985

28769 measured reflections  
 4072 independent reflections  
 2667 reflections with *I* > 2σ(*I*)  
*R<sub>int</sub>* = 0.032  
 θ<sub>max</sub> = 27.5°

Refinement

Refinement on *F*<sup>2</sup>  
*R* [*F*<sup>2</sup> > 2σ(*F*<sup>2</sup>)] = 0.056  
*wR* (*F*<sup>2</sup>) = 0.169  
*S* = 1.08  
 4072 reflections  
 243 parameters  
 H-atom parameters constrained

*w* = 1/[σ<sup>2</sup>(*F<sub>o</sub>*<sup>2</sup>) + (0.0794*P*)<sup>2</sup> + 0.4215*P*]  
 where *P* = (*F<sub>o</sub>*<sup>2</sup> + 2*F<sub>c</sub>*<sup>2</sup>)/3  
 (Δ/σ)<sub>max</sub> < 0.001  
 Δρ<sub>max</sub> = 0.28 e Å<sup>-3</sup>  
 Δρ<sub>min</sub> = -0.21 e Å<sup>-3</sup>

Table 1

Selected geometric parameters (Å, °).

C7—O1	1.209 (2)	C16—N3'	1.456 (7)
C7—N1	1.388 (2)	N3—C19	1.474 (3)
C8—N2	1.385 (2)	N3'—C19'	1.521 (8)
C16—N3	1.346 (3)	N3'—C17'	1.525 (9)
C5—C6—N1	119.61 (17)	N1—C16—N3'	114.5 (5)
O1—C7—N1	125.29 (16)	C16—N3—C17	116.59 (17)
N1—C7—C8	104.07 (14)	C16—N3—C19	123.36 (18)
N2—C8—C7	109.03 (14)	C16—N3'—C19'	123.5 (8)
N2—C16—N3	123.82 (16)	C20'—C19'—N3'	119.1 (17)
N3—C16—N1	121.76 (16)	C7—N1—C6	121.44 (14)
N3—C16—N3'	35.2 (5)	C16—N2—C8	106.04 (14)
C4—C5—C6—N1	-179.7 (2)	N2—C16—N3'—C19'	163.9 (9)
C2—C1—C6—N1	179.28 (19)	N1—C16—N3'—C19'	-54.2 (13)
O1—C7—C8—C9	3.0 (3)	N3—C16—N3'—C17'	-93.7 (16)
N1—C7—C8—C9	-178.30 (17)	C16—N3'—C19'—C20'	-41 (2)
N1—C7—C8—N2	1.30 (19)	C17'—N3'—C19'—C20'	114.5 (19)
N3'—C16—N3—C17	84.7 (7)	O1—C7—N1—C6	19.0 (3)
N1—C16—N3—C19	-31.7 (3)	N3'—C16—N1—C7	-145.4 (6)
N3'—C16—N3—C19	-119.8 (7)	C9—C8—N2—C16	177.51 (19)
C16—N3—C19—C20	127.5 (3)		

Table 2

Hydrogen-bond geometry (Å, °).

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
C15—H15...N2	0.93	2.44	3.078 (2)	126
C19—H19A...N1	0.97	2.56	2.968 (3)	105

Positional disorder was found in atoms C17—C20/N3 of the diethylamino substituent and atoms of the minor disorder component were refined isotropically. The final site-occupancy factors for the two components were 0.829 (4) and 0.171 (4). All H atoms were positioned geometrically [C—H = 0.93 (CH), 0.97 (CH<sub>2</sub>) and 0.96 Å

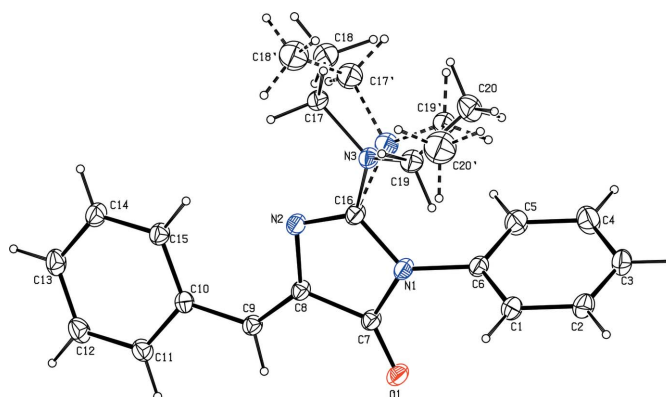


Figure 1

View of the molecule of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level. Both disorder components are shown.

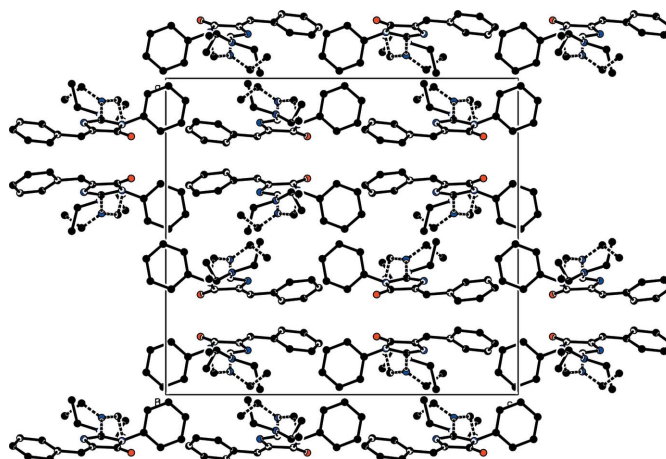


Figure 2

The crystal structure of (I), viewed along the *b* axis. H atoms bonded to C atoms have been omitted for clarity. Only one disorder component is shown.

(CH<sub>3</sub>)] and constrained to ride on their parent atoms, with *U*<sub>iso</sub>(H) values of 1.2 (1.5 for methyl) times *U*<sub>eq</sub>(C).

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT* (Bruker, 2001); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXTL* (Sheldrick, 2001).

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