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### Key indicators

Single-crystal X-ray study T = 294 K Mean  $\sigma$ (C–C) = 0.003 Å R factor = 0.047 wR factor = 0.139 Data-to-parameter ratio = 13.9

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. Ethyl 8-benzoyl-5-oxo-7-phenyl-1,2,3,5,6,7hexahydroimidazo[1,2-a]pyridine-6carboxylate

The geometric parameters for the title compound,  $C_{23}H_{22}N_2O_4$ , are normal. The molecules form dimeric associations via inversion-generated pairs of  $N-H\cdots O$  hydrogen bonds.

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# Comment

5-Oxoimidazo[1,2-*a*]pyridine derivatives are potentially biologically active compounds. Heterocyclic ketene aminals (HKAs) are useful intermediates for the preparation of these compounds. We have reported some examples of HKAs reacting with  $\alpha$ , $\beta$ -unsaturated compounds to yield 5-oxoimidazo[1,2-*a*]pyridine derivatives (Huang & Liu, 1986). Compound (I) (Fig. 1) (Table 1) resulted from a Michael addition reaction (see *Experimental*). A <sup>1</sup>H–<sup>1</sup>H nuclear Overhauser effect experiment failed to provide unambiguous evidence for the determination of a *Z* or *E* configuration about the C1=C16 double bond and its crystal structure was determined to resolve this ambiguity.



Compound (I) is chiral (in the molecule of the arbitrarily chosen asymmetric unit, C5 has an R configuration and C9 has an S configuration), but crystal symmetry generates a racemic mixture of the two enantiomers.

The six-membered C1/N2/C4/C5/C9/C16 ring in (I) adopts a twist boat conformation; the torsion angle N2-C4-C5-C9 is -34.7 (3)° and N2-C1-and C16-C9 is 3.2 (2)°. The H5-C5-C9-H9 dihedral angle of 74° is consistent with the value (64.1°) calculated by application of the Karplus equation on the basis of the NMR coupling constant of 1.9 Hz for these H atoms.

The N1-H1 group participates in a bifurcated N-H···(O,O) interaction: an intramolecular hydrogen bond to O4 (Table 2) and an intermolecular bond to a second, inversion symmetry-generated, O4 species in an adjacent molecule (bond angle sum about H1 =  $350^{\circ}$ ) (Fig. 2). The molecular packing is shown in Fig. 3.

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### Figure 1

A view of (I), showing 40% displacement ellipsoids (arbitrary spheres for the H atoms). All H atoms except H1, H5 and H9 have been omitted for clarity. The dashed line indicates a hydrogen bond.

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# Figure 2

The dimeric association of molecules of (I). Dashed lines indicate hydrogen bonds. [Symmetry code: (i) 1 - x, 1 - y, -z.]

# Experimental

A solution of 2-(benzoylmethylene)imidazolidine (1 mmol) (Huang & Wang, 1992) and diethyl 2-benzylidene malonate (1 mmol) derived from benzaldehyde (Breukelman *et al.*, 1981) in dioxane (10 ml) was stirred at 363 K for 12 h and then refluxed for 36 h. After removal of the solvent, the desired product was separated by flash chromatography (ethyl acetate–petroleum ether, 1.6:1) to yield the title compound, (I), in 71% yield (m.p. 440 K). Crystals of (I) were recrystallised from ethyl acetate–petroleum ether (ca 1:6).

### Crystal data

 $\begin{array}{l} C_{23}H_{22}N_2O_4 \\ M_r = 390.43 \\ \text{Monoclinic, } P_{2_1}^2/c \\ a = 9.644 \ (3) \text{ Å} \\ b = 11.937 \ (3) \text{ Å} \\ c = 17.998 \ (5) \text{ Å} \\ \beta = 90.691 \ (5)^\circ \\ V = 2071.8 \ (10) \text{ Å}^3 \end{array}$ 

# Data collection

Bruker SMART CCD area-detector diffractometer  $\varphi$  and  $\omega$  scans Absorption correction: multi-scan (*SADABS*; Bruker, 1998)  $T_{\min} = 0.823, T_{\max} = 1.00$ 

# Refinement

Refinement on  $F^2$   $R[F^2 > 2\sigma(F^2)] = 0.047$   $wR(F^2) = 0.139$  S = 1.06 3660 reflections 263 parameters H-atom parameters constrained Z = 4  $D_x = 1.252 \text{ Mg m}^{-3}$ Mo K\alpha radiation  $\mu = 0.09 \text{ mm}^{-1}$  T = 294 (2) KBlock, colourless  $0.30 \times 0.24 \times 0.18 \text{ mm}$ 

10276 measured reflections 3660 independent reflections 2457 reflections with  $I > 2\sigma(I)$  $R_{\text{int}} = 0.036$  $\theta_{\text{max}} = 25.0^{\circ}$ 

 $w = 1/[\sigma^{2}(F_{o}^{2}) + (0.0677P)^{2} + 0.4139P]$ where  $P = (F_{o}^{2} + 2F_{c}^{2})/3 (\Delta/\sigma)_{max} = 0.001 \Delta\rho_{max} = 0.23 \text{ e} \text{ Å}^{-3} \Delta\rho_{min} = -0.25 \text{ e} \text{ Å}^{-3}$ 



# Figure 3

The packing in (I), viewed down [100], with hydrogen bonds indicated by dashed lines. All C-bound H atoms have been omitted for clarity.

# Table 1

Selected bond lengths (Å).

C1-N1	1.329 (3)	C1-C16	1.378 (3)
C1-N2	1.390 (3)		

Table 2Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D{\cdots}A$	$D - \mathbf{H} \cdots A$
N1-H1···O4	0.86	2.20	2.720 (3)	119
$N1 - H1 \cdots O4^i$	0.86	2.19	2.854 (3)	134

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

All H atoms were placed in geometrically idealized positions (C– H = 0.95–1.00 Å) and refined as riding, with  $U_{iso}$ (H) = 1.2 $U_{eq}$ (carrier), or  $U_{iso}$ (H) = 1.5 $U_{eq}$ (methyl carrier).

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT* (Bruker, 1998); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics:

SHELXTL (Bruker, 1998); software used to prepare material for publication: SHELXTL.

Dr Hong-Geng Wang of Nankai University collected the intensity data.

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