# organic papers

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# Shazia Anjum,<sup>a</sup>\* M. Iqbal Choudhary,<sup>a</sup> Shamsher Ali,<sup>a</sup> Hoong-Kun Fun<sup>b</sup>\* and Atta-ur-Rahman<sup>a</sup>

<sup>a</sup>HEJ Research Institute of Chemistry, International Centre for Chemical Sciences, University of Karachi, Karachi 75270, Pakistan, and <sup>b</sup>X-ray Crystallography Unit, School of Physics, Universiti Sains Malaysia, 11800 USM, Penang, Malaysia

Correspondence e-mail: anjumshazia@yahoo.com, hkfun@usm.my

#### **Key indicators**

Single-crystal X-ray study T = 293 K Mean  $\sigma$ (C–C) = 0.002 Å R factor = 0.042 wR factor = 0.113 Data-to-parameter ratio = 14.8

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

# 2-(Methoxycarbonyl)succinanilic acid

The title compound,  $C_{12}H_{13}NO_5$ , was synthesized by the condensation of methyl anthranilate and succinic anhydride. The dihedral angle between the phenylacetamide and carboxylic acid (-C-COOH) planes is 80.47 (5)°. In the crystal structure, inversion-related molecules form an O-H···O hydrogen-bonded dimer. Adjacent dimers are interlinked by C-H···O hydrogen bonds to form a chain along [110].

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## Comment

Methyl(2-methoxycarbonyl)succinate, a natural aromatic amide isolated from the methanolic extract of *Jolyna laminarioides*, has shown potent chymotrypsin inhibitory activity (Atta-ur-Rahman *et al.*, 1997). Therefore, we have synthesized our desired analogue, (I), in 85% yield, by a one-step condensation of methyl anthranilate and succinic anhydride (see scheme). We report here the structure of (I).



Molecules of the title compound, (I), have normal bond lengths (Allen *et al.*, 1987). The C6–N1–C7 bond angle



#### Figure 1

The structure of (I), showing 50% probability displacement ellipsoids and the atom-numbering scheme. Dashed lines indicate intramolecular hydrogen bonds.

© 2005 International Union of Crystallography Printed in Great Britain – all rights reserved  $[130.13 (11)^{\circ}]$  is larger than the value of 126.69 (10)° observed in *N*,*N*'-diphenylsuccinimide (Anjum *et al.*, 2005). The N1–C7 and C8–C9 bonds are *trans* with respect to the C7–C8 bond for steric reasons, the N1–C7–C8–C9 torsion angle being 175.77 (11)°.

The phenylacetamide moiety is planar to within  $\pm 0.018$  (1) Å. The C9-C10-O4-O5 and C11-C12-O1-O2 planes form dihedral angles of 80.47 (5) and 9.38 (9)°, respectively, with the phenylacetamide moiety. This orientation is influenced by intramolecular C-H···O and N-H···O interactions, namely N1-H1N···O2, C5-H5···O3 and C2-H2···O1 (Table 1). As seen in Fig. 1, each of these interactions generates rings of graph-set motif S(5) or S(6) (Bernstein *et al.*, 1995).

In the crystal structure, centrosymmetrically related molecules are linked by  $O5-H1O5\cdots O4^{i}$  intermolecular hydrogen bonds to form a dimeric pair (symmetry code as in Table 1). These hydrogen bonds form an  $R_2^2(8)$  ring motif. The adjacent dimers are interlinked by  $C9-H9B\cdots O2^{ii}$  hydrogen bonds to form a chain along [110]. These hydrogen bonds form an  $R_2^2(18)$  ring motif (Fig. 2). A C $-H\cdots\pi$  interaction involving atom H8B and the C1–C6 ring is observed, with atom H8B separated from the centroid (*Cg*1) of the ring by 2.58 Å (Table 1).

## Experimental

Succinic anhydride (1.0 g, 0.01 mol) was added to methyl anthranilate (0.5 g, 0.003 mol) in a round-bottomed flask containing dry toluene (50 ml). The reaction mixture was then refluxed for 15 h using a Dean–Stark trap. The reaction mixture was quickly filtered and left for crystallization at room temperature, resulting in colourless crystals of compound (I) (1.01 g, yield 85%, m.p. 539–540 K).

## Crystal data

C <sub>12</sub> H <sub>13</sub> NO <sub>5</sub>	$D_x = 1.376 \text{ Mg m}^{-3}$
$M_r = 251.23$	Mo $K\alpha$ radiation
Monoclinic, $C2/c$	Cell parameters from 5412
a = 18.301 (2) Å	reflections
b = 12.1788 (15) Å	$\theta = 2.0-26.5^{\circ}$
c = 10.8967 (14) Å	$\mu = 0.11 \text{ mm}^{-1}$
$\beta = 93.001 (2)^{\circ}$	T = 293 (2) K
V = 2425.3 (5) Å <sup>3</sup>	Block, colourless
Z = 8	$0.42 \times 0.27 \times 0.24 \text{ mm}$
Data collection	
Siemens SMART CCD area-	2498 independent reflections
detector diffractometer	2230 reflections with $I > 2\sigma(I)$
$\omega$ scans	$R_{\rm int} = 0.019$
Absorption correction: multi-scan	$\theta_{\rm max} = 26.5^{\circ}$
(SADABS: Sheldrick, 1996)	$h = -22 \rightarrow 22$
$T_{\rm min} = 0.956, T_{\rm max} = 0.975$	$k = -15 \rightarrow 14$
6736 measured reflections	$l = -12 \rightarrow 13$
Refinement	
Refinement on $F^2$	$w = 1/[\sigma^2(F_0^2) + (0.0551P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.042$	+ 1.0027P]
$wR(F^2) = 0.113$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.04	$(\Delta/\sigma)_{\rm max} = 0.001$
2498 reflections	$\Delta \rho_{\rm max} = 0.20 \ {\rm e} \ {\rm \AA}^{-3}$
169 parameters	$\Delta \rho_{\rm min} = -0.17 \ {\rm e} \ {\rm \AA}^{-3}$
H atoms treated by a mixture of	Extinction correction: SHELXTL

(Sheldrick, 1997)

Extinction coefficient: 0.0070 (10)



### Figure 2

Part of the crystal structure of (I), showing  $R_2^2(8)$  and  $R_2^2(18)$  ring motifs. Dashed lines indicate hydrogen bonds.

## Table 1

Hydrogen-bond geometry (Å, °).

Cg1 is the centroid of ring C1–C6.

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D - \mathbf{H} \cdot \cdot \cdot A$
N1-H1N···O2	0.86	1.94	2.6534 (17)	139
$O5-H1O5\cdots O4^{i}$	0.87(2)	1.80(2)	2.6657 (15)	173 (2)
$C2-H2 \cdot \cdot \cdot O1$	0.93	2.36	2.700 (2)	101
C5-H5···O3	0.93	2.27	2.881 (2)	123
$C9-H9B\cdots O2^{ii}$	0.97	2.42	3.3184 (19)	154
$C8-H8B\cdots Cg1^{iii}$	0.97	2.58	3.4470 (15)	149

Symmetry codes: (i) -x + 1, -y, -z; (ii)  $-x + \frac{1}{2}, -y + \frac{1}{2}, -z$ ; (iii)  $x, -y, z - \frac{1}{2}$ .

Atom H1O5 was located in a difference Fourier map and refined isotropically. All other H atoms were positioned geometrically and allowed to ride on their parent atoms, with N-H = 0.86 Å and C-H = 0.93–0.97 Å, and with  $U_{iso}(H) = 1.2$  or  $1.5U_{eq}(C,N)$ . A rotating-group model was used for the methyl group.

Data collection: *SMART* (Siemens, 1997); cell refinement: *SAINT* (Siemens, 1997); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 1997); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*, *PARST* (Nardelli, 1995) and *PLATON* (Spek, 2003).

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