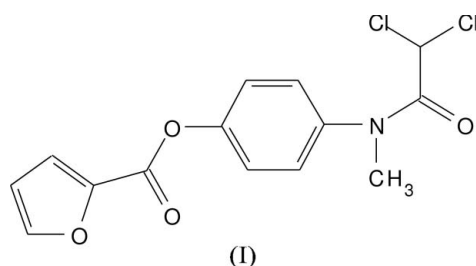


4-(2,2-Dichloro-*N*-methylacetamido)phenyl
furan-2-carboxylateS. Naveen,^a Dinesh Manvar,^b
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Key indicators

Single-crystal X-ray study
 $T = 295$ K
Mean $\sigma(\text{C}-\text{C}) = 0.004$ Å
 R factor = 0.040
 wR factor = 0.143
Data-to-parameter ratio = 13.1For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.In the title compound, $\text{C}_{14}\text{H}_{11}\text{Cl}_2\text{NO}_4$, the dihedral angle
between the two aromatic rings is $47.39(13)^\circ$. The crystal
packing is stabilized by $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds.Received 19 June 2006
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Comment

Amoebiasis is a major public health problem in tropical and
subtropical countries (McAuley *et al.*, 1992). Diloxanide
furoate, (I), is the mainstay for treating asymptomatic cyst
carriers and one of the oldest dichloroacetamide derivative
drugs for the treatment of amoebiasis. A crystal structure
determination was carried out in order to elucidate the mol-
ecular conformation.

A perspective view of (I) is shown in Fig. 1. In (I), the furan ring is planar. The dihedral angle between the furan ring and the benzene ring is $47.39(13)^\circ$. The keto group is almost coplanar with the furan ring, as indicated by the $\text{C}1-\text{C}2-\text{C}6-\text{O}7$ torsion angle of $-8.5(4)^\circ$, and is in a synperiplanar conformation [$\text{O}7-\text{C}6-\text{O}8-\text{C}9 = 8.3(3)^\circ$]. The other keto group adjacent to the *N*-methyl unit is in an antiperiplanar conformation, as indicated by the $\text{C}12-\text{N}15-\text{C}17-\text{O}18$ torsion angle of $177.9(2)^\circ$. The 2,2-dichloroacetylmethylamino group at the 4 position of the benzene ring is rotated $80.3(1)^\circ$ from the plane of the benzene ring. The crystal packing is stabilized by $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonding (Table 1). The molecules form hydrogen-bonded dimers.

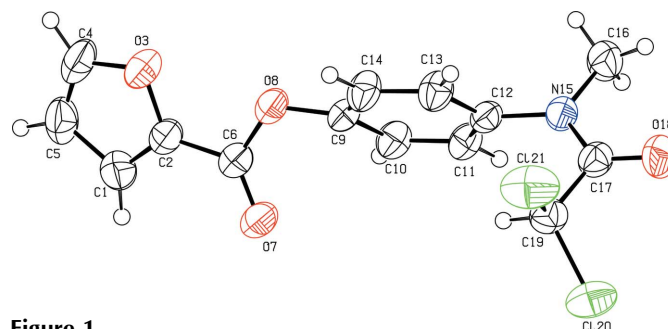


Figure 1
The molecular structure of (I), with 50% probability displacement ellipsoids.

Experimental

Diloxanide furoate (1.5 g) was dissolved in methanol (30 ml). Charcoal (3.5 g) was added and the mixture was stirred with heating for 6 minutes on a Deepali Stirrer (MS-4) at 1150 r.p.m. The hot solution was filtered through Whatmann 42 filter paper and kept in a slightly open, stoppered conical flask. Rectangular pale-green crystals grew by thin-film evaporation.

Crystal data

$C_{14}H_{11}Cl_2NO_4$	$Z = 4$
$M_r = 328.14$	$D_x = 1.522 \text{ Mg m}^{-3}$
Monoclinic, $P2_1/n$	Mo $K\alpha$ radiation
$a = 8.334 (5) \text{ \AA}$	$\mu = 0.47 \text{ mm}^{-1}$
$b = 10.778 (5) \text{ \AA}$	$T = 295 (2) \text{ K}$
$c = 16.412 (10) \text{ \AA}$	Block, pale green
$\beta = 103.761 (2)^\circ$	$0.25 \times 0.24 \times 0.22 \text{ mm}$
$V = 1431.9 (14) \text{ \AA}^3$	

Data collection

MacScience DIPLabo 32001 diffractometer	2524 independent reflections
ω scans	2209 reflections with $I > 2\sigma(I)$
Absorption correction: none	$R_{\text{int}} = 0.018$
4433 measured reflections	$\theta_{\text{max}} = 25.0^\circ$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0816P)^2 + 0.2846P]$
$R[F^2 > 2\sigma(F^2)] = 0.040$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.143$	$(\Delta/\sigma)_{\text{max}} < 0.001$
$S = 1.20$	$\Delta\rho_{\text{max}} = 0.35 \text{ e \AA}^{-3}$
2524 reflections	$\Delta\rho_{\text{min}} = -0.30 \text{ e \AA}^{-3}$
192 parameters	Extinction correction: <i>SHELXL97</i>
H-atom parameters constrained	Extinction coefficient: 0.056 (5)

Table 1

Hydrogen-bond geometry (\AA , $^\circ$).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$C4-H4\cdots O18^i$	0.93	2.41	3.285 (4)	156

Symmetry code: (i) $x - \frac{3}{2}, -y + \frac{1}{2}, z - \frac{1}{2}$.

H atoms were placed at idealized positions and allowed to ride on their parent atoms, with $C-H = 0.92-0.98 \text{ \AA}$ and $N-H = 0.86 \text{ \AA}$; $U_{\text{iso}}(\text{H})$ values were set equal to $xU_{\text{eq}}(\text{carrier atom})$, where $x = 1.5$ for methyl H atoms and $x = 1.2$ for all other H atoms.

Data collection: *XPRESS* (MacScience, 2002); cell refinement: *SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *SCALEPACK* and *DENZO* (Otwinowski & Minor, 1997); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2003) and *ORTEPII* (Johnson,

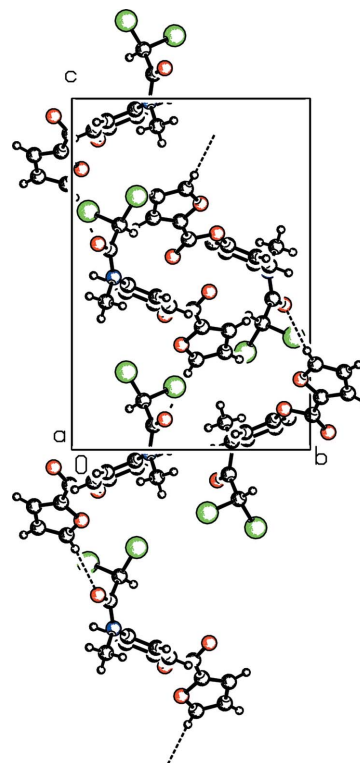


Figure 2

The packing of (I), viewed down the c axis. Dashed lines indicate hydrogen bonds.

1976); software used to prepare material for publication: *PLATON* (Spek, 2003).

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