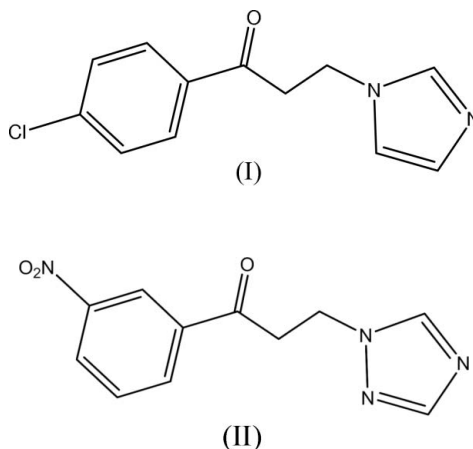
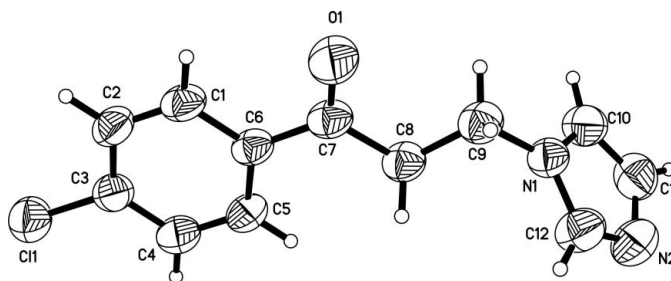


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**Key indicators**Single-crystal X-ray study  
 $T = 293\text{ K}$   
Mean  $\sigma(\text{C}-\text{C}) = 0.002\text{ \AA}$   
 $R$  factor = 0.041  
 $wR$  factor = 0.112  
Data-to-parameter ratio = 15.3For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.**1-(4-Chlorophenyl)-3-(1*H*-imidazol-1-yl)-  
propan-1-one**The molecule of the title compound,  $\text{C}_{12}\text{H}_{11}\text{ClN}_2\text{O}$ , is non-  
planar, the benzene and imidazole rings making a dihedral  
angle of  $72.5(1)^\circ$ . Molecules are linked into a three-  
dimensional framework by weak intermolecular  $\text{C}-\text{H}\cdots\text{O}$   
and  $\text{C}-\text{H}\cdots\text{N}$  hydrogen bonds.Received 7 July 2005  
Accepted 19 July 2005  
Online 23 July 2005**Comment**A molecular view of the title compound, (I), is shown in Fig. 1.  
The bond lengths and angles are within normal ranges (Allen  
*et al.*, 1987) and compare well with those reported in the  
related compound, (II) (Wan *et al.*, 2005). In contrast to the  
planar conformation of (II), compound (I) is non-planar, with  
the substituted benzene ring and the imidazole ring making a  
dihedral angle of  $72.5(1)^\circ$ .In the crystal structure, molecules are linked into a three-  
dimensional framework by weak intermolecular  $\text{C}-\text{H}\cdots\text{O}$   
and  $\text{C}-\text{H}\cdots\text{N}$  hydrogen bonds (Fig. 2 and Table 1).**Figure 1**  
The structure of the compound (I), showing 50% probability displace-  
ment ellipsoids and the atom-numbering scheme.

Experimental

To a solution of 1-(4-chlorophenyl)-3-(dimethylamino)propan-1-one (10.6 g, 0.05 mol) in water (25 ml) was added imidazole (4.4 g, 0.06 mol). The mixture was heated under reflux for 5 h, yielding a copious precipitate. Colorless single crystals suitable for X-ray diffraction study were obtained by slow evaporation of a dichloromethane–cyclohexane (1:2, v/v) solution over a period of 5 d.

Crystal data

$C_{12}H_{11}ClN_2O$   $D_x = 1.375 \text{ Mg m}^{-3}$   
 $M_r = 234.68$  Mo  $K\alpha$  radiation  
 Monoclinic,  $P2_1/n$  Cell parameters from 2278 reflections  
 $a = 10.4902 (9) \text{ \AA}$   $\theta = 2.9\text{--}25.9^\circ$   
 $b = 8.1611 (7) \text{ \AA}$   $\mu = 0.32 \text{ mm}^{-1}$   
 $c = 13.8691 (12) \text{ \AA}$   $T = 293 (2) \text{ K}$   
 $\beta = 107.252 (1)^\circ$  Plate, colorless  
 $V = 1133.94 (17) \text{ \AA}^3$   $0.49 \times 0.26 \times 0.06 \text{ mm}$   
 $Z = 4$

Data collection

Siemens SMART 1000 CCD area-detector diffractometer 2218 independent reflections  
 $\omega$  scans 1860 reflections with  $I > 2\sigma(I)$   
 Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)  $R_{int} = 0.017$   
 $T_{min} = 0.861, T_{max} = 0.981$   $\theta_{max} = 26.0^\circ$   
 6176 measured reflections  $h = -12 \rightarrow 12$   
 $k = -10 \rightarrow 9$   
 $l = -17 \rightarrow 13$

Refinement

Refinement on  $F^2$   $w = 1/[\sigma^2(F_o^2) + (0.0533P)^2 + 0.2452P]$   
 $R[F^2 > 2\sigma(F^2)] = 0.041$  where  $P = (F_o^2 + 2F_c^2)/3$   
 $wR(F^2) = 0.112$   $(\Delta\sigma)_{max} < 0.001$   
 $S = 1.05$   $\Delta\rho_{max} = 0.18 \text{ e \AA}^{-3}$   
 2218 reflections  $\Delta\rho_{min} = -0.19 \text{ e \AA}^{-3}$   
 145 parameters  
 H-atom parameters constrained

Table 1 Hydrogen-bond geometry ( $\text{\AA}, ^\circ$ ).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$C4-H4\cdots O1^i$	0.93	2.51	3.353 (2)	151
$C8-H8A\cdots N2^{ii}$	0.97	2.57	3.499 (2)	160

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

All H atoms were located in difference Fourier maps but they were treated as riding on their parent atoms [ $C-H = 0.93$  and  $0.97 \text{ \AA}$ , and  $U_{iso}(H) = 1.2U_{eq}(C)$ ].

Data collection: *SMART* (Siemens, 1996); cell refinement: *SAINT* (Siemens, 1996); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 1997); program(s) used to refine

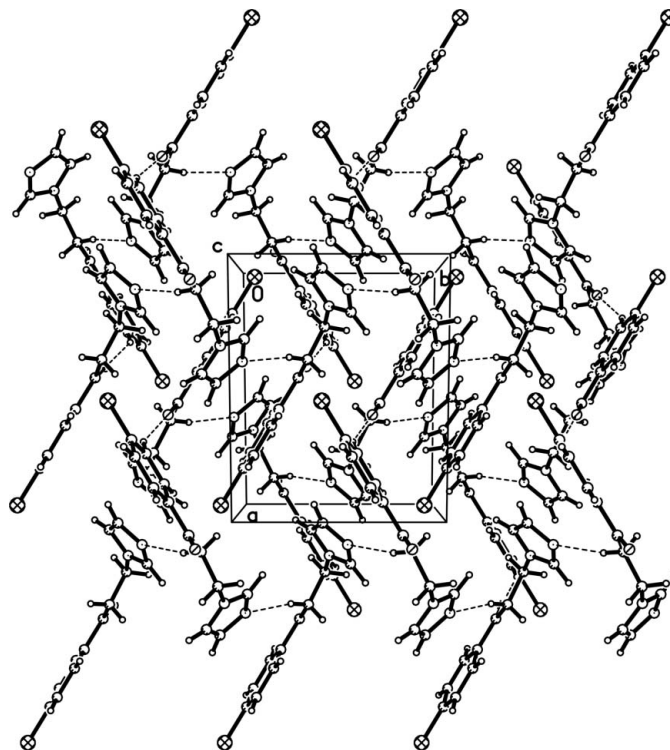


Figure 2 A view down the  $c$  axis. Hydrogen bonds are indicated by dashed lines.

structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*, *PARST* (Nardelli, 1995) and *PLATON* (Spek, 2003).

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