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Key indicatorsSingle-crystal X-ray study
 $T = 293\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.004\text{ \AA}$
 R factor = 0.038
 wR factor = 0.092
Data-to-parameter ratio = 7.5For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.**1-(3-Nitrophenyl)-3-(1*H*-1,2,4-triazol-1-yl)-propan-1-one**

In the title compound, $\text{C}_{11}\text{H}_{10}\text{N}_4\text{O}_3$, the substituted phenyl ring and the triazole ring make a dihedral angle of $59.7(2)^\circ$. The molecules are linked into zigzag chains by intermolecular $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds. These chains are further connected through $\text{C}-\text{H}\cdots\text{N}$ hydrogen bonds to build up a three-dimensional network.

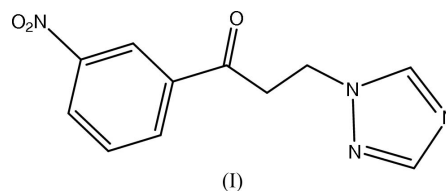
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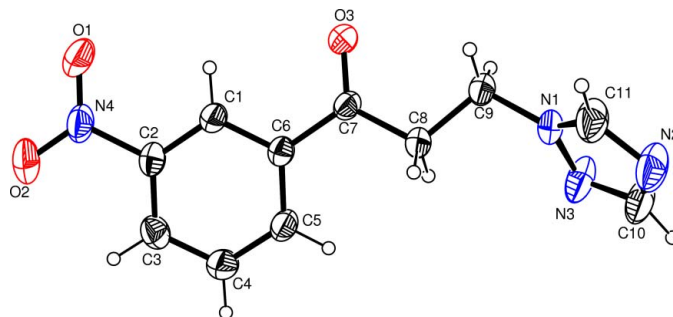
A molecular view of the title compound, (I), is shown in Fig. 1. The bond lengths and angles are within normal ranges (Allen *et al.*, 1987) and compare well with those reported for related compounds (Wan, Li, Li & Zhang, 2005; Wan, Li, Li, Zhang *et al.*, 2005). The molecule of (I) is non-planar, with the substituted phenyl ring and the triazole ring making a dihedral angle of $59.7(2)^\circ$.



In the crystal structure, the molecules of (I) are linked into zigzag chains by $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds (Fig. 2, Table 1). Further $\text{C}-\text{H}\cdots\text{N}$ hydrogen bonds connect these chains to build up a three-dimensional network which stabilizes the packing.

Experimental

To a solution of 3-(dimethylamino)-1-(3-nitrophenyl)propan-1-one hydrochloride (18 g, 0.07 mol) in water (20 ml) was added triazole (5.5 g, 0.08 mol). The mixture was heated under reflux for 5 h,

**Figure 1**

A view of compound (I), with the atom-labelling scheme. Ellipsoids are drawn at the 30% probability level.

yielding a copious precipitate. Colourless single crystals suitable for X-ray diffraction study were obtained by slow evaporation of an ethyl acetate–petroleum ether (1:1 (v/v) solution over a period of two weeks.

Crystal data

$C_{11}H_{10}N_4O_3$
 $M_r = 246.23$
 Orthorhombic, $Pca2_1$
 $a = 26.334 (8) \text{ \AA}$
 $b = 4.5962 (13) \text{ \AA}$
 $c = 9.570 (3) \text{ \AA}$
 $V = 1158.4 (6) \text{ \AA}^3$
 $Z = 4$
 $D_x = 1.412 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation
 Cell parameters from 1871 reflections
 $\theta = 2.6\text{--}22.3^\circ$
 $\mu = 0.11 \text{ mm}^{-1}$
 $T = 293 (2) \text{ K}$
 Thick plate, colourless
 $0.26 \times 0.24 \times 0.13 \text{ mm}$

Data collection

Siemens SMART 1000 CCD area-detector diffractometer
 ω scans
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.973$, $T_{\max} = 0.986$
 5794 measured reflections

1215 independent reflections
 1093 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.023$
 $\theta_{\text{max}} = 26.0^\circ$
 $h = -31 \rightarrow 32$
 $k = -5 \rightarrow 5$
 $l = -11 \rightarrow 11$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.038$
 $wR(F^2) = 0.092$
 $S = 1.14$
 1215 reflections
 163 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0426P)^2 + 0.1225P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.14 \text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.13 \text{ e \AA}^{-3}$

Table 1

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$C3-H3B\cdots O3^i$	0.93	2.48	3.400 (4)	172
$C4-H4A\cdots O1^i$	0.93	2.55	3.481 (5)	175
$C5-H5A\cdots N2^{ii}$	0.93	2.53	3.449 (4)	169

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

All H atoms were located in difference Fourier maps and constrained to ride on their parent atoms, with C–H distances in the range 0.93–0.97 \AA , and with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$. The Friedel reflections were merged before the final refinement because of the absence of any significant anomalous scattering effects.

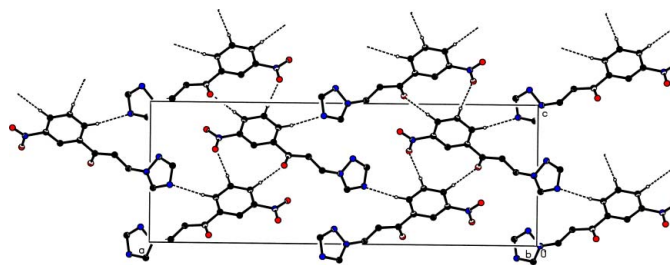


Figure 2

A view down the b axis, showing the C–H \cdots O and C–H \cdots N hydrogen bonds (dashed lines) leading to a three-dimensional network. H atoms not involved in hydrogen-bond interactions have been omitted for clarity.

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 structure: SHELXTL; molecular graphics: ORTEP3 (Farrugia, 1997), PLATON (Spek, 2003) and SHELXTL; software used to prepare material for publication: SHELXTL, PARST (Nardelli, 1995) and PLATON.

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