

Acta Crystallographica Section E

Structure Reports

Online

ISSN 1600-5368

4-Benzyl-1-*p*-tolyl-1*H*-1,2,4-triazol-5(4*H*)-oneYu-Jin Zhu^{a,b} and Hong-Quan Duan^{a*}

^aSchool of Pharmacy, Tianjin Medical University, Tianjin 300070, People's Republic of China, and ^bSchool of Chinese Armed Police Force Medical College, Tianjin 300162, People's Republic of China

Correspondence e-mail: duanhongquan2009@yahoo.cn

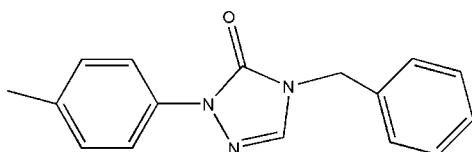
Received 20 January 2009; accepted 20 January 2009

Key indicators: single-crystal X-ray study; $T = 113$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.055; wR factor = 0.150; data-to-parameter ratio = 12.7.

In the title compound, $\text{C}_{16}\text{H}_{15}\text{N}_3\text{O}$, the triazole ring makes dihedral angles of 7.08 (2) and 74.53 (3)° with the two outer aromatic rings. The crystal packing is stabilized by very short intermolecular $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds and weak $\pi-\pi$ stacking interactions [centroid-to-centroid distance 3.632 (3) Å], resulting in the formation of zigzag chains parallel to the b axis.

Related literature

For details of the biological activity of trisubstituted triazolones, see: Chang *et al.* (1993, 1994). For bond-length data, see: Allen *et al.* (1987). For details of synthesis, see: Theodoridis (1998).



Experimental

Crystal data

$\text{C}_{16}\text{H}_{15}\text{N}_3\text{O}$
 $M_r = 265.31$
 Monoclinic, $P2_1/n$
 $a = 4.6130$ (9) Å
 $b = 25.488$ (5) Å
 $c = 11.460$ (2) Å
 $\beta = 96.18$ (3)°
 $V = 1339.6$ (5) Å³
 $Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 0.09$ mm⁻¹
 $T = 113$ (2) K
 $0.18 \times 0.04 \times 0.04$ mm

Data collection

Rigaku Saturn diffractometer
 Absorption correction: multi-scan (*CrystalClear*; Rigaku, 2005)
 $T_{\min} = 0.985$, $T_{\max} = 0.997$
 928 measured reflections
 2333 independent reflections
 1998 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.052$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.055$
 $wR(F^2) = 0.150$
 $S = 1.10$
 2333 reflections
 183 parameters
 H-atom parameters constrained
 $\Delta\rho_{\max} = 0.23$ e Å⁻³
 $\Delta\rho_{\min} = -0.26$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{C9}-\text{H9}\cdots\text{O1}^i$	0.93	2.19	3.114 (2)	174

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

Data collection: *CrystalClear* (Rigaku, 2005); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 2008); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HG2470).

References

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supporting information

Acta Cryst. (2009). E65, o384 [doi:10.1107/S1600536809002487]

4-Benzyl-1-*p*-tolyl-1*H*-1,2,4-triazol-5(4*H*)-one

Yu-Jin Zhu and Hong-Quan Duan

S1. Comment

4-Benzyl-1-*p*-tolyl-1*H*-1,2,4-triazol-5(4*H*)-one is a N-substituted triazolinone. It was reported that trisubstituted triazolinones were employed as nonpeptide angiotensin II receptor antagonists (Chang *et al.*, 1993, 1994). In our effort to further study triazolinone derivatives as novel AII antagonists, the title compound was prepared. Here, we report the crystal structure of it.

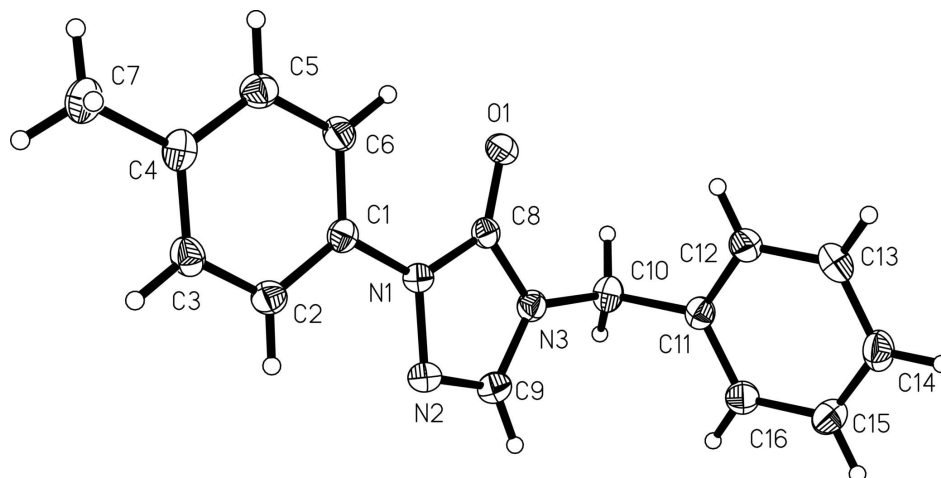
In title compound, all bond lengths in the molecule are normal (Allen *et al.*, 1987). The triazole ring N1–N3/C8–C9 makes dihedral angles of 7.08 (2) and 74.53 (3)° with the two phenyl rings (C1–C6, C11–C16). The relatively short distance of 3.632 (3) Å between the centroids of triazole ring N1–N3/C8–C9 and benzene ring C1–C6 [at $-1 + x, y, z$] indicates the presence of weak π – π interactions. The crystal packing is stabilized by intermolecular C—H \cdots O hydrogen bonds, linking the molecules into zigzag chains parallel to the *b* axis.

S2. Experimental

1-*p*-Tolyl-1*H*-1,2,4-triazol-5(4*H*)one (1.75 g, 0.01 mol) was dissolved in 30 ml of acetic anhydride, 1.38 g (0.01 mol) potassium carbonate and 0.75 ml (0.01 mol) phenylmethanol were added. The solution was heated to reflux and stirred for 2 h and then cooled to room temperature. 100 ml of water was added and the deposited precipitate filtered. The precipitate was recrystallized with acetone and dried to give 4-benzyl-1-*p*-tolyl-1*H*-1,2,4-triazol-5(4*H*)one as a colorless powder (2.40 g, yield 90.5%) (Theodoridis, 1998). Crystals suitable for X-ray diffraction were obtained through slow evaporation of the solution of the title compound in dichloromethane and ethyl acetate (v/v 1:1).

S3. Refinement

All H atoms were placed in calculated positions, with C—H = 0.93–0.97 Å, and included in the final cycles of refinement using a riding model, with $U_{\text{iso}}(\text{H}) = 1.2$ (1.5 times for methyl) times $U_{\text{eq}}(\text{C})$.

**Figure 1**

View of the title compound (I), with displacement ellipsoids drawn at the 40% probability level.

4-Benzyl-1-*p*-tolyl-1*H*-1,2,4-triazol-5(4*H*)-one

Crystal data

$C_{16}H_{15}N_3O$

$M_r = 265.31$

Monoclinic, $P2_1/n$

Hall symbol: $-P\ 2_1n$

$a = 4.6130\ (9)\ \text{\AA}$

$b = 25.488\ (5)\ \text{\AA}$

$c = 11.460\ (2)\ \text{\AA}$

$\beta = 96.18\ (3)^\circ$

$V = 1339.6\ (5)\ \text{\AA}^3$

$Z = 4$

$F(000) = 560$

$D_x = 1.315\ \text{Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073\ \text{\AA}$

Cell parameters from 2910 reflections

$\theta = 1.6\text{--}27.9^\circ$

$\mu = 0.09\ \text{mm}^{-1}$

$T = 113\ \text{K}$

Block, colourless

$0.18 \times 0.04 \times 0.04\ \text{mm}$

Data collection

Rigaku Saturn
diffractometer

Radiation source: rotating anode

Confocal monochromator

ω scans

Absorption correction: multi-scan

(*CrystalClear*; Rigaku, 2005)

$T_{\min} = 0.985$, $T_{\max} = 0.997$

9828 measured reflections

2333 independent reflections

1998 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.052$

$\theta_{\max} = 25.0^\circ$, $\theta_{\min} = 2.0^\circ$

$h = -5 \rightarrow 5$

$k = -30 \rightarrow 30$

$l = -13 \rightarrow 13$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.055$

$wR(F^2) = 0.150$

$S = 1.10$

2333 reflections

183 parameters

0 restraints

Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier
map

Hydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0922P)^2]$

where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} < 0.001$

$\Delta\rho_{\max} = 0.23\ \text{e \AA}^{-3}$

$\Delta\rho_{\min} = -0.26\ \text{e \AA}^{-3}$

Extinction correction: *SHELXTL* (Sheldrick, 2008), $F_c^* = kFc[1 + 0.001xFc^2\lambda^3/\sin(2\theta)]^{-1/4}$

Extinction coefficient: 0.030 (5)

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) *etc.* and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	U_{iso}^*/U_{eq}
O1	0.4812 (3)	0.20822 (5)	0.27648 (11)	0.0298 (4)
N1	0.5904 (3)	0.17809 (5)	0.09066 (12)	0.0250 (4)
N2	0.5011 (4)	0.19124 (6)	-0.02578 (13)	0.0328 (4)
N3	0.2711 (3)	0.24033 (5)	0.09506 (13)	0.0267 (4)
C1	0.7986 (4)	0.13699 (6)	0.11575 (15)	0.0247 (4)
C2	0.8840 (4)	0.10765 (7)	0.02333 (16)	0.0295 (5)
H2	0.8061	0.1147	-0.0533	0.035*
C3	1.0867 (4)	0.06763 (7)	0.04622 (16)	0.0310 (5)
H3	1.1441	0.0483	-0.0162	0.037*
C4	1.2065 (4)	0.05553 (7)	0.15959 (17)	0.0281 (5)
C5	1.1164 (4)	0.08567 (7)	0.25051 (17)	0.0305 (5)
H5	1.1928	0.0784	0.3272	0.037*
C6	0.9155 (4)	0.12638 (7)	0.23019 (16)	0.0284 (5)
H6	0.8605	0.1462	0.2923	0.034*
C7	1.4263 (4)	0.01155 (7)	0.18258 (18)	0.0337 (5)
H7A	1.6175	0.0243	0.1720	0.051*
H7B	1.3764	-0.0167	0.1287	0.051*
H7C	1.4243	-0.0009	0.2616	0.051*
C8	0.4527 (4)	0.20823 (6)	0.16844 (16)	0.0242 (4)
C9	0.3125 (4)	0.22863 (7)	-0.01808 (17)	0.0328 (5)
H9	0.2161	0.2456	-0.0828	0.039*
C10	0.0864 (4)	0.28203 (7)	0.13604 (18)	0.0306 (5)
H10A	0.0539	0.2752	0.2169	0.037*
H10B	-0.1016	0.2817	0.0891	0.037*
C11	0.2243 (4)	0.33566 (7)	0.12782 (15)	0.0265 (5)
C12	0.4397 (4)	0.35251 (7)	0.21405 (16)	0.0326 (5)
H12	0.4966	0.3310	0.2779	0.039*
C13	0.5705 (5)	0.40130 (8)	0.20553 (18)	0.0370 (5)
H13	0.7158	0.4121	0.2631	0.044*
C14	0.4840 (5)	0.43383 (7)	0.11114 (17)	0.0374 (5)
H14	0.5691	0.4667	0.1057	0.045*
C15	0.2713 (5)	0.41719 (8)	0.02540 (18)	0.0388 (5)
H15	0.2142	0.4389	-0.0381	0.047*

C16	0.1420 (4)	0.36866 (7)	0.03274 (17)	0.0328 (5)
H16	-0.0006	0.3579	-0.0259	0.039*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0334 (8)	0.0304 (7)	0.0250 (7)	0.0016 (5)	0.0005 (6)	-0.0033 (5)
N1	0.0277 (9)	0.0245 (8)	0.0224 (8)	0.0005 (6)	0.0004 (6)	0.0006 (6)
N2	0.0443 (10)	0.0300 (9)	0.0235 (9)	0.0029 (8)	0.0013 (7)	0.0019 (7)
N3	0.0273 (9)	0.0239 (8)	0.0284 (9)	0.0020 (6)	0.0012 (7)	0.0001 (6)
C1	0.0235 (10)	0.0226 (9)	0.0282 (10)	-0.0032 (7)	0.0032 (8)	0.0008 (7)
C2	0.0331 (11)	0.0297 (10)	0.0266 (10)	-0.0023 (8)	0.0066 (8)	0.0018 (7)
C3	0.0328 (11)	0.0295 (10)	0.0326 (11)	-0.0003 (8)	0.0125 (9)	-0.0039 (8)
C4	0.0217 (10)	0.0250 (9)	0.0384 (11)	-0.0046 (7)	0.0063 (8)	0.0005 (8)
C5	0.0273 (11)	0.0337 (11)	0.0295 (10)	-0.0012 (8)	-0.0012 (8)	0.0006 (8)
C6	0.0288 (11)	0.0274 (10)	0.0285 (11)	0.0000 (8)	0.0013 (8)	-0.0038 (7)
C7	0.0286 (11)	0.0288 (10)	0.0440 (12)	0.0015 (8)	0.0053 (9)	0.0009 (9)
C8	0.0244 (10)	0.0217 (9)	0.0258 (10)	-0.0045 (7)	0.0002 (7)	-0.0009 (7)
C9	0.0377 (12)	0.0308 (10)	0.0287 (11)	0.0021 (8)	-0.0019 (9)	0.0032 (8)
C10	0.0258 (11)	0.0271 (10)	0.0395 (11)	0.0012 (8)	0.0056 (9)	0.0002 (8)
C11	0.0265 (10)	0.0251 (9)	0.0292 (10)	0.0033 (7)	0.0088 (8)	-0.0008 (7)
C12	0.0383 (12)	0.0303 (10)	0.0292 (10)	0.0032 (8)	0.0035 (9)	-0.0023 (8)
C13	0.0376 (12)	0.0362 (11)	0.0365 (12)	-0.0032 (9)	0.0013 (9)	-0.0107 (9)
C14	0.0443 (13)	0.0280 (10)	0.0421 (13)	-0.0082 (9)	0.0151 (10)	-0.0037 (9)
C15	0.0486 (14)	0.0325 (11)	0.0360 (12)	-0.0005 (9)	0.0075 (10)	0.0066 (9)
C16	0.0332 (12)	0.0321 (10)	0.0319 (11)	0.0009 (8)	-0.0008 (9)	0.0002 (8)

Geometric parameters (Å, °)

O1—C8	1.231 (2)	C7—H7A	0.9600
N1—C8	1.382 (2)	C7—H7B	0.9600
N1—N2	1.394 (2)	C7—H7C	0.9600
N1—C1	1.429 (2)	C9—H9	0.9300
N2—C9	1.299 (2)	C10—C11	1.515 (2)
N3—C9	1.364 (2)	C10—H10A	0.9700
N3—C8	1.388 (2)	C10—H10B	0.9700
N3—C10	1.471 (2)	C11—C12	1.392 (3)
C1—C2	1.388 (2)	C11—C16	1.397 (3)
C1—C6	1.390 (2)	C12—C13	1.390 (3)
C2—C3	1.390 (3)	C12—H12	0.9300
C2—H2	0.9300	C13—C14	1.387 (3)
C3—C4	1.391 (3)	C13—H13	0.9300
C3—H3	0.9300	C14—C15	1.379 (3)
C4—C5	1.394 (3)	C14—H14	0.9300
C4—C7	1.516 (2)	C15—C16	1.379 (3)
C5—C6	1.394 (3)	C15—H15	0.9300
C5—H5	0.9300	C16—H16	0.9300
C6—H6	0.9300		

C8—N1—N2	112.01 (14)	O1—C8—N1	129.94 (16)
C8—N1—C1	128.55 (15)	O1—C8—N3	126.98 (16)
N2—N1—C1	119.43 (14)	N1—C8—N3	103.07 (15)
C9—N2—N1	104.01 (15)	N2—C9—N3	112.84 (16)
C9—N3—C8	108.06 (15)	N2—C9—H9	123.6
C9—N3—C10	127.37 (16)	N3—C9—H9	123.6
C8—N3—C10	124.41 (15)	N3—C10—C11	111.73 (14)
C2—C1—C6	120.14 (16)	N3—C10—H10A	109.3
C2—C1—N1	118.77 (16)	C11—C10—H10A	109.3
C6—C1—N1	121.10 (15)	N3—C10—H10B	109.3
C1—C2—C3	119.47 (17)	C11—C10—H10B	109.3
C1—C2—H2	120.3	H10A—C10—H10B	107.9
C3—C2—H2	120.3	C12—C11—C16	118.79 (17)
C2—C3—C4	122.05 (17)	C12—C11—C10	120.45 (16)
C2—C3—H3	119.0	C16—C11—C10	120.75 (17)
C4—C3—H3	119.0	C13—C12—C11	120.55 (18)
C3—C4—C5	117.13 (17)	C13—C12—H12	119.7
C3—C4—C7	121.17 (17)	C11—C12—H12	119.7
C5—C4—C7	121.71 (17)	C14—C13—C12	119.93 (19)
C4—C5—C6	122.13 (18)	C14—C13—H13	120.0
C4—C5—H5	118.9	C12—C13—H13	120.0
C6—C5—H5	118.9	C15—C14—C13	119.65 (18)
C1—C6—C5	119.08 (17)	C15—C14—H14	120.2
C1—C6—H6	120.5	C13—C14—H14	120.2
C5—C6—H6	120.5	C14—C15—C16	120.78 (19)
C4—C7—H7A	109.5	C14—C15—H15	119.6
C4—C7—H7B	109.5	C16—C15—H15	119.6
H7A—C7—H7B	109.5	C15—C16—C11	120.30 (18)
C4—C7—H7C	109.5	C15—C16—H16	119.8
H7A—C7—H7C	109.5	C11—C16—H16	119.8
H7B—C7—H7C	109.5		
C8—N1—N2—C9	-0.07 (19)	C9—N3—C8—O1	178.18 (18)
C1—N1—N2—C9	179.13 (15)	C10—N3—C8—O1	2.4 (3)
C8—N1—C1—C2	172.54 (16)	C9—N3—C8—N1	-0.95 (18)
N2—N1—C1—C2	-6.5 (2)	C10—N3—C8—N1	-176.71 (14)
C8—N1—C1—C6	-7.7 (3)	N1—N2—C9—N3	-0.6 (2)
N2—N1—C1—C6	173.26 (15)	C8—N3—C9—N2	1.0 (2)
C6—C1—C2—C3	0.1 (3)	C10—N3—C9—N2	176.61 (16)
N1—C1—C2—C3	179.85 (15)	C9—N3—C10—C11	-75.1 (2)
C1—C2—C3—C4	0.5 (3)	C8—N3—C10—C11	99.78 (19)
C2—C3—C4—C5	-0.5 (3)	N3—C10—C11—C12	-80.5 (2)
C2—C3—C4—C7	179.72 (15)	N3—C10—C11—C16	98.3 (2)
C3—C4—C5—C6	-0.1 (3)	C16—C11—C12—C13	-0.1 (3)
C7—C4—C5—C6	179.67 (16)	C10—C11—C12—C13	178.78 (17)
C2—C1—C6—C5	-0.7 (3)	C11—C12—C13—C14	0.7 (3)
N1—C1—C6—C5	179.57 (15)	C12—C13—C14—C15	-0.9 (3)

C4—C5—C6—C1	0.7 (3)	C13—C14—C15—C16	0.4 (3)
N2—N1—C8—O1	-178.45 (17)	C14—C15—C16—C11	0.3 (3)
C1—N1—C8—O1	2.4 (3)	C12—C11—C16—C15	-0.4 (3)
N2—N1—C8—N3	0.64 (18)	C10—C11—C16—C15	-179.28 (17)
C1—N1—C8—N3	-178.46 (15)		

Hydrogen-bond geometry (Å, °)

<i>D—H...A</i>	<i>D—H</i>	<i>H...A</i>	<i>D...A</i>	<i>D—H...A</i>
C9—H9...O1 ⁱ	0.93	2.19	3.114 (2)	174

Symmetry code: (i) $x-1/2, -y+1/2, z-1/2$.