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## Structure Reports

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## 1-(4-Methylphenyl)-1H-1,2,3,4-tetrazole

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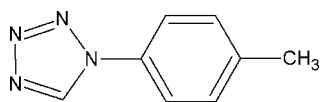
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Key indicators: single-crystal X-ray study;  $T = 293$  K; mean  $\sigma(\text{C}-\text{C}) = 0.004$  Å;  $R$  factor = 0.067;  $wR$  factor = 0.211; data-to-parameter ratio = 12.9.

In the title compound,  $\text{C}_8\text{H}_8\text{N}_4$ , the dihedral angle between the tetrazole and benzene rings is  $21.6$  ( $1$ )°. An intermolecular  $\text{C}-\text{H}\cdots\pi$  interaction is observed.

## Related literature

For background to and applications of tetrazole derivatives, see: Singh *et al.* (1980); Brown (1967); Ostrovskii *et al.* (1999). For the synthesis, see: Aridoss & Laali (2011). For related structures, see: Matsunaga *et al.* (1999); Lyakhov *et al.* (2000).



## Experimental

## Crystal data

$\text{C}_8\text{H}_8\text{N}_4$   
 $M_r = 160.18$   
Monoclinic,  $P2_1/n$   
 $a = 9.8352$  (13) Å  
 $b = 5.7244$  (6) Å  
 $c = 14.4190$  (19) Å  
 $\beta = 96.285$  ( $12$ )°

$V = 806.92$  (17) Å<sup>3</sup>  
 $Z = 4$   
Mo  $K\alpha$  radiation  
 $\mu = 0.09$  mm<sup>-1</sup>  
 $T = 293$  K  
 $0.3 \times 0.2 \times 0.1$  mm

## Data collection

Oxford Diffraction Xcalibur  
Sapphire3 diffractometer  
Absorption correction: multi-scan  
(*CrysAlis PRO*; Oxford  
Diffraction, 2010)  
 $T_{\min} = 0.762$ ,  $T_{\max} = 1.000$

14618 measured reflections  
1419 independent reflections  
936 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.054$

## Refinement

$R[F^2 > 2\sigma(F^2)] = 0.067$   
 $wR(F^2) = 0.211$   
 $S = 1.05$   
1419 reflections

110 parameters  
H-atom parameters constrained  
 $\Delta\rho_{\max} = 0.21$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.25$  e Å<sup>-3</sup>

Table 1

Hydrogen-bond geometry (Å, °).

Cg is the centroid of the C2–C7 ring.

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{C6}-\text{H6}\cdots\text{Cg}^i$	0.93	2.89	3.630 (3)	138

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

Data collection: *CrysAlis PRO* (Oxford Diffraction, 2010); cell refinement: *CrysAlis PRO*; data reduction: *CrysAlis RED* (Oxford Diffraction, 2010); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *PLATON* (Spek, 2009); software used to prepare material for publication: *PLATON*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: IS5047).

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## supporting information

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## 1-(4-Methylphenyl)-1*H*-1,2,3,4-tetrazole

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### S1. Comment

Tetrazoles are an important functional group with wide range of applications (Aridoss & Laali, 2011). They function as ligands in coordination chemistry, in medicinal chemistry as a metabolically stable surrogate for a carboxylic acid group (Singh *et al.*, 1980) and in materials science applications including propellants (Brown, 1967) and explosives (Ostrovskii *et al.*, 1999).

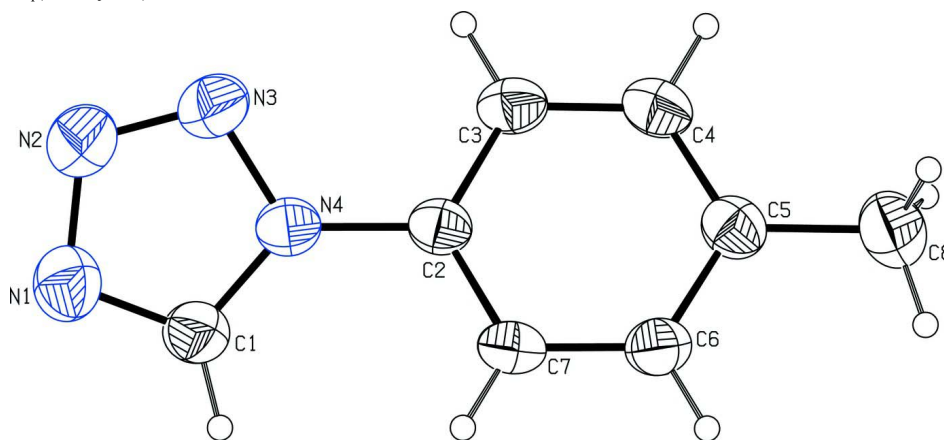
Bond lengths and bond angles are comparable with the similar crystal structures (Matsunaga *et al.*, 1999; Lyakhov *et al.*, 2000). The tetrazole and benzene rings are planar with maximum deviations of 0.006 (2) and 0.011 (2) Å (r.m.s. deviations of the rings being 0.005 and 0.008 Å), respectively. The two rings are not coplanar with the dihedral angle being 21.6 (1)°. Methyl carbon atom lies 0.014 (5) Å from the plane of the phenyl ring. Bond distances C1—N1 [1.304 (4) Å] and N2—N3 [1.289 (4) Å] indicate the consistence of the formation of double bonds.

### S2. Experimental

The title compound was synthesized from the known procedure reported elsewhere (Aridoss & Laali, 2011). Fine white diffraction quality crystals were obtained from the slow evaporation of its solution in ethanol.

### S3. Refinement

All H atoms were refined using a riding model, with C—H = 0.93 Å (aromatic) or 0.96 Å (methyl), and with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$  or  $1.5U_{\text{eq}}(\text{methyl C})$ .



**Figure 1**

The molecular structure of the title compound, showing 30% probability displacement ellipsoids.

**1-(4-Methylphenyl)-1H-1,2,3,4-tetrazole***Crystal data*C<sub>8</sub>H<sub>8</sub>N<sub>4</sub> $M_r = 160.18$ Monoclinic,  $P2_1/n$ 

Hall symbol: -P 2yn

 $a = 9.8352$  (13) Å $b = 5.7244$  (6) Å $c = 14.4190$  (19) Å $\beta = 96.285$  (12)° $V = 806.92$  (17) Å<sup>3</sup> $Z = 4$  $F(000) = 336$  $D_x = 1.319$  Mg m<sup>-3</sup>Mo  $K\alpha$  radiation,  $\lambda = 0.71073$  Å

Cell parameters from 3527 reflections

 $\theta = 3.6$ – $29.2$ ° $\mu = 0.09$  mm<sup>-1</sup> $T = 293$  K

Plate, white

 $0.3 \times 0.2 \times 0.1$  mm*Data collection*Oxford Diffraction Xcalibur Sapphire3  
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

Detector resolution: 16.1049 pixels mm<sup>-1</sup> $\omega$  scans

Absorption correction: multi-scan

(CrysAlis PRO; Oxford Diffraction, 2010)

 $T_{\min} = 0.762$ ,  $T_{\max} = 1.000$ 

14618 measured reflections

1419 independent reflections

936 reflections with  $I > 2\sigma(I)$  $R_{\text{int}} = 0.054$  $\theta_{\max} = 25.0$ °,  $\theta_{\min} = 3.8$ ° $h = -11 \rightarrow 11$  $k = -6 \rightarrow 6$  $l = -17 \rightarrow 17$ *Refinement*Refinement on  $F^2$ 

Least-squares matrix: full

 $R[F^2 > 2\sigma(F^2)] = 0.067$  $wR(F^2) = 0.211$  $S = 1.05$ 

1419 reflections

110 parameters

0 restraints

Primary atom site location: structure-invariant  
direct methodsSecondary atom site location: difference Fourier  
mapHydrogen site location: inferred from  
neighbouring sites

H-atom parameters constrained

 $w = 1/[\sigma^2(F_o^2) + (0.1133P)^2 + 0.1542P]$ where  $P = (F_o^2 + 2F_c^2)/3$  $(\Delta/\sigma)_{\max} < 0.001$  $\Delta\rho_{\max} = 0.21$  e Å<sup>-3</sup> $\Delta\rho_{\min} = -0.25$  e Å<sup>-3</sup>*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 (Å<sup>2</sup>)*

	$x$	$y$	$z$	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.7153 (3)	0.0846 (5)	0.8500 (2)	0.0726 (9)
H1	0.7271	-0.0729	0.8372	0.087*
C2	0.4656 (3)	0.0810 (4)	0.87347 (17)	0.0540 (7)
C3	0.3730 (3)	0.1934 (5)	0.92335 (18)	0.0635 (8)

H3	0.3959	0.3336	0.9537	0.076*
C4	0.2465 (3)	0.0950 (5)	0.9274 (2)	0.0672 (9)
H4	0.1838	0.1718	0.9605	0.081*
C5	0.2090 (3)	-0.1143 (5)	0.88414 (19)	0.0622 (8)
C6	0.3039 (3)	-0.2219 (5)	0.8337 (2)	0.0634 (8)
H6	0.2811	-0.3625	0.8036	0.076*
C7	0.4315 (3)	-0.1253 (4)	0.82696 (18)	0.0607 (8)
H7	0.4932	-0.1980	0.7918	0.073*
C8	0.0709 (3)	-0.2228 (6)	0.8897 (2)	0.0850 (10)
H8A	0.0534	-0.2328	0.9538	0.127*
H8B	0.0692	-0.3767	0.8631	0.127*
H8C	0.0017	-0.1283	0.8558	0.127*
N1	0.8114 (3)	0.2424 (5)	0.85255 (19)	0.0841 (9)
N2	0.7502 (3)	0.4443 (5)	0.8748 (2)	0.0874 (9)
N3	0.6234 (3)	0.4089 (4)	0.8848 (2)	0.0826 (9)
N4	0.5985 (2)	0.1809 (4)	0.86824 (14)	0.0580 (7)

*Atomic displacement parameters (Å<sup>2</sup>)*

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
C1	0.077 (2)	0.0625 (18)	0.081 (2)	0.0000 (16)	0.0255 (17)	-0.0042 (15)
C2	0.0584 (16)	0.0504 (15)	0.0528 (15)	0.0089 (11)	0.0041 (12)	0.0054 (11)
C3	0.079 (2)	0.0557 (16)	0.0568 (17)	0.0051 (14)	0.0129 (14)	-0.0082 (13)
C4	0.071 (2)	0.0697 (19)	0.0636 (18)	0.0125 (15)	0.0181 (15)	-0.0055 (14)
C5	0.0616 (18)	0.0644 (17)	0.0609 (17)	0.0079 (13)	0.0075 (14)	0.0067 (13)
C6	0.0678 (18)	0.0531 (16)	0.0686 (18)	0.0045 (13)	0.0046 (14)	-0.0049 (13)
C7	0.0661 (19)	0.0524 (15)	0.0635 (18)	0.0138 (13)	0.0073 (14)	-0.0052 (12)
C8	0.066 (2)	0.101 (3)	0.088 (2)	-0.0003 (17)	0.0104 (17)	-0.0046 (19)
N1	0.0828 (19)	0.086 (2)	0.087 (2)	-0.0101 (15)	0.0223 (15)	0.0017 (14)
N2	0.085 (2)	0.0721 (18)	0.106 (2)	-0.0069 (15)	0.0121 (16)	0.0022 (15)
N3	0.082 (2)	0.0569 (16)	0.108 (2)	0.0002 (13)	0.0060 (16)	-0.0030 (13)
N4	0.0649 (15)	0.0519 (13)	0.0568 (14)	0.0047 (11)	0.0048 (11)	0.0029 (10)

*Geometric parameters (Å, °)*

C1—N1	1.304 (4)	C5—C8	1.504 (4)
C1—N4	1.326 (3)	C6—C7	1.385 (4)
C1—H1	0.9300	C6—H6	0.9300
C2—C3	1.380 (4)	C7—H7	0.9300
C2—C7	1.381 (4)	C8—H8A	0.9600
C2—N4	1.436 (3)	C8—H8B	0.9600
C3—C4	1.373 (4)	C8—H8C	0.9600
C3—H3	0.9300	N1—N2	1.357 (4)
C4—C5	1.382 (4)	N2—N3	1.288 (4)
C4—H4	0.9300	N3—N4	1.345 (3)
C5—C6	1.389 (4)		
N1—C1—N4	110.3 (3)	C5—C6—H6	119.1

N1—C1—H1	124.8	C2—C7—C6	118.8 (3)
N4—C1—H1	124.8	C2—C7—H7	120.6
C3—C2—C7	120.8 (3)	C6—C7—H7	120.6
C3—C2—N4	119.9 (2)	C5—C8—H8A	109.5
C7—C2—N4	119.2 (2)	C5—C8—H8B	109.5
C4—C3—C2	119.0 (3)	H8A—C8—H8B	109.5
C4—C3—H3	120.5	C5—C8—H8C	109.5
C2—C3—H3	120.5	H8A—C8—H8C	109.5
C3—C4—C5	122.3 (2)	H8B—C8—H8C	109.5
C3—C4—H4	118.8	C1—N1—N2	105.0 (3)
C5—C4—H4	118.8	N3—N2—N1	110.6 (2)
C4—C5—C6	117.4 (3)	N2—N3—N4	107.0 (2)
C4—C5—C8	122.1 (3)	C1—N4—N3	107.1 (2)
C6—C5—C8	120.5 (3)	C1—N4—C2	131.2 (2)
C7—C6—C5	121.7 (3)	N3—N4—C2	121.7 (2)
C7—C6—H6	119.1		
C7—C2—C3—C4	1.0 (4)	C1—N1—N2—N3	-0.1 (4)
N4—C2—C3—C4	180.0 (2)	N1—N2—N3—N4	-0.6 (4)
C2—C3—C4—C5	0.7 (4)	N1—C1—N4—N3	-1.2 (3)
C3—C4—C5—C6	-1.4 (4)	N1—C1—N4—C2	180.0 (3)
C3—C4—C5—C8	179.3 (3)	N2—N3—N4—C1	1.1 (3)
C4—C5—C6—C7	0.3 (4)	N2—N3—N4—C2	-180.0 (2)
C8—C5—C6—C7	179.7 (3)	C3—C2—N4—C1	157.9 (3)
C3—C2—C7—C6	-2.0 (4)	C7—C2—N4—C1	-23.2 (4)
N4—C2—C7—C6	179.0 (2)	C3—C2—N4—N3	-20.7 (4)
C5—C6—C7—C2	1.3 (4)	C7—C2—N4—N3	158.2 (2)
N4—C1—N1—N2	0.9 (3)		

*Hydrogen-bond geometry* (Å, °)

C<sub>g</sub> is the centroid of the C2—C7 ring.

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
C6—H6...C <sub>g</sub> <sup>i</sup>	0.93	2.89	3.630 (3)	138

Symmetry code: (i)  $-x+1/2, y-1/2, -z+3/2$ .