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N-(Naphthalen-1-ylmethylidene)-4H-1,2,4-triazol-4-amine

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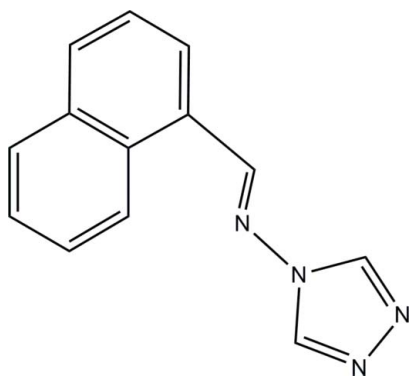
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Key indicators: single-crystal X-ray study; $T = 296$ K; mean $\sigma(\text{C}—\text{C}) = 0.002$ Å;
 R factor = 0.042; wR factor = 0.113; data-to-parameter ratio = 14.1.

In the title molecule, $\text{C}_{13}\text{H}_{10}\text{N}_4$, the dihedral angle between the triazole ring and the naphthalene ring system is $56.1(2)^\circ$. In the crystal, molecules are connected by weak $\text{C}—\text{H}\cdots\text{N}$ hydrogen bonds into chains along $[100]$. A short intramolecular $\text{C}—\text{H}\cdots\text{N}$ contact is also observed.

Related literature

For applications of triazole derivatives, see: Demirbas *et al.* (2002); Foroumadi *et al.* (2003); He *et al.* (2006); Kritsanida *et al.* (2002); Manfredini *et al.* (2000). For standard bond lengths, see: Allen *et al.* (1987).



Experimental

Crystal data

$\text{C}_{13}\text{H}_{10}\text{N}_4$
 $M_r = 222.25$
Monoclinic, $P2_1/c$
 $a = 5.499(3)$ Å

$b = 10.079(6)$ Å
 $c = 19.942(12)$ Å
 $\beta = 92.758(7)^\circ$
 $V = 1104.0(11)$ Å³

$Z = 4$
 $\text{Mo } K\alpha$ radiation
 $\mu = 0.09$ mm⁻¹

$T = 296$ K
 $0.2 \times 0.15 \times 0.1$ mm

Data collection

Bruker APEXII CCD
diffractometer
Absorption correction: multi-scan
(SADABS; Sheldrick, 1996)
 $T_{\min} = 0.5$, $T_{\max} = 1.0$

6574 measured reflections
2168 independent reflections
1787 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.019$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.042$
 $wR(F^2) = 0.113$
 $S = 1.08$
2168 reflections

154 parameters
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.15$ e Å⁻³
 $\Delta\rho_{\min} = -0.25$ e Å⁻³

Table 1
Hydrogen-bond geometry (Å, °).

$D—H\cdots A$	$D—H$	$H\cdots A$	$D\cdots A$	$D—H\cdots A$
$\text{C9}—\text{H9A}\cdots\text{N1}$	0.93	2.36	2.914 (2)	118
$\text{C13}—\text{H13A}\cdots\text{N4}^i$	0.93	2.45	3.330 (3)	157

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

Data collection: APEX2 (Bruker, 2008); cell refinement: SAINT (Bruker, 2008); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: SHELXTL (Sheldrick, 2008); software used to prepare material for publication: publCIF (Westrip, 2010).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: LH5529).

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supplementary materials

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N-(Naphthalen-1-ylmethylidene)-4H-1,2,4-triazol-4-amine

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Comment

1,2,4-Triazole is a basic aromatic ring and possesses good coordination ability due to the presence of nitrogen atoms. 1,2,4-Triazole derivatives can be used to build polymetallic complexes (He *et al.*, 2006). Compounds derived from triazole possess antimicrobial, analgesic, anti-inflammatory, local anesthetic, antineoplastic and antimalarial properties (Foroumadi *et al.*, 2003). Some triazole Schiff bases also exhibit antiproliferative and anticancer activities (Manfredini *et al.*, 2000). Due to their significant biological applications, triazoles have gained much attention in bioinorganic and metal-based drug discovery (Demirbas *et al.*, 2002; Kritsanida *et al.*, 2002).

The molecular structure of the title compound is shown in Fig. 1. The dihedral angle between the triazole and naphthalene ring system is 56.1 (2)°. The C—N and C=N bond lengths agree with standard values (Allen *et al.*, (1987) and show the obvious effects of electron delocalization. In the crystal, molecules are connected by weak C—H···N hydrogen bonds into chains along [100] (Fig. 2).

Experimental

A mixture of 1-naphthaldehyde (10 mmol) and 4-amino-4H-1,2,4-triazole (10 mmol) in ethanol (20 mL) was refluxed on a steam-bath for 30 min. The colour of the solution changed to reddish-orange and was kept under ice-cold conditions to obtain a white solid product. Single crystals were formed in the mother liquor after ten days.

Refinement

H atoms were positioned geometrically (C—H = 0.93 Å) and allowed to ride on their parent atoms, with $U_{\text{iso}}(\text{H}) = 1.2 U_{\text{eq}}(\text{C})$.

Computing details

Data collection: *APEX2* (Bruker, 2008); cell refinement: *SAINT* (Bruker, 2008); data reduction: *SAINT* (Bruker, 2008); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *publCIF* (Westrip, 2010).

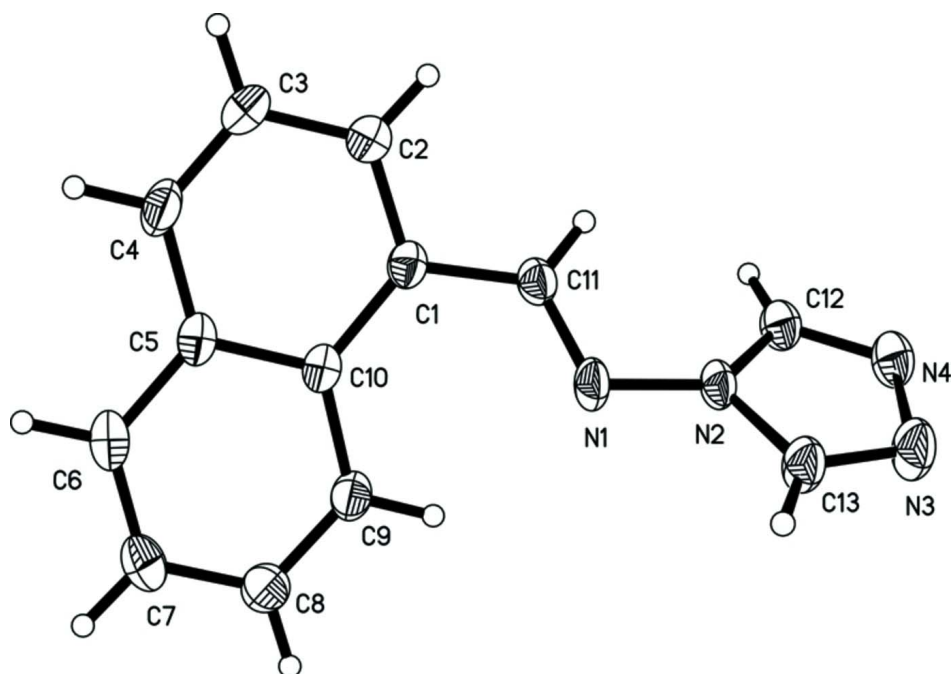


Figure 1

The molecular structure of the title compound. Displacement ellipsoids are drawn at the 30% probability level.

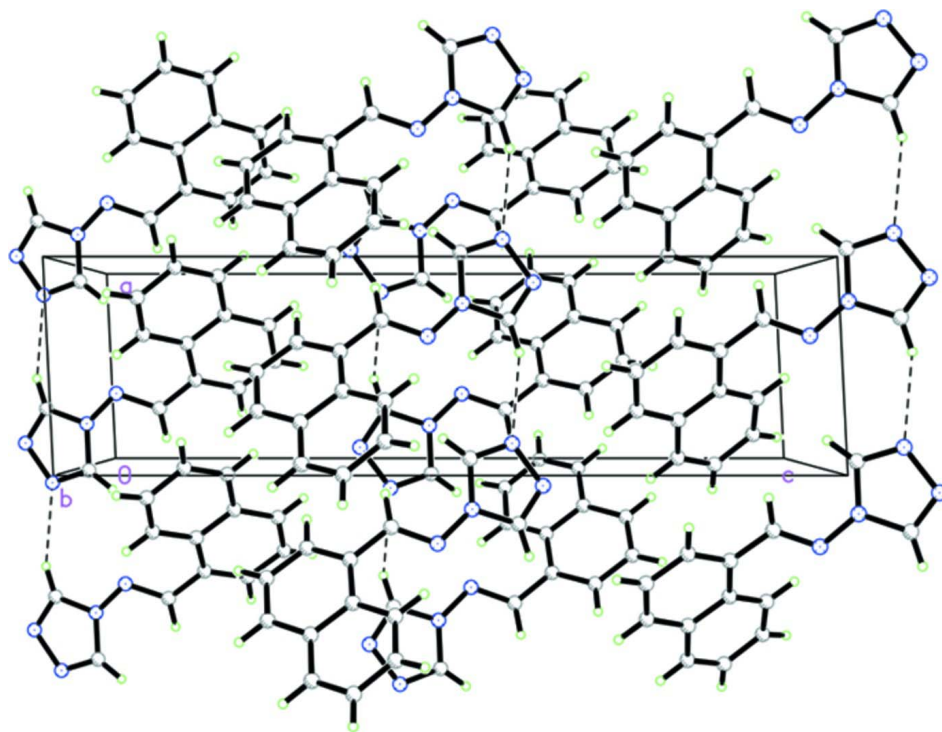


Figure 2

Part of the crystal structure of the title compound, showing weak hydrogen bonds as dashed lines.

N-(Naphthalen-1-ylmethylidene)-4*H*-1,2,4-triazol-4-amine

Crystal data

$C_{13}H_{10}N_4$	$F(000) = 464$
$M_r = 222.25$	$D_x = 1.337 \text{ Mg m}^{-3}$
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
Hall symbol: $-P 2_1bc$	Cell parameters from 6574 reflections
$a = 5.499 (3) \text{ \AA}$	$\theta = 2.0\text{--}26^\circ$
$b = 10.079 (6) \text{ \AA}$	$\mu = 0.09 \text{ mm}^{-1}$
$c = 19.942 (12) \text{ \AA}$	$T = 296 \text{ K}$
$\beta = 92.758 (7)^\circ$	Block, colorless
$V = 1104.0 (11) \text{ \AA}^3$	$0.2 \times 0.15 \times 0.1 \text{ mm}$
$Z = 4$	

Data collection

Bruker APEXII CCD	6574 measured reflections
diffractometer	2168 independent reflections
Radiation source: fine-focus sealed tube	1787 reflections with $I > 2\sigma(I)$
Graphite monochromator	$R_{\text{int}} = 0.019$
φ and ω scans	$\theta_{\text{max}} = 26.0^\circ$, $\theta_{\text{min}} = 2.0^\circ$
Absorption correction: multi-scan	$h = -6 \rightarrow 6$
(<i>SADABS</i> ; Sheldrick, 1996)	$k = -12 \rightarrow 12$
$T_{\text{min}} = 0.5$, $T_{\text{max}} = 1.0$	$l = -24 \rightarrow 24$

Refinement

Refinement on F^2	0 restraints
Least-squares matrix: full	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.042$	$w = 1/[\sigma^2(F_o^2) + (0.0598P)^2 + 0.1522P]$
$wR(F^2) = 0.113$	where $P = (F_o^2 + 2F_c^2)/3$
$S = 1.08$	$(\Delta/\sigma)_{\text{max}} = 0.001$
2168 reflections	$\Delta\rho_{\text{max}} = 0.15 \text{ e \AA}^{-3}$
154 parameters	$\Delta\rho_{\text{min}} = -0.25 \text{ e \AA}^{-3}$

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 ($\text{\AA}^2$)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
N1	0.6439 (2)	0.79790 (12)	0.97839 (6)	0.0429 (3)
N2	0.8048 (2)	0.84041 (11)	1.02686 (6)	0.0384 (3)
N3	0.9117 (2)	0.91095 (15)	1.12480 (6)	0.0570 (4)
N4	1.1212 (2)	0.88352 (14)	1.08514 (6)	0.0517 (3)
C1	0.5720 (2)	0.77204 (14)	0.86143 (6)	0.0382 (3)
C2	0.6174 (3)	0.84224 (16)	0.80284 (7)	0.0494 (4)
H2A	0.7474	0.9012	0.8024	0.059*
C3	0.4736 (3)	0.82456 (17)	0.74739 (8)	0.0595 (5)

H3A	0.5001	0.8713	0.7082	0.071*
C4	0.2871 (3)	0.73591 (17)	0.75027 (7)	0.0544 (4)
H4A	0.1853	0.7253	0.7120	0.065*
C5	0.2370 (3)	0.65767 (13)	0.80837 (7)	0.0400 (3)
C6	0.0469 (3)	0.56297 (15)	0.81066 (8)	0.0489 (4)
H6A	−0.0576	0.5540	0.7729	0.059*
C7	0.0069 (3)	0.48337 (16)	0.86520 (8)	0.0539 (4)
H7A	−0.1223	0.4238	0.8643	0.065*
C8	0.1586 (3)	0.49371 (15)	0.91971 (8)	0.0518 (4)
H8A	0.1405	0.4386	0.9565	0.062*
C9	0.3399 (3)	0.58640 (14)	0.92032 (7)	0.0445 (4)
H9A	0.4410	0.5938	0.9589	0.053*
C10	0.3841 (2)	0.67329 (13)	0.86498 (6)	0.0356 (3)
C11	0.7224 (3)	0.80772 (13)	0.91754 (7)	0.0398 (3)
H11A	0.8792	0.8386	0.9114	0.048*
C12	1.0518 (3)	0.84233 (14)	1.02727 (7)	0.0437 (4)
H12A	1.1494	0.8185	0.9924	0.052*
C13	0.7272 (3)	0.88263 (17)	1.08876 (7)	0.0489 (4)
H13A	0.5666	0.8887	1.1011	0.059*

Atomic displacement parameters ($\text{\AA}^2$)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
N1	0.0343 (6)	0.0541 (7)	0.0390 (6)	−0.0046 (5)	−0.0129 (5)	−0.0057 (5)
N2	0.0317 (6)	0.0439 (6)	0.0385 (6)	−0.0033 (5)	−0.0114 (5)	−0.0025 (5)
N3	0.0428 (8)	0.0794 (10)	0.0471 (7)	−0.0031 (7)	−0.0143 (6)	−0.0101 (7)
N4	0.0383 (7)	0.0617 (8)	0.0534 (7)	−0.0040 (6)	−0.0154 (6)	−0.0039 (6)
C1	0.0372 (7)	0.0403 (7)	0.0362 (7)	0.0026 (6)	−0.0062 (5)	−0.0008 (5)
C2	0.0502 (9)	0.0541 (9)	0.0432 (8)	−0.0083 (7)	−0.0050 (7)	0.0064 (6)
C3	0.0709 (11)	0.0685 (11)	0.0378 (8)	−0.0079 (9)	−0.0108 (7)	0.0141 (7)
C4	0.0592 (10)	0.0630 (10)	0.0389 (8)	0.0006 (8)	−0.0188 (7)	0.0019 (7)
C5	0.0385 (8)	0.0413 (7)	0.0393 (7)	0.0056 (6)	−0.0082 (6)	−0.0065 (6)
C6	0.0434 (8)	0.0501 (9)	0.0516 (8)	0.0021 (7)	−0.0130 (7)	−0.0127 (7)
C7	0.0499 (9)	0.0480 (9)	0.0631 (10)	−0.0109 (7)	−0.0042 (7)	−0.0123 (7)
C8	0.0653 (10)	0.0427 (8)	0.0471 (8)	−0.0082 (7)	−0.0005 (7)	0.0007 (6)
C9	0.0536 (9)	0.0406 (8)	0.0384 (7)	−0.0030 (7)	−0.0088 (6)	−0.0012 (6)
C10	0.0364 (7)	0.0361 (7)	0.0337 (7)	0.0052 (5)	−0.0045 (5)	−0.0046 (5)
C11	0.0350 (7)	0.0406 (7)	0.0429 (8)	−0.0030 (6)	−0.0078 (6)	0.0002 (6)
C12	0.0337 (7)	0.0489 (8)	0.0475 (8)	−0.0005 (6)	−0.0079 (6)	−0.0018 (6)
C13	0.0356 (8)	0.0692 (10)	0.0411 (8)	−0.0031 (7)	−0.0073 (6)	−0.0067 (7)

Geometric parameters ($\text{\AA}$, $^\circ$)

N1—C11	1.311 (2)	C4—H4A	0.9300
N1—N2	1.3486 (16)	C5—C10	1.3661 (19)
N2—C12	1.3577 (19)	C5—C6	1.418 (2)
N2—C13	1.392 (2)	C6—C7	1.378 (2)
N3—C13	1.2481 (19)	C6—H6A	0.9300
N3—N4	1.455 (2)	C7—C8	1.342 (2)
N4—C12	1.2679 (19)	C7—H7A	0.9300

C1—C2	1.399 (2)	C8—C9	1.366 (2)
C1—C11	1.4059 (19)	C8—H8A	0.9300
C1—C10	1.438 (2)	C9—C10	1.439 (2)
C2—C3	1.340 (2)	C9—H9A	0.9300
C2—H2A	0.9300	C11—H11A	0.9300
C3—C4	1.363 (2)	C12—H12A	0.9300
C3—H3A	0.9300	C13—H13A	0.9300
C4—C5	1.439 (2)		
C11—N1—N2	113.90 (12)	C5—C6—H6A	117.9
N1—N2—C12	129.10 (12)	C8—C7—C6	118.53 (15)
N1—N2—C13	120.92 (12)	C8—C7—H7A	120.7
C12—N2—C13	109.85 (11)	C6—C7—H7A	120.7
C13—N3—N4	106.65 (13)	C7—C8—C9	119.02 (15)
C12—N4—N3	110.20 (12)	C7—C8—H8A	120.5
C2—C1—C11	114.43 (13)	C9—C8—H8A	120.5
C2—C1—C10	123.27 (12)	C8—C9—C10	124.04 (13)
C11—C1—C10	122.29 (12)	C8—C9—H9A	118.0
C3—C2—C1	120.01 (15)	C10—C9—H9A	118.0
C3—C2—H2A	120.0	C5—C10—C9	116.64 (13)
C1—C2—H2A	120.0	C5—C10—C1	115.93 (12)
C2—C3—C4	117.87 (15)	C9—C10—C1	127.39 (12)
C2—C3—H3A	121.1	N1—C11—C1	120.65 (13)
C4—C3—H3A	121.1	N1—C11—H11A	119.7
C3—C4—C5	124.52 (13)	C1—C11—H11A	119.7
C3—C4—H4A	117.7	N4—C12—N2	105.52 (13)
C5—C4—H4A	117.7	N4—C12—H12A	127.2
C10—C5—C6	117.34 (13)	N2—C12—H12A	127.2
C10—C5—C4	118.24 (14)	N3—C13—N2	107.76 (14)
C6—C5—C4	124.40 (13)	N3—C13—H13A	126.1
C7—C6—C5	124.29 (13)	N2—C13—H13A	126.1
C7—C6—H6A	117.9		

Hydrogen-bond geometry (Å, °)

<i>D</i> —H $\cdots$ <i>A</i>	<i>D</i> —H	H $\cdots$ <i>A</i>	<i>D</i> $\cdots$ <i>A</i>	<i>D</i> —H $\cdots$ <i>A</i>
C9—H9A $\cdots$ N1	0.93	2.36	2.914 (2)	118
C13—H13A $\cdots$ N4 ⁱ	0.93	2.45	3.330 (3)	157

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