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# 4-Amino-3,5-diethyl-4H-1,2,4-triazole at 100 K: chains of edge-fused $R_4^4(10)$ and $R_4^4(20)$ rings

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The title compound,  $C_6H_{12}N_4$ , has four crystallographically independent molecules in the asymmetric unit. Intermolecular  $N-H\cdots N$  hydrogen bonds involving amino groups and triazole N atoms form a three-dimensional framework involving  $R_4^4(10)$  and  $R_4^4(20)$  rings. The hydrogen bonding is supported by weak  $C-H\cdots \pi$  interactions.

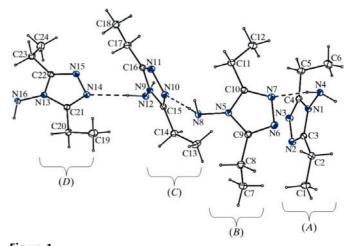
## Comment

1,2,4-Triazole and its derivatives have been used as starting materials for the synthesis of many heterocycles (Desenko, 1995). The triazole ring, having strong  $\sigma$ -donor and weak  $\pi$ -acceptor properties, potentially has two different coordination modes through three N donor atoms coordinating to metal ions (Van Diemen et al., 1991; Ding et al., 2004; Yi et al., 2004; Ren et al., 2006). Recent interest in substituted 1,2,4triazoles has arisen in part from their transition metal complexes with intriguing structures and specific magnetic properties (Zhou et al., 2005, 2006). Many metal complexes containing substituted 1,2,4-triazole have potential applications in molecular-based memory devices, displays and optical switches owing to their spin crossover properties (Garcia et al., 1997; Kahn & Martinez, 1998). Apart from their chemical significance, 1,2,4-triazole derivatives have been found to be associated with diverse pharmacological properties, such as anti-inflammatory, antifungal and antiviral activities (Massa et al., 1992; Mahomed et al., 1993; Mullican et al., 1993). Some are also known to exhibit analgesic, anticonvulsant, tranquilizing, antidepressant, anxiolytic (Bradbury & Rivett, 1991; Sughen & Yoloye, 1978; Stillings et al., 1986; Kane et al., 1988) or even antitumour activities (Hatheway et al., 1978) and are applied in therapy (e.g. alprazolam, estazolam, triazolam and adinazolam; Budavari et al., 1996). In spite of the chemical and medicinal importance of this class of compounds, relatively few crystal structures of 1,2,4-triazole derivatives have been reported (Cambridge Structural Database, Version 5.27 of

November, 2005; Allen, 2002). In order to clarify the structure of this type of compound, an X-ray structure determination of the title compound, (I), has been carried out, and the results are presented here. The structure has been confirmed by IR, <sup>1</sup>H NMR and <sup>13</sup>C NMR spectroscopies and also by elemental analysis.

The molecular structure and atom-numbering scheme for (I) are shown in Fig. 1; selected bond lengths are given in Table 1. Compound (I) crystallizes in the space group C2/c with Z' = 4, and the hydrogen bonding was analysed with the aid of PLATON (Spek, 2003). The asymmetric unit contains four independent molecules with statistically equivalent metrical parameters but different conformations. The N1-N4, N5-N8, N9-N12 and N13-N16 bond lengths (Table 1) indicate single-bond character, whereas the N2-N3, N6-N7, N10-N11 and N14-N15 bond lengths are indicative of significant double-bond character. Similar N-N and N=N bond-length values have been observed in 4-amino-3-methyl-5-(p-tolyl)-4H-1,2,4-triazole and 4-amino-3-methyl-5-phenyl-4H-1,2,4-triazole [N-N = 1.4090 (16) and 1.4081 (18) Å, and N=N = 1.3859 (19) and 1.396 (2) Å; Şahin *et al.*, 2006]. The H atoms of the amino group form hydrogen bonds with the N atoms of neighbouring triazole rings. The geometric parameters of the N-H···N hydrogen-bonding interactions are given in Table 2.

Amino atom N4 in the reference molecule at (x, y, z) acts as a hydrogen-bond donor, via H4B, to atom N7 within the asymmetric unit and, via H4A, to atom N6 $^{\rm v}$  (symmetry codes are defined in the footnote of Table 2), so forming a centrosymmetric  $R_4^4(10)(A)$  (Bernstein *et al.*, 1995) ring centred at  $(\frac{1}{4}, \frac{3}{4}, \frac{1}{2})$ . Similarly, amino atom N12 acts as a hydrogen-bond



**Figure 1**The molecular structure of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 40% probability level.

## organic compounds

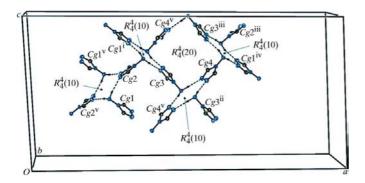


Figure 2 The packing of (I), showing the  $R_4^4(10)$  and  $R_4^4(20)$  ring patterns. Dashed lines indicate hydrogen bonds. H atoms not involved in these interactions and ethyl groups have been omitted for clarity. (Symmetry codes are provided in Table 2.)

donor, via H12D, to atom N14 within the asymmetric unit and, via H12E, to atom N15<sup>ii</sup>, so forming a second centrosymmetric ring motif, this time of  $R_4^4(10)(B)$  type, centred at  $(\frac{1}{2}, 0, \frac{1}{2})$ . The arrangement of the N8-H8 $C \cdot \cdot$ N10, N8-H8 $D \cdot \cdot \cdot$ N3<sup>i</sup>, N16<sup>iii</sup>-H16 $A^{iii} \cdot \cdot \cdot$ N11 and N16<sup>iii</sup>-H16 $B^{iii} \cdot \cdot \cdot$ N2<sup>i</sup> interactions can be described by the graph-set notation  $R_4^4(10)(C)$ . At the same time, the N16-H16 $A \cdot \cdot \cdot$ N11<sup>iii</sup>, N16-H16 $B \cdot \cdot \cdot \cdot$ N2<sup>iv</sup>, N8<sup>iii</sup>-H8 $C^{iii} \cdot \cdot \cdot$ N10<sup>iii</sup> and N8<sup>iii</sup>-H8 $D^{iii} \cdot \cdot \cdot$ N3<sup>iv</sup> interactions constitute an  $R_4^4(10)(D)$  ring. Finally, the N12-H12 $D \cdot \cdot \cdot \cdot$ N14, N16-H16 $A \cdot \cdot \cdot \cdot$ N11<sup>iii</sup>, N12<sup>iii</sup>-H12 $D^{iii} \cdot \cdot \cdot \cdot$ N14 interactions produce an  $R_4^4(20)(E)$  ring (Fig. 2).

Propagation of eight hydrogen bonds then forms a chain of edge-fused rings, containing  $R_4^4(10)(A)R_4^4(10)(B)R_4^4(10)(C)$ - $R_4^4(10)(D)$  sequences of four edge-fused rings. Similarly, edge-fused  $R_4^4(10)(C)$  and  $R_4^4(20)(E)$  rings form a chain running along the c axis. In compound (I), interlinked  $C_4^4(20)$  antiparallel chains zigzagging along the a axis are formed through  $N4^v-H4A^v\cdots N6$ ,  $N8-H8C\cdots N10$ ,  $N12-H12D\cdots N14$  and  $N16-H16B\cdots N2^{iv}$  interactions. Amino atom N16 in the molecule at  $(x,-y,-\frac{1}{2}+z)$  acts as hydrogen-bond donor, via H16A, to  $N11^{ii}$ , while  $N12^{ii}$  acts as donor to N15, and in this manner a  $C_2^2(10)$  chain running parallel to the [001] direction is generated.

These intermolecular interactions, namely an extensive network of hydrogen bonds and  $\pi$ -ring interactions, are responsible for constructing an infinite three-dimensional framework.

## **Experimental**

Propionic acid (18.5 g, 0.25 mol) was added to a solution of hydrazine hydrate (21.5 g, 0.4 mol) and the mixture was refluxed for 5 h. On cooling, a precipitate was formed, and this product was filtered off and dried. Recrystallization from ethyl acetate gave a colourless product (yield 62%). Single crystals of (I) were obtained from ethyl acetate at room temperature by slow evaporation (m.p. 438–439 K). IR (KBr, cm<sup>-1</sup>): 3235–3120 (v<sub>NH2</sub>), 1664 (v<sub>C=N</sub>);  $^{1}$ H NMR (CDCl<sub>3</sub>):  $\delta$  1.30 (t, 6H, 2CH<sub>3</sub>), 2.76 (g, 4H, 2CH<sub>2</sub>), 5.02 (g, 2H, NH<sub>2</sub>);  $^{13}$ C NMR (CDCl<sub>3</sub>):  $\delta$  156.21 (triazole C<sub>3</sub> and triazole C<sub>5</sub>), 17.87 (CH<sub>2</sub>), 11.40 (CH<sub>3</sub>). Elemental analysis calculated for C<sub>6</sub>H<sub>12</sub>N<sub>4</sub>: C 51.41, H 8.63, N 39.97%; found: C 52.40, H 8.61, N 39.75%.

Cr	etal	data
$c_{I}$	vsiai	аши

$C_6H_{12}N_4$	$V = 6457.6 (7) \text{ Å}^3$
$M_r = 140.20$	Z = 32
Monoclinic, C2/c	Mo $K\alpha$ radiation
a = 37.782 (2) Å	$\mu = 0.08 \text{ mm}^{-1}$
b = 9.2996 (4) Å	T = 100  K
c = 18.4055 (12)  Å	$0.50 \times 0.48 \times 0.41 \text{ mm}$
$\beta = 93.067 (5)^{\circ}$	

#### Data collection

Stoe IPDSII diffractometer 4902 reflections with  $I>2\sigma(I)$  6583 measured reflections  $R_{\rm int}=0.027$ 

### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.035$	H atoms treated by a mixture of
$wR(F^2) = 0.091$	independent and constrained
S = 1.04	refinement
6331 reflections	$\Delta \rho_{\text{max}} = 0.16 \text{ e Å}^{-3}$
393 parameters	$\Delta \rho_{\min} = -0.23 \text{ e Å}^{-3}$

Table 1 Selected bond lengths (Å).

N1-N4	1.4130 (14)	N9-N12	1.4109 (14)
N2-N3	1.3997 (15)	N10-N11	1.4016 (14)
N5-N8	1.4118 (14)	N13-N16	1.4119 (14)
N6-N7	1.3981 (15)	N14-N15	1.3984 (15)

**Table 2** Hydrogen-bond geometry (Å, °).

$D-\mathbf{H}\cdot\cdot\cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D - H \cdot \cdot \cdot A$
$\begin{array}{c} B-11 \cdot VA \\ N8-H8C \cdots N10 \\ N8-H8D \cdots N3^{i} \\ N12-H12D \cdots N14 \\ N12-H12E \cdots N15^{ii} \\ N16-H16A \cdots N11^{iii} \\ N16-H16B \cdots N2^{iv} \\ N4-H4A \cdots N6^{v} \end{array}$	0.910 (17) 0.923 (19) 0.929 (16) 0.952 (16) 0.931 (17) 0.942 (17) 0.888 (16)	2.192 (17) 2.088 (19) 2.108 (16) 2.127 (17) 2.112 (18) 2.070 (17) 2.195 (17)	3.0975 (15) 2.9645 (16) 3.0295 (15) 3.0650 (16) 3.0256 (16) 3.0066 (15) 3.0803 (15)	173.4 (14) 158.2 (15) 171.6 (14) 168.0 (13) 166.7 (14) 172.9 (14) 174.8 (14)
$N4-H4B\cdots N7$	0.925 (18)	2.161 (18)	3.0736 (16)	168.8 (14)

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

All H atoms bound to carbon were refined using a riding model, with C—H distances of 0.97 Å [ $U_{\rm iso}({\rm H})=1.2U_{\rm eq}({\rm parent\ atom})$ ] for methylene H atoms and 0.96 Å [ $U_{\rm iso}({\rm H})=1.5U_{\rm eq}({\rm parent\ atom})$ ] for methyl H atoms. The amino H atoms were located in a difference map and were refined freely (distances are given in Table 2).

Data collection: *X-AREA* (Stoe & Cie, 2002); cell refinement: *X-AREA*; data reduction: *X-RED* (Stoe & Cie, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: GZ3081). Services for accessing these data are described at the back of the journal.

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