

## 4-(2-Hydroxyethyl)-4*H*-1,2,4-triazole, an intermediate in the synthesis of iron–triazole spin-crossover compounds

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### Key indicators

Single-crystal X-ray study  
T = 150 K  
Mean  $\sigma(C-C) = 0.002 \text{ \AA}$   
R factor = 0.041  
wR factor = 0.100  
Data-to-parameter ratio = 12.9

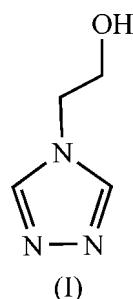
For details of how these key indicators were automatically derived from the article, see  
<http://journals.iucr.org/e>.

The title 1,2,4-triazole derivative,  $C_4H_7N_3O$ , has been synthesized and characterized. Neighboring molecules are connected by intermolecular O–H···N hydrogen bonds, forming a linear chain arrangement along the crystallographic [101] direction.

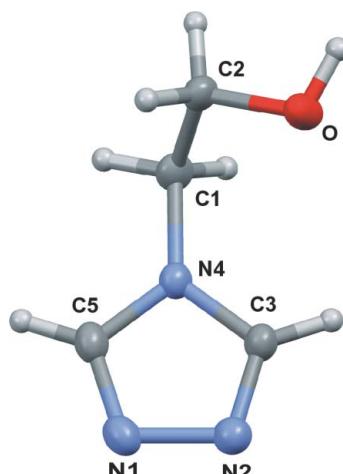
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### Comment

The title compound, (hyetrz, I), has been used to react with divalent Fe or Cu complexes in order to investigate the interesting spin-crossover phenomena of  $Fe^{2+}$  compounds (Garcia, van Koningsbruggen, Bravic *et al.*, 1997; Garcia *et al.*, 2000, 2002) and to establish the structural model based on  $Cu^{2+}$  analogs (Garcia, van Koningsbruggen, Codjovi *et al.*, 1997; Garcia *et al.*, 2003). Although three metal complexes of the ligand have been structurally determined, the crystal structure of the free ligand itself is yet unknown. Here we report its structure.



The hydroxyethyl group represents the flexible part of the molecule (Fig. 1), and bends out of the plane of the triazole



**Figure 1**  
Displacement ellipsoid plot (50% probability level) for (I).

ring. The resulting torsion angle N4—C1—C2—O is 61.3 (2)°. This value is in the range of those observed in the metal coordination complexes: 48.2–64.8° in [Cu(hyetrz)<sub>3</sub>]·(ClO<sub>4</sub>)<sub>2</sub>·3H<sub>2</sub>O (Garcia, van Koningsbruggen, Bravic *et al.*, 1997), 52.9–66.3° in [Cu(hyetrz)<sub>3</sub>]·(CF<sub>3</sub>SO<sub>3</sub>)<sub>2</sub>·H<sub>2</sub>O (Garcia *et al.*, 2003), and 68.0 and 68.3° in [Fe<sub>3</sub>(hyetrz)<sub>6</sub>]·(H<sub>2</sub>O)<sub>6</sub>]·(CF<sub>3</sub>SO<sub>3</sub>)<sub>6</sub> (Garcia *et al.*, 2000). Obviously it is affected by the intermolecular interactions. The bond lengths and angles are almost identical to those in the above metal complexes.

In the crystal structure of (I), the molecules are connected by hydrogen bonds between N atoms of the triazole rings and H atoms of the hydroxy groups in neighboring molecules (Table 2), leading to a linear chain arrangement along the [101] direction (Fig. 2).

## Experimental

The compound was prepared from monoformyl hydrazine, triethyl orthoformate and 2-ethanolamine according to the general method described by Bayer *et al.* (1974). m.p.: 357 K by DTA. <sup>1</sup>H NMR (250 MHz) (DMSO): 3.66 (*t*, 2H, CH<sub>2</sub>), 4.09 (*t*, 2H, CH<sub>2</sub>), 5.10 (*s*, 1H, OH), 8.48 (*s*, 2H, CH). Recrystallization from methanol results in single crystals suitable for X-ray measurement.

### Crystal data

C <sub>4</sub> H <sub>7</sub> N <sub>3</sub> O	$D_x = 1.400 \text{ Mg m}^{-3}$
$M_r = 113.13$	Mo $\text{K}\alpha$ radiation
Monoclinic, $P2_1/c$	Cell parameters from 39 reflections
$a = 5.5894 (18) \text{ \AA}$	$\theta = 2.8\text{--}28.0^\circ$
$b = 14.444 (5) \text{ \AA}$	$\mu = 0.11 \text{ mm}^{-1}$
$c = 7.013 (4) \text{ \AA}$	$T = 150 (2) \text{ K}$
$\beta = 108.56 (3)^\circ$	Block, colourless
$V = 536.7 (4) \text{ \AA}^3$	$0.23 \times 0.18 \times 0.18 \text{ mm}$
$Z = 4$	

### Data collection

Siemens SMART 1K CCD area-detector diffractometer	976 reflections with $I > 2\sigma(I)$
$\omega$ scans	$R_{\text{int}} = 0.095$
Absorption correction: none	$\theta_{\text{max}} = 28.0^\circ$
8131 measured reflections	$h = -7 \rightarrow 7$
1301 independent reflections	$k = -19 \rightarrow 18$
All H-atom parameters refined	$l = -9 \rightarrow 9$

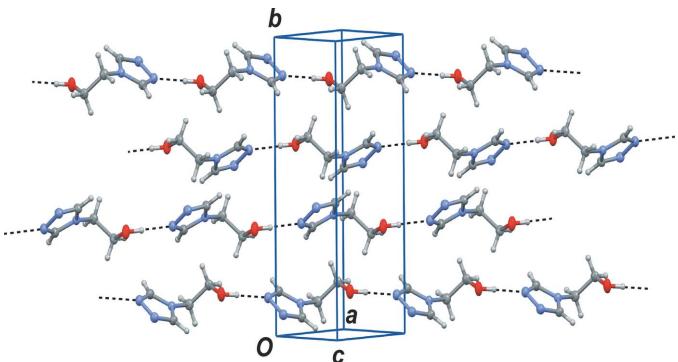
### Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F_o^2) + (0.0324P)^2 + 0.1227P]$
$R[F^2 > 2\sigma(F^2)] = 0.041$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.100$	$(\Delta/\sigma)_{\text{max}} < 0.001$
$S = 1.08$	$\Delta\rho_{\text{max}} = 0.20 \text{ e \AA}^{-3}$
1301 reflections	$\Delta\rho_{\text{min}} = -0.25 \text{ e \AA}^{-3}$
101 parameters	
All H-atom parameters refined	

**Table 1**

Selected bond length.

N1—N2	1.3923 (17)	N1—C5	1.3031 (19)
N2—C3	1.308 (2)	N4—C1	1.466 (2)
C3—N4	1.3505 (19)	C1—C2	1.509 (2)
N4—C5	1.3594 (17)	C2—O	1.4197 (17)



**Figure 2**

MERCURY (Bruno *et al.*, 2002) representation of the intermolecular hydrogen-bond (dashed lines) arrangement (C, grey; H, light gray; N, blue; O, red).

**Table 2**

Hydrogen-bond geometry (Å, °).

$D—H \cdots A$	$D—H$	$H \cdots A$	$D \cdots A$	$D—H \cdots A$
O—H1···N2 <sup>i</sup>	0.90 (2)	1.85 (2)	2.754 (2)	178 (2)

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

All H atoms were located in difference Fourier syntheses and were refined freely with isotropic displacement parameters.

Data collection: SMART (Siemens, 1995); cell refinement: SMART; data reduction: SAINT (Siemens, 1995); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: MERCURY (Bruno *et al.*, 2002); software used to prepare material for publication: SHELXL97.

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