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Synthesis and crystal structure of 1*H*-1,2,4-triazole-3,5-diamine monohydrate

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The title compound, a hydrate of 3,5-diamino-1,2,4-triazole (DATA), $C_2H_5N_5$ · H_2O , was synthesized in the presence of sodium perchlorate. The evaporation of H_2O from its aqueous solution resulted in anhydrous DATA, suggesting that sodium perchlorate was required to precipitate the DATA hydrate. The DATA hydrate crystallizes in the $P2_1/c$ space group in the form of needle-shaped crystals with one DATA and one water molecule in the asymmetric unit. The water molecules form a three-dimensional network in the crystal structure. Hirshfeld surface analysis revealed that 8.5% of the intermolecular interactions originate from H···O contacts derived from the incorporation of the water molecules.

1. Chemical context

Researchers have focused on the development of less sensitive and highly energetic materials. The sensitivity of energetic materials is related to their crystal structures and intermolecular interactions (Kuklja & Rashkeev, 2007). Additionally, the density, which strongly influences the detonation performance, can be calculated from the crystal structure. Hence, the determination of the crystal structure can elucidate the characteristics of energetic materials.

Azole derivatives have been recognized as promising frameworks for energetic materials because of their high heats of formation (Fisher *et al.*, 2012; Kumasaki *et al.*, 2021; Inoue *et al.*, 2022*a*). Tetrazoles, triazoles, and imidazoles are N-rich heterocyclic azole derivatives. Several energetic materials, including organic explosives, energetic salts, and co-crystals, have been synthesized using azole compounds (Kumasaki *et al.*, 2011; Mori *et al.*, 2021; Inoue *et al.*, 2022*b*).



In our previous study, 1*H*-tetrazole was co-crystallized with NaClO₄; NaClO₄ is an oxidizer that is sometimes used in pyrotechnics (Inoue *et al.*, 2022*b*). The co-crystal exhibited high sensitivity, which was comparable to that of typical primary explosives (Inoue *et al.*, 2022*a*). Subsequently, 3,5-diamino-1,2,4-triazole (guanazole, DATA) was selected as the target material for co-crystallization with NaClO₄. However, our attempt to prepare a co-crystal of DATA and NaClO₄ resulted instead in crystals of DATA hydrate, which has not been previously been reported. Although DATA is used as a





 Table 1

 Bond lengths (Å) in hydrated and non-hydrated DATA.

Bond	DATA	DATA hydrate
C7-N2	1.3544 (16)	1.3670 (15)
N2-C8	1.3339 (16)	1.3400 (15)
C8-N4	1.3356 (16)	1.3337 (15)
N4-N3	1.3951 (15)	1.3943 (13)
N3-C7	1.3238 (16)	1.3185 (15)
C7-N6	1.3747 (17)	1.3797 (15)
C8-N5	1.3502 (17)	1.3547 (15)

raw material in the synthesis of various energetic compounds (Khan *et al.*, 2024; Zhang *et al.*, 2010; Yin *et al.*, 2015), understanding its hydration is valuable for its treatment.

2. Structural commentary

DATA hydrate (Fig. 1) crystallizes in the $P2_1/c$ space group with one DATA molecule and one water molecule in the asymmetric unit. The two N atoms of the amino groups are not coplanar with the mean plane of the ring structure, the distances between the mean plane of the ring structure and N5 and N6 being 0.0719 (19) and 0.1038 (19) Å, respectively. The N4–N3 bond [1.3943 (13) Å] is longer than all of the C–N bonds [the range is 1.3797 (15) for C7–N6 to 1.3185 (15) Å for N3–C7]. Furthermore, the N2–C8 double bond [1.3400 (15) Å] is longer than C8–N4 [1.3337 (15) Å], which is a single bond. Table 1 compares the bond lengths of hydrated and non-hydrated DATA (Klapötke *et al.*, 2010). Evidently, the N2–C8 and N2–C7 bonds in DATA hydrate are longer than those of non-hydrated DATA. The differences between all of the bond lengths are statistically significant.

3. Supramolecular features

In the crystal, the DATA and H_2O molecules form a layered structure (Fig. 2), with layers parallel to the (102) plane and an



Figure 1

Displacement ellipsoid plot (probability level of 50%) of DATA hydrate showing the atom-numbering scheme. The hydrogen bond is represented by a dashed blue line.

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

, , ,				
$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$N5-H5A\cdotsO1^{i}$	0.895 (17)	2.080 (17)	2.9173 (14)	155.4 (14)
$O1 - H1A \cdots N3^{n}$ $O1 - H1B \cdots N2$	0.8/1 (19) 0.90 (2)	1.988 (19) 1.95 (2)	2.8558 (13) 2.8410 (13)	174.1 (16) 171.5 (18)
N4−H4···N3 ⁱⁱⁱ	0.883 (17)	2.306 (16)	2.9947 (14)	134.9 (13)
Symmetry codes: -x+2, -y+1, -z+1	(i) $x, -y + \frac{3}{2}$	$\frac{1}{2}, z + \frac{1}{2};$ (ii)	$-x+1, y+\frac{1}{2}, -$	$z + \frac{1}{2};$ (iii)

interlayer distance of 3.26969 (4) Å. The O1-H1A···N3 and O1···H5A-N5 hydrogen bonds form the layers while the O1-H1B···N2 and N4-H4···N3 hydrogen bonds connect adjacent layers (Table 2, Fig. 3). The water molecules produce a 3D network within the crystal. A water molecule forms two hydrogen bonds with two DATA molecules from the same layer [O1-H1A···N3(1 - x, $\frac{1}{2}$ + y, $\frac{1}{2}$ - z) and O1···H5A-N5(x, $\frac{1}{2}$ - y, $\frac{1}{2}$ + z)] and one with that from an





(a) The crystal structure viewed along the *b* axis. The intra- and interlayer hydrogen bonds are indicated by blue and red lines, respectively, and the layer structure is shown in blue. (b) The intra- and interlayer hydrogen bonds with atom numbers.



Figure 3

Hydrogen bonding in DATA hydrate viewed along the *a* axis. Symmetry codes: (i) $x, \frac{1}{2} - y, \frac{1}{2} + z$; (ii) $1 - x, \frac{1}{2} + y, \frac{1}{2} - z$; (iii) -x, -y, -z.



Figure 4 Fingerprint plots for (*a*) DATA hydrate and (*b*) anhydrous DATA.

adjacent layer (O1-H1B···N2). Atoms N5 of the amino group and N2 from the ring form hydrogen bonds with each water molecule; however, N6 is not involved in hydrogen bonding. Two DATA molecules in two adjacent layers are mutually connected by two N4-H4···N3(-x, -y, -z) hydrogen bonds.

The supramolecular interactions in DATA hydrate were further investigated through Hirshfeld surface analysis using *Crystal Explorer 21* (Spackman *et al.*, 2021). Fig. 4 shows the fingerprint plots for a molecule of DATA non-hydrate (Klapötke *et al.*, 2010) and DATA hydrate. The third spike concerning the H···O interaction was observed upon hydration, whereas the DATA non-hydrate exhibited two spikes of N···H and H···N. The dominant interaction of DATA nonhydrate was N···H/H···N of 53.0%, which decreased to 39.3% with the incorporation of water molecules, and the H···O interaction contributes 8.5% to the crystal packing. The contribution of H···H interactions in DATA hydrate is 37.9%, which is higher than that of DATA non-hydrate (34.2%).

4. Database survey

Previous studies on DATA were explored in the Cambridge Structural Database (CSD, June 2024; Groom *et al.*, 2016). The search resulted in four reports: DAMTRZ11 (Klapötke *et al.*, 2010), DAMTRZ22 (Ivanova & Spiteller, 2017), DAMTRZ10 (Starova *et al.*, 1979) and DAMTRZ20 (Starova *et al.*, 1980). The title DATA hydrate forms a layered structure in a monoclinic space group, whereas anhydrous DATA was reported to form a herringbone structure.

5. Synthesis and crystallization

DATA was purchased from Tokyo Chemical Industry Co., Ltd. Sodium perchlorate was obtained from Kanto Chemical Co., Inc. DATA (1 mmol) and sodium perchlorate (1 mmol) were dissolved in deionized water, and the solvent was removed in a

Table 3	
Experimental	details.

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Crystal data	
Chemical formula	$C_2H_5N_5 \cdot H_2O$
Mr	117.13
Crystal system, space group	Monoclinic, $P2_1/c$
Temperature (K)	123
a, b, c (Å)	3.80560 (5), 9.49424 (11), 14.01599 (15)
β (°)	92.9639 (11)
$V(A^3)$	505.74 (1)
Z	4
Radiation type	Cu Ka
$\mu (\text{mm}^{-1})$	1.07
Crystal size (mm)	$0.2 \times 0.1 \times 0.1$
Data collection	
Diffractometer	XtaLAB AFC12 (RINC): Kappa dual home/near
Absorption correction	Multi-scan (<i>CrysAlis PRO</i> ; Rigaku OD, 2019)
T_{\min}, T_{\max}	0.883, 1.000
No. of measured, independent and observed $[I > 2\sigma(I)]$ reflections	2894, 989, 956
R _{int}	0.019
$(\sin \theta / \lambda)_{\max} (\text{\AA}^{-1})$	0.619
Refinement	
$R[F^2 > 2\sigma(F^2)], wR(F^2), S$	0.031, 0.081, 1.11
No. of reflections	989
No. of parameters	102
H-atom treatment	All H-atom parameters refined
$\Delta \rho_{\rm max}, \Delta \rho_{\rm min} \ ({\rm e} \ {\rm \AA}^{-3})$	0.19, -0.22

Computer programs: CrysAlis PRO (Rigaku OD, 2019), SHELXT2018/2 (Sheldrick, 2015a), SHELXL2019/3 (Sheldrick, 2015b), Mercury (Macrae et al., 2020), publCIF (Westrip, 2010) and OLEX2 (Dolomanov et al., 2009).

silica gel desiccator. After one week, needle-shaped crystals were precipitated (yield: 40.81%). Interestingly, the evaporation of water from an aqueous solution of DATA generated block-shaped non-hydrated DATA crystals. Therefore, sodium perchlorate was required to precipitate the DATA hydrate. The mass proportion of H₂O in the crystals was measured using thermogravimetry. A Thermoplus TG8120 (Rigaku) was used with an Al₂O₃ open cell. The heating rate was set to 10 K min⁻¹. Flow gas was not used to prevent dehydration under the dried flow gas. The measured mass proportion of H₂O in the crystal was slightly lower than the theoretical mass content of H₂O (15.38%).

6. Dehydration behavior

After the storage of DATA hydrate over one night at 33% RH (saturated salt method (Greenspan, 1977); MgCl, 295 K), the water in the hydrate was removed and DATA hydrate turned into DATA. In contrast, after the storage at room temperature (approximately 295 K) and 60% RH for one night, the crystals remained as hydrates.

7. Refinement

The crystal data, data collection, and structural refinement details are summarized in Table 3. The H atoms were identified using difference-Fourier maps and all H-atom parameters were refined.

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Synthesis and crystal structure of 1H-1,2,4-triazole-3,5-diamine monohydrate

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Computing details

1H-1,2,4-Triazole-3,5-diamine monohydrate

Crystal data

C₂H₅N₅·H₂O $M_r = 117.13$ Monoclinic, $P2_1/c$ a = 3.80560 (5) Å b = 9.49424 (11) Å c = 14.01599 (15) Å $\beta = 92.9639$ (11)° V = 505.74 (1) Å³ Z = 4

Data collection

XtaLAB AFC12 (RINC): Kappa dual home/near diffractometer
Radiation source: micro-focus sealed X-ray tube, Rigaku (Cu) X-ray Source
Mirror monochromator
Detector resolution: 5.8140 pixels mm⁻¹ ω scans
Absorption correction: multi-scan (CrysAlisPro; Rigaku OD, 2019)

Refinement

Refinement on F^2 Least-squares matrix: full $R[F^2 > 2\sigma(F^2)] = 0.031$ $wR(F^2) = 0.081$ S = 1.11989 reflections 102 parameters 0 restraints Primary atom site location: dual Hydrogen site location: difference Fourier map F(000) = 248 $D_x = 1.538 \text{ Mg m}^{-3}$ Cu Ka radiation, $\lambda = 1.54184 \text{ Å}$ Cell parameters from 2506 reflections $\theta = 6.3-72.5^{\circ}$ $\mu = 1.07 \text{ mm}^{-1}$ T = 123 KBlock, clear light colourless $0.2 \times 0.1 \times 0.1 \text{ mm}$

 $T_{\min} = 0.883, T_{\max} = 1.000$ 2894 measured reflections 989 independent reflections 956 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.019$ $\theta_{\max} = 72.7^{\circ}, \theta_{\min} = 5.6^{\circ}$ $h = -4 \rightarrow 4$ $k = -6 \rightarrow 11$ $l = -17 \rightarrow 16$

All H-atom parameters refined $w = 1/[\sigma^2(F_o^2) + (0.0403P)^2 + 0.1848P]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{max} < 0.001$ $\Delta\rho_{max} = 0.19 \text{ e } \text{Å}^{-3}$ $\Delta\rho_{min} = -0.22 \text{ e } \text{Å}^{-3}$ Extinction correction: *SHELXL2019/2* (Sheldrick, 2015b), Fc*=kFc[1+0.001xFc^2\lambda^3/sin(2\theta)]^{-1/4} Extinction coefficient: 0.0108 (15)

Special details

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

	x	У	Ζ	$U_{ m iso}$ */ $U_{ m eq}$	
01	0.3849 (2)	0.75354 (9)	0.11841 (6)	0.0209 (3)	
N2	0.7285 (2)	0.72150 (10)	0.30194 (7)	0.0148 (3)	
N4	0.7684 (2)	0.62922 (10)	0.44551 (7)	0.0157 (3)	
N3	0.9151 (3)	0.52932 (10)	0.38648 (7)	0.0164 (3)	
N6	0.9638 (3)	0.52599 (11)	0.21830 (7)	0.0172 (3)	
N5	0.5282 (3)	0.86004 (11)	0.43097 (8)	0.0186 (3)	
C8	0.6633 (3)	0.74110 (12)	0.39403 (8)	0.0145 (3)	
C7	0.8788 (3)	0.59072 (12)	0.30217 (8)	0.0141 (3)	
H4	0.765 (4)	0.6124 (16)	0.5074 (12)	0.021 (4)*	
H5A	0.454 (4)	0.8506 (17)	0.4902 (12)	0.023 (4)*	
H6A	1.064 (4)	0.5847 (19)	0.1783 (12)	0.032 (4)*	
H6B	1.085 (4)	0.448 (2)	0.2284 (12)	0.031 (4)*	
H5B	0.402 (4)	0.9142 (19)	0.3880 (12)	0.032 (4)*	
H1A	0.280 (5)	0.835 (2)	0.1149 (13)	0.041 (5)*	
H1B	0.495 (5)	0.753 (2)	0.1769 (16)	0.045 (5)*	

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

Atomic displacement parameters $(Å^2)$

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
01	0.0267 (5)	0.0192 (5)	0.0162 (5)	0.0054 (4)	-0.0036 (4)	-0.0011 (3)
N2	0.0157 (5)	0.0133 (5)	0.0150 (5)	0.0000 (4)	-0.0010 (4)	0.0002 (4)
N4	0.0199 (5)	0.0147 (5)	0.0123 (5)	0.0009 (4)	-0.0014 (4)	0.0003 (4)
N3	0.0181 (5)	0.0142 (5)	0.0165 (5)	0.0011 (4)	-0.0018 (4)	-0.0009(4)
N6	0.0211 (5)	0.0136 (5)	0.0169 (5)	0.0012 (4)	0.0003 (4)	-0.0005 (4)
N5	0.0239 (5)	0.0164 (5)	0.0156 (5)	0.0037 (4)	0.0011 (4)	0.0001 (4)
C8	0.0130 (5)	0.0144 (6)	0.0157 (5)	-0.0021 (4)	-0.0022 (4)	0.0002 (4)
C7	0.0128 (5)	0.0128 (5)	0.0165 (6)	-0.0018 (4)	-0.0018 (4)	-0.0001 (4)
N5 C8 C7	0.0239 (5) 0.0130 (5) 0.0128 (5)	0.0164 (5) 0.0144 (6) 0.0128 (5)	0.0156 (5) 0.0157 (5) 0.0165 (6)	0.0037 (4) -0.0021 (4) -0.0018 (4)	0.0011 (4) -0.0022 (4) -0.0018 (4)	0.0001 (4) 0.0002 (4) -0.0001 (

Geometric parameters (Å, °)

O1—H1A	0.87 (2)	N3—C7	1.3185 (15)	
O1—H1B	0.90 (2)	N6—C7	1.3797 (15)	
N2—C8	1.3400 (15)	N6—H6A	0.891 (19)	
N2—C7	1.3670 (15)	N6—H6B	0.884 (19)	
N4—N3	1.3943 (13)	N5—C8	1.3547 (15)	
N4—C8	1.3337 (15)	N5—H5A	0.896 (17)	
N4—H4	0.883 (16)	N5—H5B	0.910 (18)	
H1A—O1—H1B	104.3 (17)	C8—N5—H5A	114.5 (10)	

C8—N2—C7	102.83 (9)	C8—N5—H5B	114.6 (10)
N3—N4—H4	119.2 (10)	H5A—N5—H5B	119.2 (14)
C8—N4—N3	109.80 (9)	N2—C8—N5	125.22 (10)
C8—N4—H4	131.0 (10)	N4—C8—N2	110.20 (10)
C7—N3—N4	101.79 (9)	N4—C8—N5	124.49 (11)
C7—N6—H6A	112.7 (11)	N2—C7—N6	121.27 (10)
C7—N6—H6B	112.5 (11)	N3—C7—N2	115.36 (10)
H6A—N6—H6B N4—N3—C7—N2 N4—N3—C7—N6 N3—N4—C8—N2 N3—N4—C8—N5 C8—N2—C7—N3	1.14 (12) -174.87 (10) 0.66 (13) -176.03 (10) -0.79 (12)	N3-C7-N6 C8-N2-C7-N6 C8-N4-N3-C7 C7-N2-C8-N4 C7-N2-C8-N5	123.25 (11) 175.30 (10) -1.05 (12) 0.04 (12) 176.69 (11)

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D····A	D—H···A
N5—H5A····O1 ⁱ	0.895 (17)	2.080 (17)	2.9173 (14)	155.4 (14)
O1—H1A····N3 ⁱⁱ	0.871 (19)	1.988 (19)	2.8558 (13)	174.1 (16)
O1—H1 <i>B</i> …N2	0.90 (2)	1.95 (2)	2.8410 (13)	171.5 (18)
N4—H4…N3 ⁱⁱⁱ	0.883 (17)	2.306 (16)	2.9947 (14)	134.9 (13)

Symmetry codes: (i) x, -y+3/2, z+1/2; (ii) -x+1, y+1/2, -z+1/2; (iii) -x+2, -y+1, -z+1.