

Bis(5-amino-3-carboxy-1*H*-1,2,4-triazol-4-ium) sulfate dihydrate

Amira Ouakkaf,^a Fadila Berrah,^{a*‡} Sofiane Bouacida^{b‡} and Thierry Roisnel^c

^aLaboratoire de Chimie Appliquée et Technologie des Matériaux LCATM, Université Larbi Ben M'Hidi, 04000 Oum El Bouaghi, Algeria, ^bUnité de Recherche de Chimie de l'Environnement et Moléculaire Structurale, CHEMS, Faculté des Sciences Exactes, Université Mentouri Constantine 25000, Algeria, and ^cCentre de Difractométrie X, UMR 6226 CNRS Unité Sciences Chimiques de Rennes, Université de Rennes I, 263 Avenue du Général Leclerc, 35042 Rennes, France
Correspondence e-mail: fadilaber@yahoo.fr

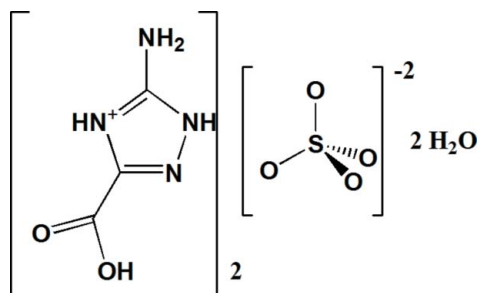
Received 3 April 2011; accepted 12 April 2011

Key indicators: single-crystal X-ray study; $T = 150$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.029; wR factor = 0.077; data-to-parameter ratio = 13.4.

The crystal structure of the title compound, $2\text{C}_3\text{H}_5\text{N}_4\text{O}_2^+ \cdot \text{SO}_4^{2-} \cdot 2\text{H}_2\text{O}$, displays a three-dimensional framework in which mixed infinite chains [oriented parallel to (510) and $(\bar{5}10)$] interfere, forming tunnels extending parallel to the c axis. Intermolecular $\text{O}-\text{H} \cdots \text{O}$, $\text{N}-\text{H} \cdots \text{O}$ and $\text{O}-\text{H} \cdots \text{N}$ hydrogen bonds ensure the unity of the structure and generate centrosymmetric $R_4^4(14)$ and $R_4^2(8)$ rings. The S atom lies on a twofold axis.

Related literature

For uses of 1,2,4 triazole derivatives, see: Beckmann & Brooker, (2003); Bhargava *et al.* (1981); Fujigaya *et al.* (2003); Hirota *et al.* (1991); Li *et al.* (2004); Matulková *et al.* (2008). For related hybrid compounds, see: Berrah *et al.* (2011*a,b,c*). For hydrogen-bond motifs, see: Bernstein *et al.* (1995); Etter *et al.* (1990).



‡ Current address: Département Sciences de la Matière, Faculté des Sciences Exactes et Sciences de la Nature et de la Vie, Université Larbi Ben M'hidi, 04000 Oum El Bouaghi, Algeria.

Experimental

Crystal data

$2\text{C}_3\text{H}_5\text{N}_4\text{O}_2^+ \cdot \text{SO}_4^{2-} \cdot 2\text{H}_2\text{O}$
 $M_r = 390.31$
Monoclinic, $C2/c$
 $a = 19.4350$ (9) Å
 $b = 5.8467$ (2) Å
 $c = 13.3343$ (6) Å
 $\beta = 109.981$ (2)°

$V = 1423.98$ (10) Å³
 $Z = 4$
Mo $K\alpha$ radiation
 $\mu = 0.31$ mm⁻¹
 $T = 150$ K
 $0.48 \times 0.08 \times 0.06$ mm

Data collection

Bruker APEXII diffractometer
11018 measured reflections
1620 independent reflections

1470 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.032$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.029$
 $wR(F^2) = 0.077$
 $S = 1.08$
1620 reflections
121 parameters

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\text{max}} = 0.37$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.41$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

| $D-\text{H} \cdots A$ | $D-\text{H}$ | $\text{H} \cdots A$ | $D \cdots A$ | $D-\text{H} \cdots A$ |
|---|--------------|---------------------|--------------|-----------------------|
| $\text{O1W}-\text{H2W} \cdots \text{N2}$ | 0.83 (2) | 2.16 (2) | 2.9774 (16) | 169 (2) |
| $\text{O1W}-\text{H1W} \cdots \text{O1}^{\text{i}}$ | 0.89 (2) | 1.93 (2) | 2.7941 (17) | 163 (2) |
| $\text{O2}-\text{H2} \cdots \text{O1W}^{\text{ii}}$ | 0.82 | 1.74 | 2.5403 (16) | 165 |
| $\text{N1}-\text{H1} \cdots \text{O4}^{\text{iii}}$ | 0.86 | 1.94 | 2.7107 (16) | 149 |
| $\text{N3}-\text{H3} \cdots \text{O4}^{\text{iv}}$ | 0.86 | 1.79 | 2.6436 (15) | 171 |
| $\text{N4}-\text{H4A} \cdots \text{O3}^{\text{v}}$ | 0.86 | 2.17 | 2.8452 (17) | 136 |
| $\text{N4}-\text{H4B} \cdots \text{O3}^{\text{vi}}$ | 0.86 | 2.08 | 2.9255 (16) | 168 |

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

Data collection: APEX2 (Bruker, 2001); cell refinement: SAINT (Bruker, 2001); data reduction: SAINT; program(s) used to solve structure: SIR2002 (Burla *et al.*, 2005); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: ORTEP-3 for Windows (Farrugia, 1997) and DIAMOND (Brandenburg & Berndt, 2001); software used to prepare material for publication: WinGX (Farrugia, 1999).

We are grateful to the LCATM Laboratory, Université Larbi Ben M'Hidi, Oum El Bouaghi, Algeria, for financial support.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BQ2295).

References

- Beckmann, U. & Brooker, S. (2003). *Coord. Chem. Rev.* **245**, 17–29.
Bernstein, J., Davis, R. E., Shimon, L. & Chang, N.-L. (1995). *Angew. Chem. Int. Ed. Engl.* **34**, 1555–1573.
Berrah, F., Ouakkaf, A., Bouacida, S. & Roisnel, T. (2011*a*). *Acta Cryst.* **E67**, o525–o526.
Berrah, F., Ouakkaf, A., Bouacida, S. & Roisnel, T. (2011*b*). *Acta Cryst.* **E67**, o677–o678.
Berrah, F., Ouakkaf, A., Bouacida, S. & Roisnel, T. (2011*c*). *Acta Cryst.* **E67**, o953–o954.
Bhargava, K. P., Tandon, M. & Bhalla, T. N. (1981). *Indian J. Chem. Sect. B*, **20**, 1017–1018.

- Brandenburg, K. & Berndt, M. (2001). *DIAMOND*. Crystal Impact, Bonn, Germany.
- Bruker (2001). *APEX2* and *SAINT*. Bruker AXS Inc., Madison, Wisconsin, USA.
- Burla, M. C., Caliandro, R., Camalli, M., Carrozzini, B., Cascarano, G. L., De Caro, L., Giacovazzo, C., Polidori, G. & Spagna, R. (2005). *J. Appl. Cryst.* **38**, 381–388.
- Etter, M. C., MacDonald, J. C. & Bernstein, J. (1990). *Acta Cryst.* **B46**, 256–262.
- Farrugia, L. J. (1997). *J. Appl. Cryst.* **30**, 565.
- Farrugia, L. J. (1999). *J. Appl. Cryst.* **32**, 837–838.
- Fujigaya, T., Jiang, D. L. & Aida, T. (2003). *J. Am. Chem. Soc.* **125**, 14690–14691.
- Hirota, T., Sasaki, K., Yamamoto, T. & Nakayama, T. (1991). *J. Heterocycl. Chem.* **28**, 257–261.
- Li, W., Wu, Q., Yu, Y., Luo, M., Hu, L., Gu, Y., Niu, F. & Hu, J. (2004). *Spectrochim. Acta Part A*, **60**, 2343–2354.
- Matulková, I., Nemeč, I., Teubner, K., Nemeč, P. & Micka, Z. (2008). *J. Mol. Struct.* **873**, 46–60.
- Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.

supplementary materials

Acta Cryst. (2011). E67, o1171-o1172 [doi:10.1107/S1600536811013882]

Bis(5-amino-3-carboxy-1*H*-1,2,4-triazol-4-ium) sulfate dihydrate

A. Ouakkaf, F. Berrah, S. Bouacida and T. Roisnel

Comment

1,2,4-triazole derivatives are remarkable substances exhibiting a wide range of properties; while several compounds incorporating 1,2,4-triazole moieties show pharmacological and biological activities (antidepressants, anti-inflammatory, fungicides) (Bhargava *et al.*, 1981; Hirota *et al.*, 1991; Li *et al.*, 2004), a number of them find use in material field and coordination chemistry (magnetic and non-linear optics properties, multidentate ligands) (Beckmann & Brooker, 2003; Fujigaya *et al.*, 2003; Matulková *et al.*, 2008).

Our recent investigation on some N-heterocycle compounds with inorganic acids (Berrah *et al.*, 2011*a,b,c*), has revealed their ability to generate original networks stabilized in particular by hydrogen bonds; we report herein the structure of a new hybrid compound obtained from a disubstituted 1,2,4-triazole derivative (5-amino-1,2,4 triazol-1*H*-3-carboxylic acid hydrate) and the sulfuric acid.

The molecular structure and the atom-numbering scheme of the title compound are shown in Fig. 1. The sulfur atom lies in the twofold axis and the sulfate anion shows a quite regular geometry compared with that seen in similar compounds (Berrah *et al.*, 2011*b,c*). The triazole ring exhibits a short distance of 1.3046 (18) Å revealing the double-bond character of the C2=N2 bond, two long distances 1.3648 (18) Å and 1.3774 (17) Å related to the single bonds C2—N3 and N1—N2, respectively. The C3—N1 and C3=N3 bonds lengths are respectively 1.3420 (18) Å and 1.3480 (18) Å which suggests the delocalization of the double bond (N1 \cdots C3 \cdots N3).

The crystal structure of the title compound (Fig. 2) displays a three-dimensional framework where mixed infinite chains interfere to form tunnels parallel the *c* axis. According to their orientation, we can distinguish two kinds of chains: the first one directed along the (510) plane and the second along the ($\bar{5}$ 10) plane (Fig. 2). In a mixed chain, the cations adopt a head to head configuration in a way that NH₂, of two adjacent cations, are linked (*via* N—H \cdots O hydrogen bonds) to two sulfate anions and C—OH are linked (by O—H \cdots O hydrogen bonds) to two water molecules (Fig. 3) (Table 1). Consequently, centro-symmetric $R^4_4(14)$ and $R^2_4(8)$ graph-set rings are created (Etter *et al.*, 1990; Bernstein *et al.*, 1995).

Experimental

The title compound was synthesized by reacting 5-amino-1,2,4 triazol-1*H*-3-carboxylic acid hydrate with some excess of sulfuric acid in aqueous solution. Slow evaporation leads to well crystallized colorless needles.

Refinement

The H atoms of the water molecules were located in difference Fourier maps and were refined with $U_{\text{iso}}(\text{H}) = 1.5 U_{\text{eq}}(\text{O})$. The remaining H atoms were located in difference Fourier maps but introduced in calculated positions and treated as riding on their parent atoms (N or O) with, O—H = 0.82 Å and N—H = 0.86 Å with $U_{\text{iso}}(\text{H}) = 1.2 U_{\text{eq}}(\text{N})$ and $U_{\text{iso}}(\text{H}) = 1.5 U_{\text{eq}}(\text{O})$.

Figures

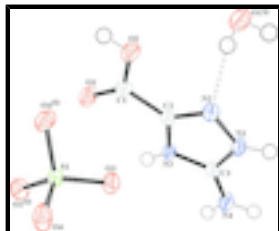


Fig. 1. The molecular structure of the title compound with the atomic labeling scheme. Displacement are drawn at the 50% probability level. Hydrogen bonds are shown as dashed lines.

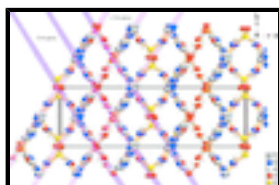


Fig. 2. (Brandenburg & Berndt, 2001) A diagram of the three-dimensional packing of (I). (510) and (-510) planes have been presented to illustrate the mixed infinite chains. Hydrogen bonds are shown as dashed lines.

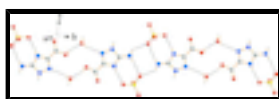
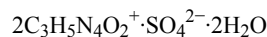


Fig. 3. (Brandenburg & Berndt, 2001) A view of the mixed infinite chain. Hydrogen bonds are shown as dashed lines.

Bis(5-amino-3-carboxy-1*H*-1,2,4-triazol-4-ium) sulfate dihydrate

Crystal data



$M_r = 390.31$

Monoclinic, $C2/c$

$a = 19.4350$ (9) Å

$b = 5.8467$ (2) Å

$c = 13.3343$ (6) Å

$\beta = 109.981$ (2)°

$V = 1423.98$ (10) Å³

$Z = 4$

$F(000) = 808$

$D_x = 1.821$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 1620 reflections

$\theta = 3.3\text{--}27.4^\circ$

$\mu = 0.31$ mm⁻¹

$T = 150$ K

Needle, colourless

$0.48 \times 0.08 \times 0.06$ mm

Data collection

Bruker APEXII
diffractometer

graphite

CCD rotation images, thin slices scans

11018 measured reflections

1620 independent reflections

1470 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.032$

$\theta_{\text{max}} = 27.5^\circ$, $\theta_{\text{min}} = 3.3^\circ$

$h = -24 \rightarrow 25$

$k = -7 \rightarrow 7$

$l = -17 \rightarrow 15$

Refinement

Refinement on F^2

Least-squares matrix: full

Primary atom site location: structure-invariant direct methods

Secondary atom site location: difference Fourier map

$$R[F^2 > 2\sigma(F^2)] = 0.029$$

$$wR(F^2) = 0.077$$

$$S = 1.08$$

1620 reflections

121 parameters

0 restraints

Hydrogen site location: inferred from neighbouring sites

H atoms treated by a mixture of independent and constrained refinement

$$w = 1/[\sigma^2(F_o^2) + (0.0343P)^2 + 1.9446P]$$

$$\text{where } P = (F_o^2 + 2F_c^2)/3$$

$$(\Delta/\sigma)_{\max} = 0.009$$

$$\Delta\rho_{\max} = 0.37 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\min} = -0.41 \text{ e } \text{\AA}^{-3}$$

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 (\AA^2)

| | x | y | z | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|-----|-------------|--------------|--------------|----------------------------------|
| C1 | 0.32927 (7) | 0.3566 (2) | 0.60560 (11) | 0.0136 (3) |
| C2 | 0.35578 (7) | 0.2304 (2) | 0.52873 (11) | 0.0123 (3) |
| C3 | 0.40656 (7) | -0.0381 (2) | 0.46589 (11) | 0.0129 (3) |
| N1 | 0.38002 (6) | 0.1265 (2) | 0.39270 (9) | 0.0138 (3) |
| H1 | 0.3827 | 0.125 | 0.3296 | 0.017* |
| N2 | 0.34797 (6) | 0.2981 (2) | 0.43222 (9) | 0.0138 (3) |
| N3 | 0.39065 (6) | 0.0245 (2) | 0.55278 (9) | 0.0123 (2) |
| H3 | 0.4005 | -0.0502 | 0.6116 | 0.015* |
| N4 | 0.44239 (7) | -0.2252 (2) | 0.45703 (10) | 0.0169 (3) |
| H4A | 0.451 | -0.2519 | 0.399 | 0.02* |
| H4B | 0.4571 | -0.3201 | 0.5093 | 0.02* |
| O1 | 0.33947 (6) | 0.28073 (19) | 0.69430 (8) | 0.0188 (2) |
| O2 | 0.29439 (6) | 0.54386 (18) | 0.56322 (8) | 0.0186 (2) |
| H2 | 0.2817 | 0.612 | 0.6077 | 0.028* |
| O3 | 0.48758 (5) | 0.50548 (17) | 0.65377 (8) | 0.0158 (2) |
| O4 | 0.56503 (6) | 0.79464 (19) | 0.76558 (8) | 0.0190 (2) |
| S1 | 0.5 | 0.64445 (8) | 0.75 | 0.01101 (13) |
| O1W | 0.25419 (6) | 0.69140 (19) | 0.32536 (9) | 0.0194 (2) |
| H1W | 0.2751 (11) | 0.724 (4) | 0.2772 (17) | 0.029* |
| H2W | 0.2786 (11) | 0.586 (4) | 0.3624 (17) | 0.029* |

supplementary materials

Atomic displacement parameters (\AA^2)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|-----|------------|------------|------------|-------------|--------------|-------------|
| C1 | 0.0118 (6) | 0.0150 (7) | 0.0148 (7) | -0.0021 (5) | 0.0054 (5) | -0.0017 (5) |
| C2 | 0.0117 (6) | 0.0128 (6) | 0.0122 (7) | -0.0015 (5) | 0.0037 (5) | -0.0008 (5) |
| C3 | 0.0119 (6) | 0.0151 (7) | 0.0117 (6) | -0.0026 (5) | 0.0040 (5) | -0.0008 (5) |
| N1 | 0.0177 (6) | 0.0155 (6) | 0.0097 (6) | 0.0011 (5) | 0.0066 (5) | -0.0003 (4) |
| N2 | 0.0150 (6) | 0.0143 (6) | 0.0126 (6) | -0.0001 (4) | 0.0053 (5) | -0.0008 (4) |
| N3 | 0.0145 (5) | 0.0138 (6) | 0.0093 (5) | 0.0010 (4) | 0.0052 (4) | 0.0015 (4) |
| N4 | 0.0220 (6) | 0.0162 (6) | 0.0145 (6) | 0.0034 (5) | 0.0087 (5) | -0.0002 (5) |
| O1 | 0.0230 (5) | 0.0213 (5) | 0.0155 (5) | 0.0024 (4) | 0.0108 (4) | 0.0015 (4) |
| O2 | 0.0227 (5) | 0.0175 (5) | 0.0170 (5) | 0.0062 (4) | 0.0086 (4) | -0.0006 (4) |
| O3 | 0.0182 (5) | 0.0164 (5) | 0.0125 (5) | -0.0004 (4) | 0.0051 (4) | -0.0035 (4) |
| O4 | 0.0221 (5) | 0.0243 (6) | 0.0133 (5) | -0.0105 (4) | 0.0094 (4) | -0.0049 (4) |
| S1 | 0.0129 (2) | 0.0119 (2) | 0.0090 (2) | 0 | 0.00474 (17) | 0 |
| O1W | 0.0228 (6) | 0.0202 (5) | 0.0187 (6) | 0.0065 (4) | 0.0117 (5) | 0.0044 (4) |

Geometric parameters (\AA , $^\circ$)

| | | | |
|-------------|--------------|-------------------------------------|--------------|
| C1—O1 | 1.2144 (18) | N3—H3 | 0.86 |
| C1—O2 | 1.3098 (17) | N4—H4A | 0.86 |
| C1—C2 | 1.4905 (19) | N4—H4B | 0.86 |
| C2—N2 | 1.3046 (18) | O2—H2 | 0.82 |
| C2—N3 | 1.3648 (18) | O3—S1 | 1.4674 (10) |
| C3—N4 | 1.3236 (19) | O4—S1 | 1.4941 (10) |
| C3—N1 | 1.3420 (18) | S1—O3 ⁱ | 1.4674 (10) |
| C3—N3 | 1.3480 (18) | S1—O4 ⁱ | 1.4941 (10) |
| N1—N2 | 1.3774 (17) | O1W—H1W | 0.89 (2) |
| N1—H1 | 0.86 | O1W—H2W | 0.83 (2) |
| O1—C1—O2 | 127.86 (13) | C3—N3—H3 | 126.9 |
| O1—C1—C2 | 120.61 (13) | C2—N3—H3 | 126.9 |
| O2—C1—C2 | 111.50 (12) | C3—N4—H4A | 120 |
| N2—C2—N3 | 112.37 (12) | C3—N4—H4B | 120 |
| N2—C2—C1 | 125.24 (13) | H4A—N4—H4B | 120 |
| N3—C2—C1 | 122.38 (12) | C1—O2—H2 | 109.5 |
| N4—C3—N1 | 127.67 (13) | O3—S1—O3 ⁱ | 112.76 (9) |
| N4—C3—N3 | 125.69 (13) | O3—S1—O4 | 109.00 (6) |
| N1—C3—N3 | 106.63 (12) | O3 ⁱ —S1—O4 | 108.98 (6) |
| C3—N1—N2 | 110.85 (11) | O3—S1—O4 ⁱ | 108.98 (6) |
| C3—N1—H1 | 124.6 | O3 ⁱ —S1—O4 ⁱ | 109.00 (6) |
| N2—N1—H1 | 124.6 | O4—S1—O4 ⁱ | 108.01 (9) |
| C2—N2—N1 | 103.94 (11) | H1W—O1W—H2W | 106.3 (19) |
| C3—N3—C2 | 106.20 (11) | | |
| O1—C1—C2—N2 | -179.54 (13) | C1—C2—N2—N1 | -179.34 (12) |
| O2—C1—C2—N2 | 2.06 (19) | C3—N1—N2—C2 | -0.42 (14) |
| O1—C1—C2—N3 | 1.6 (2) | N4—C3—N3—C2 | 177.99 (13) |

| | | | |
|-------------|--------------|-------------|--------------|
| O2—C1—C2—N3 | -176.82 (12) | N1—C3—N3—C2 | -1.19 (14) |
| N4—C3—N1—N2 | -178.13 (13) | N2—C2—N3—C3 | 1.00 (16) |
| N3—C3—N1—N2 | 1.03 (15) | C1—C2—N3—C3 | -179.99 (12) |
| N3—C2—N2—N1 | -0.37 (15) | | |

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

Hydrogen-bond geometry ($\text{\AA}, ^\circ$)

| <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> |
|-----------------------------------|-------------|---------------------|----------------------------|-------------------------------|
| O1W—H2W \cdots N2 | 0.83 (2) | 2.16 (2) | 2.9774 (16) | 169 (2) |
| O1W—H1W \cdots O1 ⁱⁱ | 0.89 (2) | 1.93 (2) | 2.7941 (17) | 163 (2) |
| O2—H2 \cdots O1W ⁱⁱⁱ | 0.82 | 1.74 | 2.5403 (16) | 165 |
| N1—H1 \cdots O4 ^{iv} | 0.86 | 1.94 | 2.7107 (16) | 149 |
| N3—H3 \cdots O4 ^v | 0.86 | 1.79 | 2.6436 (15) | 171 |
| N4—H4A \cdots O3 ^{vi} | 0.86 | 2.17 | 2.8452 (17) | 136 |
| N4—H4B \cdots O3 ^{vii} | 0.86 | 2.08 | 2.9255 (16) | 168 |

Symmetry codes: (ii) $x, -y+1, z-1/2$; (iii) $-x+1/2, -y+3/2, -z+1$; (iv) $-x+1, -y+1, -z+1$; (v) $-x+1, y-1, -z+3/2$; (vi) $-x+1, -y, -z+1$; (vii) $x, y-1, z$.

Fig. 1

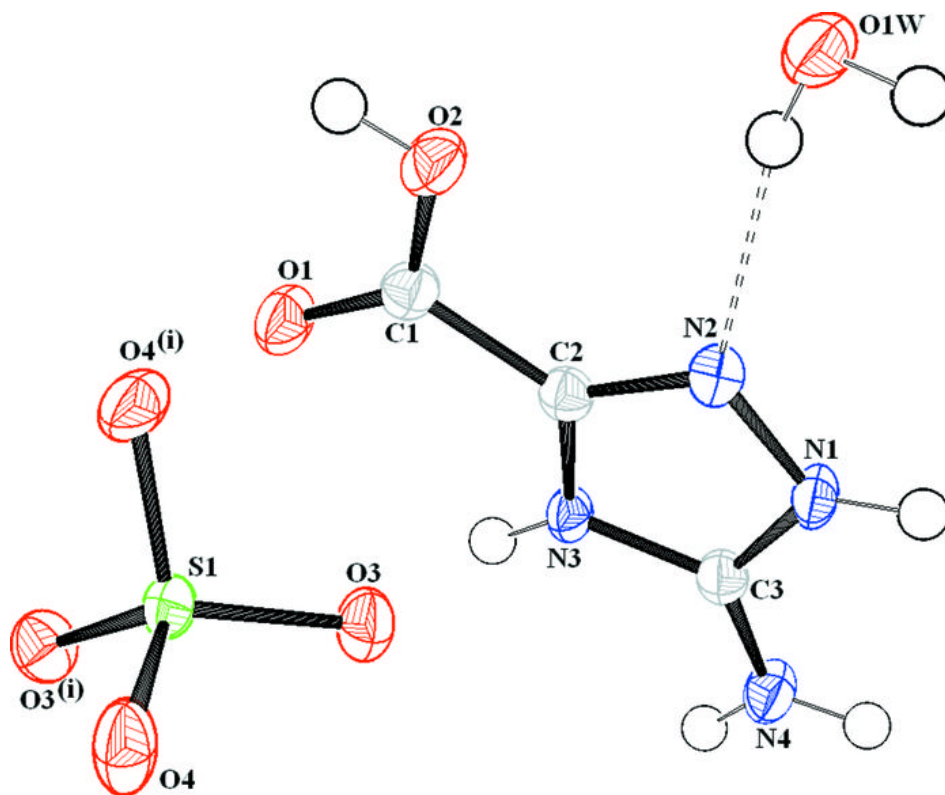


Fig. 2

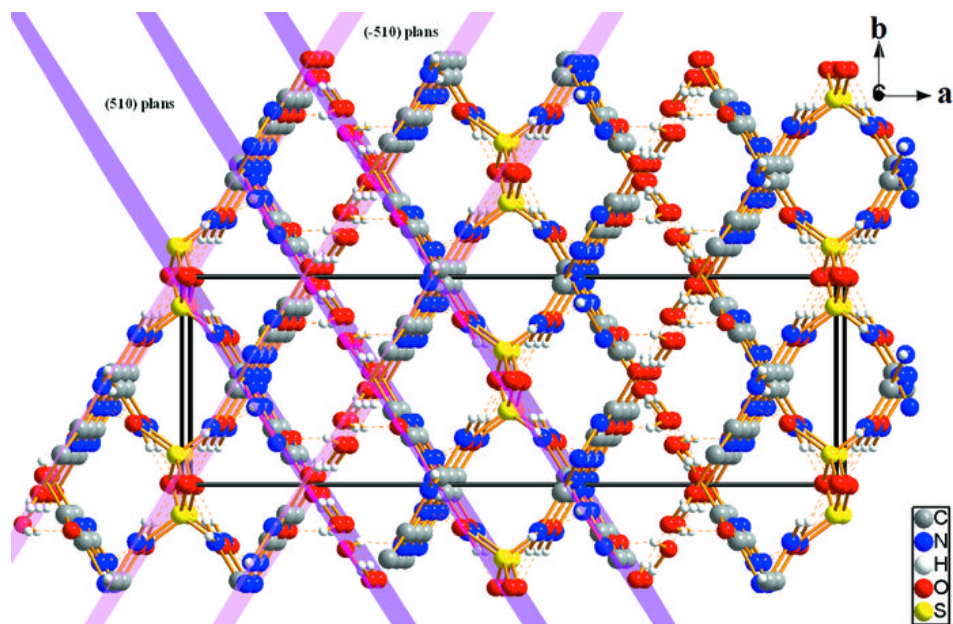


Fig. 3

