organic papers

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

Mohd Mustaqim Rosli,^a Hoong-Kun Fun,^a* Beck Sim Lee,^a Pumsak Ruanwas^b and Suchada Chantrapromma^b*

^aX-ray Crystallography Unit, School of Physics, Universiti Sains Malaysia, 11800 USM, Penang, Malaysia, and ^bDepartment of Chemistry, Faculty of Science, Prince of Songkla University, Hat-Yai, Songkhla 90112, Thailand

Correspondence e-mail: hkfun@usm.my, suchada.c@psu.ac.th

Key indicators

Single-crystal X-ray study T = 297 KMean $\sigma(C-C) = 0.001 \text{ Å}$ Disorder in solvent or counterion R factor = 0.055 wR factor = 0.175 Data-to-parameter ratio = 31.5

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. Hexamethylenetetraminium 2,4-dinitrobenzoate monohydrate

In the crystal structure of the proton-transfer compound, $C_6H_{13}N_4^+ \cdot C_7H_3N_2O_6^- \cdot H_2O$, the hexamethylenetetraminium and 2,4-dinitrobenzoate ions are linked by bifurcated N-H···O hydrogen bonds. The crystal structure is stabilized by $O-H \cdot \cdot \cdot N$ hydrogen bonds and weak $C-H \cdot \cdot \cdot O$ interactions to form a three-dimensional network. The benzene rings of the 2,4-dinitrobenzoate ion make a π - π interaction with a centroid-centroid distance of 3.689 (1) Å.

Comment

The interactions of organic acids with amine bases in the solid phase have been of interest owing to their hydrogen-bonding interactions (Coupar *et al.*, 1997; Sobczyk *et al.*, 2000; Smith, Wermuth & White, 2001; Smith, Wermuth, Bott *et al.*, 2001; Fun *et al.*, 2003; Chantrapromma & Fun, 2004) and their important role in biological systems (Blow, 1976). Following our continuing studies of hydrogen-bonding interactions and phase transitions due to hydrogen bonding (Usman *et al.*, 2001; Fun *et al.*, 2003; How *et al.*, 2005; Chantrapromma *et al.*, 2006), we present here another proton-transfer compound of hexamethylenetetramine with a nitro-substituted aromatic carboxylic acid, (I).



In the asymmetric unit of (I), all bond lengths and angles show normal values (Allen *et al.*, 1987). The carboxylate and the neighbouring nitro group are twisted away from the plane of the benzene ring, with torsion angles O5-C7-C4-C3 = $40.66 (16)^{\circ}$, $O6-C7-C4-C5 = 43.87 (15)^{\circ}$, O3-N5-C3- $C2 = 53.26 (15)^{\circ}$ and $O4-N5-C3-C4 = 51.83 (15)^{\circ}$, while the other nitro group is located almost in the plane of the benzene ring, with torsion angles O1-N6-C1-C6 = $0.03 (14)^{\circ}$ and $O1-N6-C1-C2 = 179.99 (9)^{\circ}$. The dihedral angle between the planes of the adjacent carboxylate and nitro groups is $51.3 (1)^{\circ}$. All the six-membered N/C/N/C/N/C rings of the hexamethylenetetraminium ion adopt a chair conformation, the puckering parameters Q, θ and φ being Received 16 May 2006 Accepted 22 June 2006

© 2006 International Union of Crystallography All rights reserved



Figure 1

The asymmetric unit of (I), showing 30% probability displacement ellipsoids and the atomic numbering. Hydrogen bonds are shown as dashed lines. Both disorder components are shown.



Figure 2

The crystal packing of (I), viewed down the a axis. Hydrogen bonds are shown as dashed lines. Only one disorder component (O1W) is shown.

0.598 (1) Å, 179.2 (1)° and 25 (10)°, respectively, for N1/C8/N2/C11/N3/C9, 0.579 (1) Å, 0.8 (1)° and 262 (7)°, respectively, for N1/C8/N2/C12/N4/C10, 0.590 (1) Å, 178.7 (1)° and 259 (5)°, respectively, for N1/C9/N3/C13/N4/C10, and 0.597 (1) Å, 1.4 (1)° and 153 (5)°, respectively, for N2/C11/N3/C13/N4/C12.

The hexamethylenetetraminium and 2.4-dinitrobenzoate ions are connected by bifurcated hydrogen bonds, N3-H13...O5 and N3-H13...O6 (Table 1). The 2,4-dinitrobenzoate ion also forms a hydrogen bond $O1W - H1W1 \cdots O5$ $(O2W-H1W2\cdots O5)$ with the disordered water molecule. The crystal structure of (I) is stabilized by intermolecular hydrogen bonds $O1W - H1W1 \cdots N4^{i}$, $C11 - H9 \cdots O1^{ii}$ C13-H12···O4ⁱⁱⁱ, C13-H12···O1 W^i and C5-H16···O6^{iv} (symmetry codes as in Table 1), which form chains in the three-dimensional molecular network. A short contact interaction of O1···N1(1 - x, -1 - y, -z) [2.926 (2) Å] and a π - π interaction between the benzene rings of 2,4-dinitrobenzoate ions with a centroid-centroid distance of 3.689 (1) Å also contribute to the stabilization of the crystal structure.



Figure 3

The crystal packing of (I), viewed down the a axis. Hydrogen bonds are shown as dashed lines. Only the second disorder component (O2W) is shown.

Experimental

Compound (I) was prepared by mixing thoroughly hexamethylenetetramine (0.70 g, 5 mmol) and 2,4-dinitrobenzoic acid (1.06 g, 5 mmol), which were dissolved in acetone (100 ml), adding a few drops of water. The resulting mixture was warmed to the temperature range 313–318 K, whereupon a clear solution was obtained. The solution was then filtered and left to evaporate slowly at room temperature. Colourless single crystals suitable for X-ray diffraction studies were obtained from the solution after a few days (m.p. 447–449 K).

Crystal data

$C_{6}H_{13}N_{4}^{+} \cdot C_{7}H_{3}N_{2}O_{6}^{-} \cdot H_{2}O_{6}$	V = 796.09 (2) Å ³
$M_r = 370.33$	Z = 2
Friclinic, $P\overline{1}$	$D_x = 1.545 \text{ Mg m}^{-3}$
a = 6.4118 (1) Å	Mo $K\alpha$ radiation
p = 9.1042 (2) Å	$\mu = 0.13 \text{ mm}^{-1}$
z = 14.1521 (2) Å	T = 297 (2) K
$\alpha = 77.986 \ (1)^{\circ}$	Block, colourless
$\beta = 80.464 \ (1)^{\circ}$	$0.55 \times 0.43 \times 0.33 \text{ mm}$
$\chi = 85.585 (1)^{\circ}$	

Data collection

```
Bruker SMART APEX2 CCD area-
detector diffractometer \omega scans
Absorption correction: multi-scan
(SADABS; Bruker, 2005)
T_{\min} = 0.855, T_{\max} = 0.959
```

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.055$ $wR(F^2) = 0.175$ S = 1.067817 reflections 248 parameters H atoms treated by a mixture of independent and constrained refinement 23506 measured reflections 7817 independent reflections 5034 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.023$ $\theta_{\text{max}} = 37.2^{\circ}$

```
\begin{split} w &= 1/[\sigma^2(F_{\rm o}^2) + (0.0818P)^2 \\ &+ 0.0892P] \\ \text{where } P &= (F_{\rm o}^2 + 2F_{\rm c}^2)/3 \\ (\Delta/\sigma)_{\rm max} &< 0.001 \\ \Delta\rho_{\rm max} &= 0.30 \text{ e } \text{ Å}^{-3} \\ \Delta\rho_{\rm min} &= -0.27 \text{ e } \text{ Å}^{-3} \end{split}
```

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

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
O1W-H1W1O5	0.84 (1)	1.80(1)	2.633 (2)	176 (1)
$O2W-H1W2\cdots O5$	0.87(1)	2.06 (1)	2.912 (3)	167 (1)
N3-H13···O5	0.88	2.18	2.935 (2)	144
N3-H13···O6	0.88	2.20	2.916 (2)	138
$O1W - H2W1 \cdot \cdot \cdot N4^{i}$	0.83(1)	2.26(1)	3.053 (3)	158 (1)
C11−H9···O1 ⁱⁱ	0.97	2.42	3.352 (2)	159
C13-H12···O4 ⁱⁱⁱ	0.97	2.51	3.207 (2)	128
$C13-H12\cdots O1W^{i}$	0.97	2.59	3.069 (3)	110
$C5-H16\cdots O6^{iv}$	0.93	2.53	3.264 (2)	136

-x+2, -y, -z.

All C-bound H atoms were placed in calculated positions with C– H distances in the range 0.93–0.97 Å and refined as riding, with $U_{iso}(H) = 1.2U_{eq}(C)$. The O-bound and N-bound H atoms were located in a difference map. For the O-bound H atoms the positions were freely refined with $U_{iso}(H) = 1.5U_{eq}(O)$, while for the N-bound H atom the positional and isotropic displacement parameters were constrained as riding, with N–H = 0.88 Å and $U_{iso}(H) = 0.063$. The water molecule is disordered over two sites with relative occupancies of 0.574 (5) and 0.426 (5).

Data collection: *APEX2* (Bruker, 2005); cell refinement: *APEX2*; data reduction: *SAINT* (Bruker, 2005); program(s) used to solve structure: *SHELXTL* (Sheldrick, 1998); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*, *PARST* (Nardelli, 1995) and *PLATON* (Spek, 2003).

The authors thank the Malaysian Government and Universiti Sains Malaysia for Scientific Advancement Grant Allocation (SAGA) No. 304/PFIZIK/653003/A118. SC also thanks the Prince of Songkla University for partial financial support.

References

Allen, F. H., Kennard, O., Watson, D. G., Brammer, L., Orpen, A. G. & Taylor, R. (1987). J. Chem. Soc. Perkin Trans. 2, pp. S1–19.

Blow, D. M. (1976). Acc. Chem. Res. 9, 145-156.

Bruker (2005). *APEX2* (Version 1.27), *SAINT* (Version 7.12A) and *SADABS* (Version 2004/1). Bruker AXS Inc., Madison, Wisconsin, USA.

Chantrapromma, S. & Fun, H.-K. (2004). Acta Cryst. E60, o1250-o1252.

Chantrapromma, S., Fun, H.-K. & Usman, A. (2006). J. Mol. Struct. **789**, 30–36. Coupar, P. I., Glidewell, C. & Ferguson, G. (1997). Acta Cryst. B**53**, 521–533.

Fun, H. K., Usman, A., Chantrapromma, S., Osman, J., Ong, L. H., Tilley, D. R. & Ishibashi, Y. (2003). Solid State Commun. 127, 677–682.

How, P. T., Lee, B. S., Fun, H.-K., Razak, I. A. & Chantrapromma, S. (2005). *Phys. Rev. B*, **71**, 174109.

Nardelli, M. (1995). J. Appl. Cryst. 28, 659.

Sheldrick, G. M. (1998). SHELXTL. Version 5.1. Bruker AXS Inc., Madison, Wisconsin, USA.

Smith, G., Wermuth, U. D., Bott, R. C., White, J. M. & Willis, A. C. (2001). Aust. J. Chem. 54, 165–170.

Smith, G., Wermuth, U. D. & White, J. M. (2001). Aust. J. Chem. 54, 171–175. Sobczyk, L., Lis, T., Olejnik, Z. & Majerz, I. (2000). J. Mol. Struct. 552, 233–

- 241. Spek, A. L. (2003). J. Appl. Cryst. 36, 7-13.
- Usman, A., Chantrapromma, S. & Fun, H.-K. (2001). Acta Cryst. C57, 1443–1446.