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Key indicators

Single-crystal X-ray study T = 298 KMean $\sigma(C-C) = 0.003 \text{ Å}$ R factor = 0.037 wR factor = 0.105 Data-to-parameter ratio = 12.7

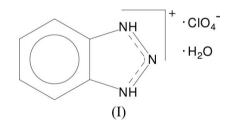
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

Benzotriazolium perchlorate monohydrate

The crystal structure of the title compound, $C_6H_6N_3^+$.-ClO₄⁻·H₂O, consists of cations, anions and water molecules linked by N-H···O and O-H···O hydrogen bonds into sheets of alternating $R_4^4(12)$ and $R_6^6(20)$ rings, which form chains running along the [101] direction. Received 14 March 2007 Accepted 21 March 2007

Comment

Benzotriazole and its derivatives are used mainly as corrosion inhibitors for metals, antifreeze agents, antidust agents for photography and polymer stabilizers, as well as in industrial cooling systems. They are also widely used in chemical syntheses (Patsalides & Robards, 1985; Gruden *et al.*, 2001; Cancilla, *et al.*, 2003; Katritzky & Rogovoy, 2003). The title compound, (I), was investigated as part of a structural study on hydrogen-bonding patterns in *N*-heterocyclic perchlorate salts (Sieroń, 2005*a*,*b*, 2007).



In (I), there are benzotriazolium cations, perchlorate anions and water molecules (Fig. 1). All the interatomic distances and angles are normal (Allen *et al.*, 1987). One of the H atoms of the benzotriazolium ion and one of the water molecule are

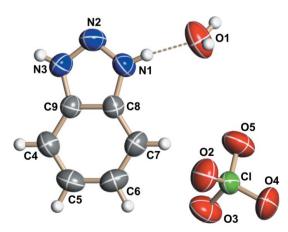


Figure 1

The asymmetric unit of (I) with the atom-numbering scheme. The displacement ellipsoids for the non-H atoms are drawn at the 50% probability level. The dashed line indicates a hydrogen bond.

© 2007 International Union of Crystallography All rights reserved engaged in a bifurcated unsymmetrical (strong and weak) hydrogen bond. Each of these hydrogen bonds involves two perchlorate O atoms (Table 1). A bifurcation is confirmed by the sums of angles about atoms H3 and H11, which are 354 and 359°, respectively (Jeffrey & Saenger, 1991). The ions and water molecules are linked by N-H···O and O-H···O hydrogen bonds, forming fused 12- and 20-membered cetrosymmetric rings, described by the graph-set notations as $R_4^4(12)$ and $R_6^6(20)$, respectively (Etter *et al.*, 1990). The substructure based on these motifs propagates along the $[10\overline{1}]$ direction (Fig. 2).

In addition, the benzotriazole rings are engaged in $\pi - \pi$ stacking interactions, with distances between ring centroids of ca 3.7 Å, and with perpendicular distances between equivalent planes of ca 3.5 Å. The closest interatomic distances $C5 \cdots C9(2 - x, 1 - y, -z)$ and $C7 \cdots C9(1 - x, 1 - y, -z)$ are 3.495 (3) and 3.498 (3) Å, respectively.

Experimental

The title compound was prepared by dissolving benzotriazole (1 mmol) in perchloric acid (4 mmol) in water (20 ml). After a few days, prism-shaped colourless crystals up to 1 mm in size were obtained at room temperature.

Crystal data

$C_6H_6N_3^+ \cdot \text{ClO}_4^- \cdot \text{H}_2\text{O}$ $M_r = 237.60$	$\gamma = 87.246 \ (2)^{\circ}$ V = 492.07 (3) Å ³
Triclinic, $P\overline{1}$	<i>Z</i> = 2
a = 7.4227 (2) Å b = 8.0524 (2) Å	Mo <i>K</i> α radiation $\mu = 0.40 \text{ mm}^{-1}$
c = 8.8973 (3) Å	T = 298 K
$\alpha = 74.023 \ (3)^{\circ}$ $\beta = 74.335 \ (3)^{\circ}$	$0.50 \times 0.50 \times 0.30 \text{ mm}$

Data collection

Kuma KM-4 CCD diffractometer	5415 measured reflections
Absorption correction: multi-scan	1925 independent reflections
(CrysAlis RED; Oxford Diffrac-	1842 reflections with $I > 2\sigma(I)$
tion, 2006)	$R_{\rm int} = 0.024$
$T_{\min} = 0.827, T_{\max} = 0.893$	

Refinement

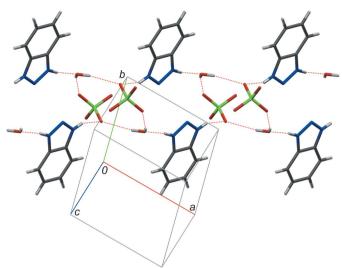
$R[F^2 > 2\sigma(F^2)] = 0.037$ $wR(F^2) = 0.105$ S = 1.09 1925 reflections	H atoms treated by a mixture of independent and constrained refinement $\Delta a = 0.26 \text{ e}^{-3}$
1925 reflections	$\Delta \rho_{\rm max} = 0.26 \ {\rm e} \ {\rm \AA}^{-3}$
151 parameters	$\Delta \rho_{\rm min} = -0.32 \ {\rm e} \ {\rm \AA}^{-3}$

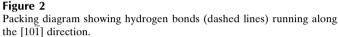
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Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdots A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$\begin{array}{c} \hline & \\ \hline N1-H1\cdots O1 \\ N3-H3\cdots O2^{i} \\ N3-H3\cdots O4^{ii} \\ O1-H12\cdots O2^{iii} \\ O1-H11\cdots O3^{iv} \\ O1-H11\cdots O3^{v} \\ \end{array}$	0.82 (3) 0.75 (3) 0.75 (3) 0.82 0.82 0.82 0.82	1.83 (3) 2.32 (3) 2.54 (3) 2.21 2.18 2.52	2.646 (2) 2.920 (3) 3.027 (3) 2.972 (3) 2.943 (3) 3.031 (3)	171 (3) 139 (3) 124 (2) 154 155 121

Symmetry codes: (i) -x + 1, -y + 1, -z; (ii) x + 1, y + 1, z - 1;(iii) -x, -y + 1, -z + 1; (iv) x, y + 1, z; (v) -x + 1, -y + 1, -z + 1.





All H atoms were initially located in a difference Fourier map. The arvl H atoms were repositioned with idealized geometry keeping C-H = 0.93 Å and they were refined using a riding model. The H atoms of the water molecule were refined with the O-H distances restrained to 0.82 Å initially, and then refined using a riding model. The positions of the amine H atoms and the atomic displacement parameters of all the H atoms were refined freely.

Data collection: CrysAlis CCD (Oxford Diffraction, 2006); cell refinement: CrysAlis RED (Oxford Diffraction, 2006); data reduction: CrysAlis RED; program(s) used to solve structure: SIR97 (Altomare et al., 1999); program(s) used to refine structure: SHELXTL (Sheldrick, 2001); molecular graphics: SHELXTL and Mercury (Macrae et al., 2006); software used to prepare material for publication: PLATON (Spek, 2003).

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.
- Altomare, A., Burla, M. C., Camalli, M., Cascarano, G. L., Giacovazzo, C., Guagliardi, A., Moliterni, A. G. G., Polidori, G. & Spagna, R. (1999). J. Appl. Cryst. 32, 115-119.
- Cancilla, D. A., Baird, J. C., Geis, S. W. & Corsi, S. R. (2003). Environ. Toxicol. Chem. 22, 134-140.
- Etter, M. C., MacDonald, J. C. & Bernstein, J. (1990). Acta Cryst. B46, 256-262.
- Gruden, C. L., Dow, S. M. & Hernandez, M. T. (2001). Water Environ. Res. 73, 72-79
- Jeffrey, G. A. & Saenger, W. (1991). Hydrogen Bonding in Biological Structures. New York: Springer-Verlag.
- Katritzky, A. R. & Rogovoy, B. V. (2003). Chem. Eur. J. 9, 4586-4593.
- Macrae, C. F., Edgington, P. R., McCabe, P., Pidcock, E., Shields, G. P., Taylor, R., Towler, M. & van de Streek, J. (2006). J. Appl. Cryst. 39, 453-457.
- Oxford Diffraction (2006). CrysAlis CCD and CrysAlis RED. Version 1.171. Oxford Diffraction Ltd, Abingdon, Oxfordshire, England.
- Patsalides, E. & Robards, K. (1985). J. Chromatogr. 331, 149-160.
- Sheldrick, G. M. (2001). SHELXTL. Version 6.12. Bruker AXS Inc., Madison, Wisconsin, USA.
- Sieroń, L. (2005a). Acta Cryst. E61, o2091-o2092.
- Sieroń, L. (2005b). Anal. Sci. (X-Ray Struct. Anal. Online), 21, x179-x180.
- Sieroń, L. (2007). Acta Cryst. E63, o1199-o1200.
- Spek, A. L. (2003). J. Appl. Cryst. 36, 7-13.