organic papers

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 288 KMean σ (C–C) = 0.003 Å R factor = 0.040 wR factor = 0.127 Data-to-parameter ratio = 15.6

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

1,10-Phenanthrolinium perchlorate

The proton-transfer title compound, $C_{12}H_9N_2^+ \cdot ClO_4^-$, has been obtained from 1,10-phenanthroline and HClO₄. In the crystal structure, the molecules stack along the *a* axis *via* hydrogen-bond interactions and van der Waals forces.

Comment

1,10-Phenanthroline (phen) has been recognized as a good proton acceptor and it is considered a suitable agent in the synthesis of proton-transfer systems. Some proton-transfer complexes of 1,10-phenanthroline have been obtained, such as (pydaH₂)(phendc) (pydaH₂ is protonated 2,6-pyridinedianine; Moghimi *et al.*, 2003) and (phenH⁺)₂(pydc²⁻) (pydc²⁻ is the pyridinedicarboxylic acid dianion; Moghimi *et al.*, 2005). We report here the synthesis and characterization of the title complex, (I).



Vigorous reaction of 1,10-phenanthroline with $HClO_4$ in water resulted in the proton-transfer compound (phenH⁺)(ClO₄⁻) (Fig. 1). In the compound, phen acts as proton acceptor and $HClO_4$ acts as proton donor (Etter, 1990). Selected bond lengths and angles are shown in Table 1.

Hydrogen bonds play a very important role in the formation of proton-transfer compounds. There are two kinds of hydrogen bond in the title complex. One is formed between atom O2 of ClO_4^- and the protonated phen (N1-H1B), and the other is formed within 1,10-phen (Table 2). A onedimensional chain along the *a* axis results from the N-H···O hydrogen bond (Fig. 2). In the crystal structure, the monoprotonated phen cations are parallel to each other, showing π - π interactions with a centroid–centroid distance of 3.37 (1) Å.

Experimental

All chemicals were obtained from commercial sources and used without further purification. 1,10-Phenanthroline (0.1 mmol, 0.0198 g) was dissolved in 10 ml water and HClO₄ (1 ml, 1 mol 1^{-1}) was added with stirring. The resulting solution was filtered after 0.5 h, then upon slow evaporation of the filtrate at room temperature, pink crystals of (I) were obtained after three weeks. Initial and final pH values of the solution were 1.6 and 1.0, respectively.

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Received 18 January 2006 Accepted 9 June 2006

Crystal data

 $C_{12}H_9N_2^+ \cdot CIO_4^ M_r = 280.66$ Triclinic, $P\overline{1}$ a = 5.8054 (12) Å b = 9.778 (2) Å c = 10.628 (2) Å $\alpha = 93.16$ (3)° $\beta = 101.41$ (3)° $\gamma = 98.08$ (3)°

Data collection

Bruker SMART CCD area-detector diffractometer φ and ω scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996) $T_{\rm min} = 0.900, T_{\rm max} = 0.938$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.040$ $wR(F^2) = 0.128$ S = 1.012767 reflections 177 parameters H atoms treated by a mixture of independent and constrained refinement

Table 1

Selected geometric parameters (Å, °).

| Cl1-O3 | 1.4230 (15) | N2-C5 | 1.325 (2) |
|---------------|-------------|---------------|-------------|
| Cl1-O4 | 1.4244 (14) | N2-C12 | 1.359 (2) |
| Cl1-O2 | 1.4251 (18) | N1-C2 | 1.326 (2) |
| Cl1-O1 | 1.4249 (19) | N1-C11 | 1.357 (2) |
| 03 - C11 - O4 | 110.13 (10) | O2 - C11 - O1 | 110.53 (15) |
| O3-Cl1-O2 | 109.38 (11) | C5-N2-C12 | 116.31 (16) |
| O4-Cl1-O2 | 109.44 (12) | C2-N1-C11 | 123.35 (16) |
| O3-Cl1-O1 | 108.78 (12) | C2-N1-H1A | 123.4 (15) |
| O4-Cl1-O1 | 108.57 (11) | C11-N1-H1A | 113.3 (15) |

V = 583.4 (2) Å³

 $D_x = 1.598 \text{ Mg m}^{-3}$

 $0.30 \times 0.25 \times 0.20$ mm

5890 measured reflections

2767 independent reflections

2345 reflections with $I > 2\sigma I$

 $w = 1/[\sigma^2(F_0^2) + (0.0799P)^2]$

where $P = (F_o^2 + 2F_c^2)/3$

+ 0.1598P]

 $(\Delta/\sigma)_{\rm max} < 0.001$ $\Delta \rho_{\rm max} = 0.38 \text{ e } \text{\AA}^{-3}$

 $\Delta \rho_{\rm min} = -0.43 \ {\rm e} \ {\rm \AA}^{-3}$

Mo $K\alpha$ radiation $\mu = 0.34 \text{ mm}^{-1}$

T = 288 (2) K

Block, pink

 $\begin{array}{l} R_{\rm int}=0.016\\ \theta_{\rm max}=28.3^\circ\end{array}$

Z = 2

Table 2

Hydrogen-bond geometry (Å, $^\circ).$

| $D - H \cdots A$ | D-H | $H \cdot \cdot \cdot A$ | $D \cdots A$ | $D - \mathbf{H} \cdots A$ |
|---------------------------------|----------|-------------------------|--------------|---------------------------|
| $N1-H1A\cdots O2^{i}$ | 0.84 (2) | 2.32 (2) | 2.944 (2) | 132 (2) |
| $N1 - H1A \cdot \cdot \cdot N2$ | 0.84 (2) | 2.33 (2) | 2.741 (2) | 111 (2) |

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

H atoms on the C atoms were located in a difference Fourier map and refined as riding on their parent atoms, with C-H = 0.93 Å and $U_{iso}(H) = 1.2U_{eq}(C)$. H atoms bonded to N atoms were located in a difference Fourier map and refined with the N-H distance restrained to 0.84 (2) Å.



Figure 1

The asymmetric unit of the title complex, showing the atom-labelling scheme and 30% probability displacement ellipsoids.





View along the c axis of the packing, with hydrogen bonds shown as dashed lines.

Data collection: *SMART* (Bruker, 2002); cell refinement: *SAINT* (Bruker, 2002); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997*a*); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997*a*); molecular graphics: *SHELXTL* (Sheldrick, 1997*b*); software used to prepare material for publication: *SHELXTL*.

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