Acta Crystallographica Section C Crystal Structure Communications ISSN 0108-2701

# Two hydration products of 3,4,5,6tetrachloro-*N*-(methyl-2-pyridyl)phthalmic acids

Paul G. Waddell,<sup>a,b</sup> Jeremy O. S. Hulse<sup>a</sup> and Jacqueline M. Cole<sup>a,b</sup>\*

<sup>a</sup>Cavendish Laboratory, University of Cambridge, J. J. Thomson Avenue, Cambridge CB3 0HE, England, and <sup>b</sup>Department of Chemistry, University of New Brunswick, Fredericton NB, E3B 5A3, Canada Correspondence e-mail: jmc61@cam.ac.uk

Received 26 April 2011 Accepted 9 June 2011 Online 23 June 2011

In 2-amino-6-methylpyridin-1-ium 2-carboxy-3,4,5,6-tetrachlorobenzoate,  $C_6H_9N_2^+ \cdot C_8HCl_4O_4^-$ , there are two perpendicular chains of hydrogen-bonded ions, one arising from the interaction between 2-carboxy-3,4,5,6-tetrachlorobenzoate ions and the other from the interaction between the 2-amino-6methylpyridin-1-ium and 2-carboxy-3,4,5,6-tetrachlorobenzoate ions. These chains combine to form a two-dimensional network of hydrogen-bonded ions. Cocrystals of bis(2-amino-3methylpyridin-1-ium) 3,4,5,6-tetrachlorophthalate–3,4,5,6tetrachlorophthalic acid (1/1),  $2C_6H_9N_2^+ \cdot C_8Cl_4O_4^{-2-} \cdot C_8H_2^ Cl_4O_4$ , form finite aggregates of hydrogen-bonded ions.  $\pi - \pi$ interactions are observed between 2-amino-3-methylpyridin-1-ium cations. Both structures exhibit the characteristic  $R_2^2(8)$ motif as a result of the hydrogen bonding between the 2-aminopyridinium and carboxylate units.

## Comment

N-(3-Methyl-2-pyridyl)-3,4,5,6-tetrachlorophthalmic acid and N-(6-methyl-2-pyridyl)-3,4,5,6-tetrachlorophthalmic acid are known to be pharamacologically active having been shown to exhibit a hypertensive effect in biological systems (Dolzhenko et al., 2003). In the context of this study, these materials are of interest for their potential as a UV-active dye for dye-sensitized solar-cell applications. Heating these compounds to 333 K in hydrated methanol produces crystals of two salts, namely 2-amino-6-methylpyridin-1-ium 2-carboxy-3,4,5,6tetrachlorobenzoate, (I) (Fig. 1), and bis(2-amino-3-methylpyridin-1-ium) 3,4,5,6-tetrachlorophthalate-3,4,5,6-tetrachlorophthalic acid (1/1), (II) (Fig. 2). These salts are the result of the reaction of the starting material with water present in the methanol solution and the equilibrium that exists between amides and water and the corresponding amines and carboxylic acids. In (I) and (II), protonation of the pyridyl N atom results in pyridinium salts stabilized by imino resonance.

Interestingly, as 3,4,5,6-tetrachlorophthalic acid cocrystallizes with 2-amino-3-methylpyridin-1-ium 3,4,5,6-tetrachlorophthalate all the products of this reaction are represented stoichiometrically in the crystal structure.



The molecular geometry of the 2-amino-6-methylpyridin-1ium cation in the structure of (I) (Table 1) can be compared with that of the nonhalogenated 2-amino-6-methylpyridinium 2-formylbenzoate monohydrate (Büyükgüngör & Odabaşoğlu, 2006). This geometry is similar in both compounds, with the characteristic bond-length alternation within the pyridyl ring, which demonstrates the imino resonance stabilizing the positive charge (Fig. 3) (Zhi *et al.*, 2002). The bond geometry of the aromatic ring in the 2-carboxy-3,4,5,6-tetrachlorobenzoate anion in (I) resembles more closely that of the hemihydrate of 3,4,5,6-tetrachlorophthalic acid (Ito *et al.*, 1975) than that of the 2-carboxy-3,4,5,6-tetrachlorobenzoate in a similar salt, 2-methyl-5-ethylpyridinium 3,4,5,6-tetrachlorophthalate (Galloy *et al.*, 1976). This indicates a contribution from the neutral canonical form, which is also observed







Figure 2

The structure of the asymmetric unit of (II), with atomic displacement ellipsoids drawn at the 50% probability level.



**Figure 3** The resonance exhibited by (I).

in the carboxylate group, where O3-C8 is observed to be shorter than O4-C8. The bond distances in the aromatic ring of 2-carboxy-3,4,5,6-tetrachlorobenzoate in (I) range from 1.374 (9) to 1.403 (9) Å.

Compound (I) forms continuous sheets of hydrogenbonded ions parallel to (010) (Table 2). These sheets contain the characteristic rings having graph set  $R_2^2(8)$  (Etter, 1990; Bernstein et al., 1995) with the amine and pyridinium N atoms acting as donors and the two carboxylate O atoms acting as acceptors (N1-H1N···O4 and N2-H2B···O3), as is well documented in this type of compound (Quah et al., 2010; Hemamalini & Fun, 2010). These rings are linked by chain motifs to form the sheets. The 2-carboxy-3,4,5,6-tetrachlorobenzoate anions form chains parallel to the [001] direction through  $O1 - H1 \cdots O4^{ii}$  interactions to give a graph-set motif of C(7) (symmetry codes as in Table 2). The 2-amino-6methylpyridin-1-ium cations link via the anions forming chains with the graph sets  $C_2^2(9)$  (through N2-H2B···O3 and N2- $H2A \cdots O2^{i}$ ) and  $C_{2}^{2}(11)$  (through N1-H1N···O4 and N2-H2A···O2<sup>i</sup>), giving an overall  $C_2^2(9)C_2^2(11)[R_2^2(8)]$  chain of rings parallel to the [100] direction (Fig. 5).

The aromatic rings in the 2-amino-3-methylpyridin-1-ium cations of compound (II) exhibit similar geometry to those in (I) with regard to the bond distances (Table 3). The distances in the benzene rings of the 3,4,5,6-tetrachlorophthalate dianion and the 3,4,5,6-tetrachlorophthalic acid molecule are in the ranges 1.387 (3)–1.403 (2) and 1.389 (3)–1.399 (3) Å,





View of the C(7) hydrogen-bonding motif in the [001] direction in (I). H atoms not involved in hydrogen bonding (dashed lines) have been omitted.





View of the  $C_2^2(9)C_2^2(11)[R_2^2(8)]$  chain of rings along [100] in (I). H atoms not involved in hydrogen bonding (dashed lines) have been omitted.

respectively, altogether more consistent than the bond distances of the 2-carboxy-3,4,5,6-tetrachlorobenzoate anion in (I) or the hemihydrate of 3,4,5,6-tetrachlorophthalic acid. This similarity in bond geometry between the dianion and the neutral acid in (II) strengthens the argument for the contribution of the neutral canonical forms in these compounds. As



Figure 6

Stereoview of the finite hydrogen-bonding network in the structure of (II). H atoms not involved in hydrogen bonding (dashed lines) have been omitted.

is observed in similar structures (Ni et al., 2007; Zhi et al., 2002), there are  $\pi - \pi$  interactions between 2-amino-3-methylpyridin-1-ium cations; the dihedral angle between the two pyridinium rings in the selected asymmetric unit is only  $3.8 (2)^{\circ}$  and the corresponding centroid–centroid separation is 3.834 (2) Å.

Unlike the two-dimensional network observed in (I), the hydrogen-bonded system in (II) consists of a finite array of four 2-amino-3-methylpyridin-1-ium cations, two 3,4,5,6tetrachlorophthalate dianions and two molecules of 3,4,5,6tetrachlorophthalic acid (Fig. 6 and Table 4). The  $R_2^2(8)$  ring motifs formed between the pyridinium and phthalate ions are once again present, with each phthalate dianion forming two such rings parallel to each other because of the  $\pi$ - $\pi$  interactions between the pyridinium cations, *i.e.* through N2- $H2A \cdots O1$  and  $N1 - H1N \cdots O2$  for one ring, and  $N4 - H4A \cdots$ O3 and N3-H3N $\cdots$ O4 for the other. In addition to this, each of these rings is connected to an adjacent  $R_2^2(8)$  ring through N2-H2B···O3<sup>i</sup> and N4-H4B···O1<sup>i</sup> forming an  $[R_2^2(8)$ - $R_4^2(8)R_2^2(8)$ ] motif within an outer  $R_4^4(16)$  ring (symmetry code is as in Table 4). Further motifs are observed when considering that there are two parallel  $[R_2^2(8)R_4^2(8)R_2^2(8)]$  motifs linked by the phthalate dianions; this gives rise to rings with graph sets  $R_4^4(18)$  and  $R_4^4(22)$ .

The 3,4,5,6-tetrachlorophthalic acid molecules and 3,4,5,6tetrachlorophthalate dianions are also connected by hydrogen bonds with the acid protons donating to carboxylate O-atom acceptors through O5-H5O···O2 and O7-H7O···O4 to create  $R_2^2(14)$  motifs. Though there is no direct hydrogen bonding between the acid molecules and the pyridinium cations, rings with the graph set  $R_8^6(34)$  are formed between acid molecules through the  $[R_2^2(8)R_4^2(8)R_2^2(8)]$  motif.

# **Experimental**

N-(3-Methyl-2-pyridyl)-3,4,5,6-tetrachlorophthalmic acid (10 mg, 0.026 mmol) and N-(6-methyl-2-pyridyl)-3,4,5,6-tetrachlorophthalmic acid (10 mg, 0.026 mmol) were heated to 333 K in hydrated methanol (5 ml) until a clear solution was obtained. Colourless platelike crystals of (I) were grown upon cooling to room temperature and colourless prism-like crystals of (II) grew after the solution was allowed to stand for one week.

## Compound (I)

#### Crvstal data

| $C_6H_9N_2^+ \cdot C_8HCl_4O_4^-$ |
|-----------------------------------|
| $M_r = 412.04$                    |
| Orthorhombic, Pca2 <sub>1</sub>   |
| a = 9.441 (14)  Å                 |
| b = 12.56 (2) Å                   |
| c = 13.69 (2) Å                   |

## Data collection

Rigaku Saturn724+ (2  $\times$  2 bin mode) diffractometer Absorption correction: multi-scan (ABSCOR; Higashi, 1995)  $T_{\min} = 0.866, T_{\max} = 0.956$ 

## Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.057$  $wR(F^2) = 0.127$ S = 1.102495 reflections 218 parameters 1 restraint

# Compound (II)

## Crystal data

| $2C_{6}H_{9}N_{2}^{+} \cdot C_{8}Cl_{4}O_{4}^{2-} \cdot C_{8}H_{2}Cl_{4}O_{4}$ |  |
|--|--|
| $M_r = 824.08$   |  |
| Friclinic, P1  |  |
| a = 8.6972 (17)  Å   |  |
| p = 13.762 (3) Å   |  |
| c = 15.381 (3)  Å  |  |
| $\alpha = 69.388 \ (9)^{\circ}$  |  |
| $\beta = 75.342 \ (10)^{\circ}$  |  |

#### Data collection

```
Rigaku Saturn724+ (2 \times 2 \text{ bin})
  mode) diffractometer
Absorption correction: multi-scan
  (ABSCOR; Higashi, 1995)
  T_{\rm min}=0.881,\;T_{\rm max}=0.914
```

## Table 1

Selected geometric parameters (Å, °) for (I).

| C7-O2    | 1.205 (7) | C9-N1     | 1.361 (9)  |
|----------|-----------|-----------|------------|
| C7-O1    | 1.413 (8) | C9-N2     | 1.402 (10) |
| C8-O3    | 1.248 (8) | C13-N1    | 1.427 (10) |
| C8-O4    | 1.344 (8) |           |            |
| O2-C7-O1 | 127.0 (6) | N1-C9-N2  | 124.9 (6)  |
| O3-C8-O4 | 134.0 (5) | C9-N1-C13 | 129.2 (5)  |

Table 2

Hydrogen-bond geometry (Å, °) for (I).

| $D - H \cdots A$       | $D-{\rm H}$ | $H \cdot \cdot \cdot A$ | $D \cdots A$ | $D - \mathbf{H} \cdots A$ |
|------------------------|-------------|-------------------------|--------------|---------------------------|
| $N2-H2B\cdots O3$      | 0.86        | 1.81                    | 2.666 (8)    | 173                       |
| $N2-H2A\cdots O2^{i}$  | 0.86        | 2.11                    | 2.931 (8)    | 159                       |
| $N1 - H1N \cdots O4$   | 0.86        | 1.71                    | 2.559 (7)    | 171                       |
| $O1-H1O\cdots O4^{ii}$ | 0.82        | 1.83                    | 2.604 (7)    | 158                       |
|                        |             |                         |              |                           |

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

V = 1623 (4) Å<sup>3</sup> Z = 4Mo  $K\alpha$  radiation  $\mu = 0.75 \text{ mm}^-$ T = 120 K $0.33 \times 0.16 \times 0.06 \text{ mm}$ 

5083 measured reflections 2495 independent reflections 2302 reflections with  $I > 2\sigma(I)$  $R_{\rm int} = 0.048$ 

H-atom parameters constrained  $\Delta \rho_{\rm max} = 0.44 \ {\rm e} \ {\rm \AA}^{-3}$  $\Delta \rho_{\rm min} = -0.50 \text{ e } \text{\AA}^{-3}$ Absolute structure: Flack (1983) Flack parameter: -0.31 (13)

 $\gamma = 72.618 \ (1)^{\circ}$ V = 1621.7 (6) Å<sup>3</sup> Z = 2Mo  $K\alpha$  radiation  $\mu = 0.75 \text{ mm}^{-1}$ T = 120 K $0.41 \times 0.14 \times 0.12 \text{ mm}$ 

```
9810 measured reflections
5964 independent reflections
5551 reflections with I > 2\sigma(I)
R_{\rm int} = 0.033
```

## Table 3

|          |           |              | 0     |         |      |    |
|----------|-----------|--------------|-------|---------|------|----|
| Selected | geometric | parameters ( | (Å, ° | ) for ( | (II) | ). |

| C7-O1     | 1.235 (3)   | C16-O7     | 1.301 (3)   |
|-----------|-------------|------------|-------------|
| C7-O2     | 1.272 (3)   | C17-N2     | 1.330 (3)   |
| C8-O3     | 1.245 (3)   | C17-N1     | 1.346 (3)   |
| C8-O4     | 1.258 (3)   | C21-N1     | 1.358 (3)   |
| C15-O6    | 1.211 (3)   | C23-N4     | 1.333 (3)   |
| C15-O5    | 1.303 (3)   | C23-N3     | 1.349 (3)   |
| C16-O8    | 1.218 (3)   | C27-N3     | 1.363 (3)   |
| 01-C7-O2  | 125.39 (19) | N2-C17-N1  | 118.96 (19) |
| 03-C8-O4  | 124.88 (19) | C17-N1-C21 | 123.66 (18) |
| O6-C15-O5 | 126.7 (2)   | N4-C23-N3  | 118.35 (18) |
| O8-C16-O7 | 126.1 (2)   | C23-N3-C27 | 122.89 (18) |
|           |             |            |             |

#### Table 4

Hydrogen-bond geometry (Å, °) for (II).

| $D - H \cdot \cdot \cdot A$     | D-H  | $H \cdot \cdot \cdot A$ | $D \cdots A$ | $D - H \cdot \cdot \cdot A$ |
|---------------------------------|------|-------------------------|--------------|-----------------------------|
| $05 - H50 \cdots 02$            | 0.84 | 1.80                    | 2.558 (2)    | 149                         |
| O7−H7O···O4                     | 0.84 | 1.83                    | 2.593 (2)    | 151                         |
| $N1 - H1N \cdot \cdot \cdot O2$ | 0.88 | 1.83                    | 2.708 (2)    | 173                         |
| $N2-H2A\cdots O1$               | 0.88 | 2.07                    | 2.943 (2)    | 171                         |
| N3-H3N···O4                     | 0.88 | 2.01                    | 2.891 (2)    | 175                         |
| $N4-H4A\cdots O3$               | 0.88 | 1.93                    | 2.805 (2)    | 175                         |
| $N2-H2B\cdots O3^{i}$           | 0.88 | 1.99                    | 2.823 (2)    | 157                         |
| $N4-H4B\cdotsO1^{i}$            | 0.88 | 2.07                    | 2.892 (2)    | 156                         |

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

#### Refinement

| $R[F^2 > 2\sigma(F^2)] = 0.036$ | 435 parameters   |
|---------------------------------|--|
| $wR(F^2) = 0.090$               | H-atom parameters constrained                              |
| S = 1.07                        | $\Delta \rho_{\rm max} = 0.43 \ {\rm e} \ {\rm \AA}^{-3}$  |
| 5964 reflections                | $\Delta \rho_{\rm min} = -0.34 \text{ e } \text{\AA}^{-3}$ |

H atoms were positioned geometrically and refined as riding on their parent atoms, with C-H = 0.95 Å and  $U_{iso}(H) = 1.2U_{eq}(C)$ , and N-H = 0.88 Å and  $U_{iso}(H) = 1.2U_{eq}(N)$ . Hydroxy and methyl H atoms were modelled in a similar fashion, with O-H = 0.84 Å and  $U_{iso}(H) = 1.5U_{eq}(C)$ , and C-H = 0.98 Å and  $U_{iso}(H) = 1.5U_{eq}(N)$ . The most disagreeable reflections were omitted and those exhibiting a  $\Delta(F^2)$  value greater than 5 s.u. were removed; 5 from (I) and 31 from (II). The refinement was further improved by restricting the reflections considered to those with  $\theta \le 25.68^{\circ}$ . The Flack parameter for (I) gives the expected values for a correct absolute structure within 3 s.u. Nonetheless since the s.u. is moderate, the inverted structure was tested. This yielded a Flack parameter of x = 1.21 (13) by the 'hole-in-one' method and of x = 1.34 (13) using TWIN/BASF, giving us confidence that we have presented the correct absolute structure with respect to the polar-axis direction. These checks were particularly important given that the precision of the Flack *x* parameter is poor owing to a low Friedel coverage of 60%. Refinement for (II) was limited to those reflections with  $\theta < 25.68^{\circ}$  reducing the number of missing data; however, a number of missing data remain (201 reflections between  $\theta_{\min}$  and  $\sin\theta/\lambda = 0.600$ ). Analysis of reciprocal-space plots reveal that these missing portions are fairly randomly dispersed which gives us confidence that this is not a systematic error. Moreover, the missing data were comprised of high-angle reflections that were just outside the reach of the data collection strategy.

For both compounds, data collection: *CrystalClear* (Rigaku, 2008); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

JMC thanks the Royal Society for a University Research Fellowship, the University of New Brunswick for the UNB Vice-Chancellor's Research Chair (JMC), and NSERC for Discovery Grant No. 355708 (for PGW).

Supplementary data for this paper are available from the IUCr electronic archives (Reference: GD3389). Services for accessing these data are described at the back of the journal.

## References

- Bernstein, J., Davis, R. E., Shimoni, L. & Chang, N.-L. (1995). Angew. Chem. Int. Ed. Engl. 34, 1555–1573.
- Büyükgüngör, O. & Odabaşoğlu, M. (2006). Acta Cryst. E62, o2749-o2750.
- Dolzhenko, A. V., Syropyatov, B. Ya., Koz'minykh, V. O., Kolotova, N. V., Zakhmatov, A. V. & Borodin, A. Yu. (2003). *Pharm. Chem. J.* **37**, 407–408.
- Etter, M. C. (1990). Acc. Chem. Res. 23, 121-126.
- Farrugia, L. J. (1999). J. Appl. Cryst. 32, 837-838.
- Flack, H. D. (1983). Acta Cryst. A39, 876-881.
- Galloy, J., Putzeys, J.-P., Germain, G. & Van Meerssche, M. (1976). *Acta Cryst.* B**32**, 2718–2720.
- Hemamalini, M. & Fun, H.-K. (2010). Acta Cryst. E66, o2192-o2193.
- Higashi, T. (1995). ABSCOR. Rigaku Corporation, Tokyo, Japan.
- Ito, K., Moriya, K., Kashino, S. & Haisa, M. (1975). Bull. Chem. Soc. Jpn, 48, 3078–3084.

Ni, S.-F., Feng, W.-J., Guo, H. & Jin, Z.-M. (2007). Acta Cryst. E63, 03866. Quah, C. K., Hemamalini, M. & Fun, H.-K. (2010). Acta Cryst. E66, 01932.

Rigaku (2008). CrystalClear. Rigaku Corporation, Tokyo, Japan.

- Sheldrick, G. M. (2008). Acta Cryst. A64, 112-122.
- Zhi, M.-J., Yuan, J.-P., Mao, L.-H. & Liang, S. (2002). J. Chem. Crystallogr. 31, 191–195.