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#### Key indicators

Single-crystal X-ray study T = 300 KMean  $\sigma(C-C) = 0.004 \text{ Å}$  R factor = 0.043 wR factor = 0.159 Data-to-parameter ratio = 18.1

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

# Pyrrolidinium chloranilate

In the title compound,  $2C_4H_{10}N^+ \cdot C_6Cl_2O_4^{2-}$ , the pyrrolidinium and chloranilate ions are connected by N-H···O hydrogen bonds and arranged in an alternating manner to give an infinite molecular tape.

#### Comment

The title compound, (I), was investigated as part of a study on  $D-H\cdots A$  hydrogen bonding (D = N, O, or C; A = N, O, Cl) in chloranilic acid-amine 1:1 and 1:2 systems (Ishida & Kashino, 1999*a*,*b*,*c*, 2000, 2001, 2002; Ishida *et al.*, 2001; Fukunaga & Ishida, 2003; Fukunaga *et al.*, 2003). Crystal structures of the chloranilic acid-secondary amine 1:1 system have been analyzed for the morpholinium, diethylammonium and piperidinium salts (Ishida & Kashino, 1999*c*, 2000; Fukunaga & Ishida, 2003), but the structure of the chloranilic acid-secondary amine 1:2 system has not yet been reported. In the present study, we prepared the 1:2 salt, (I), using pyrrolidine as a counter-base and determined its crystal structure.



In (I), the asymmetric unit is composed of  $2C_4H_8NH_2^+ \cdot C_6O_4Cl_2^{2-}$ ; an acid-base interaction involving a proton transfer is observed between the chloranilic acid and



#### Figure 1

 $O\overline{RTEP}$ -3 (Farrugia, 1997) drawing of (I), with the atom-labeling scheme. Displacement ellipsoids of non-H atoms are drawn at the 50% probability level. N-H···O hydrogen bonds are indicated by dashed lines [symmetry code: (i) x - 1, y, z].

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pyrrolidine (Fig. 1). The chloranilate ion shows a characteristic structure having four short C–C bonds [1.382 (3)– 1.416 (3) Å] and two extremely long C–C bonds [1.540 (3) and 1.544 (3) Å], which is explainable in terms of the double  $\pi$  system of the anion (Andersen, 1967; Benchekroun & Savariault, 1995). The chloranilate and pyrrolidinium ions are connected by N–H···O hydrogen bonds (Table 2) and arranged in an alternating manner to give a molecular tape running along the *a* axis (Fig. 2). The tapes are stacked along the *b* axis, such that the chloranilate ions related by an inversion center are stacked through  $\pi$ – $\pi$  interactions, with an interplanar spacing of 3.346 Å and a centroid offset of 3.209 Å ( $Cg \cdots Cg = 4.636$  Å; Cg is the centroid of each anion).

The pyrrolidinium cation, which is isoelectronic with cyclopentane, is expected to have ring flexibility as high as cyclopentane. Molecular-orbital calculations for the isolated cation in the gas phase showed that the twist form ( $C_2$  form) is the stable conformation and that the envelope on N atom form ( $C_s$  form) is a saddle point of the first order (Ishida, 2000). Other twist conformations and envelope on C-atom conformations have not been optimized as stationary points. The difference in the electronic energy between the  $C_2$  and  $C_s$  forms calculated at the HF/6–31G(d,p) level is 2.0 kJ mol<sup>-1</sup>, which is small compared to the energies of intermolecular interactions in the solid. In the present salt, the N1–C7–C8–C9–C10 cation ring is close to the  $C_2$  form, while the N2–C11–C12–C13–C14 ring is close to the envelope on C13 form.

### **Experimental**

Crystals were obtained by slow evaporation of an acetonitrile solution of pyrrolidine and chloranilic acid in a 2:1 molar ratio.

#### Crystal data

 $2C_4H_{10}N^+ \cdot C_6Cl_2O_4^{2-}$   $M_r = 351.23$ Triclinic,  $P\overline{1}$  a = 9.337 (3) Å b = 9.601 (4) Å c = 9.947 (5) Å  $\alpha = 68.18 (3)^{\circ}$   $\beta = 84.25 (3)^{\circ}$   $\gamma = 71.52 (2)^{\circ}$   $V = 785.0 (6) \text{ Å}^3$ 

#### Data collection

Rigaku AFC-5*R* diffractometer  $\omega$ -2 $\theta$  scans Absorption correction:  $\psi$  scan (North *et al.*, 1968)  $T_{\min} = 0.91, T_{\max} = 0.94$ 3821 measured reflections 3606 independent reflections 2522 reflections with  $I > 2\sigma(I)$ 

#### Refinement

Refinement on  $F^2$   $R[F^2 > 2\sigma(F^2)] = 0.043$   $wR(F^2) = 0.159$  S = 1.093606 reflections 199 parameters Z = 2  $D_x = 1.486 \text{ Mg m}^{-3}$ Mo K $\alpha$  radiation Cell parameters from 25 reflections  $\theta = 11.2 - 11.8^{\circ}$   $\mu = 0.43 \text{ mm}^{-1}$ T = 300 K Prism, dark red  $0.40 \times 0.20 \times 0.15 \text{ mm}$ 

$$\begin{split} R_{\rm int} &= 0.014 \\ \theta_{\rm max} &= 27.5^{\circ} \\ h &= -12 \rightarrow 11 \\ k &= -12 \rightarrow 0 \\ l &= -12 \rightarrow 11 \\ 3 \text{ standard reflections} \\ \text{every 97 reflections} \\ \text{intensity decay: } 1.7\% \end{split}$$

H-atom parameters constrained  $w = 1/[\sigma^2(F_o^2) + (0.09P)^2]$ where  $P = (F_o^2 + 2F_c^2)/3$   $(\Delta/\sigma)_{max} = 0.001$   $\Delta\rho_{max} = 0.57 \text{ e} \text{ Å}^{-3}$  $\Delta\rho_{min} = -0.36 \text{ e} \text{ Å}^{-3}$ 



#### Figure 2

Packing diagram showing a molecular tape formed *via*  $N-H\cdots O$  hydrogen bonds (shown as dashed lines). H atoms except those attached to N atoms have been omitted for clarity.

## Table 1

Selected geometric parameters (Å, °).

Cl1-C2	1.746 (3)	C1-C6	1.544 (3)
Cl2-C5	1.742 (2)	C2-C3	1.382 (3)
O1-C1	1.230 (3)	C3-C4	1.540 (3)
O2-C3	1.256 (3)	C4-C5	1.415 (3)
O3-C4	1.238 (3)	C5-C6	1.384 (3)
O4-C6	1.259 (3)	C7-C8	1.487 (4)
N1-C7	1.495 (3)	C8-C9	1.498 (5)
N1-C10	1.495 (3)	C9-C10	1.502 (5)
N2-C11	1.497 (3)	C11-C12	1.482 (4)
N2-C14	1.499 (3)	C12-C13	1.510 (5)
C1-C2	1.416 (3)	C13-C14	1.516 (4)
C1-C2-Cl1	117.3 (2)	O2-C3-C4	116.1 (2)
C4-C5-Cl2	117.4 (2)	O3-C4-C3	116.9 (2)
O1-C1-C6	117.1 (2)	O4-C6-C1	116.1 (2)

Table 2		
Hydrogen-bonding geometr	y (Å,	°).

$D-\mathrm{H}\cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
N1-H1···O2	0.89	1.91	2.791 (3)	169
$N1 - H2 \cdot \cdot \cdot O1^{i}$	0.89	2.39	2.962 (3)	122
$N1 - H2 \cdot \cdot \cdot O4^{i}$	0.89	1.98	2.824 (3)	158
$N2-H11\cdots O2$	0.89	2.12	2.841 (3)	138
$N2-H11\cdots O3$	0.89	2.26	2.930 (2)	132
$N2-H12\cdots O4^{i}$	0.89	1.96	2.783 (3)	153

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

H atoms were treated as riding atoms, with C-H = 0.97 Å and N-H = 0.89 Å.  $U_{iso}(H)$  values were set at  $1.5U_{eq}$  of the parent atom.

Data collection: *MSC/AFC Diffractometer Control Software* (Molecular Structure Corporation, 1990); cell refinement: *MSC/AFC Diffractometer Control Software*; data reduction: *TEXSAN for Windows* (Molecular Structure Corporation, 1997–1999); program(s) used to solve structure: *SIR*92 (Altomare *et al.*, 1994); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *ORTEP3* (Farrugia, 1997); software used to prepare material for publication: *TEXSAN for Windows*.

X-ray measurements were made at the X-ray Laboratory of Okayama University.

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