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Key indicators

Single-crystal X-ray study T = 300 KMean σ (C–C) = 0.004 Å R factor = 0.048 wR factor = 0.134 Data-to-parameter ratio = 15.8

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

Benzimidazolium 2-chloro-4-nitrobenzoate

In the title compound, $C_7H_7N_2^+ \cdot C_7H_3CINO_4^-$, the cations and anions are connected by $N-H \cdot \cdot \cdot O$ hydrogen bonds to afford a 2₁ helical chain.

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Comment

The title compound, (I), was investigated as part of a study on $D - H \cdot \cdot \cdot A$ hydrogen bonding (D = N, O or C; A = N, O or Cl) in chloro- and nitro-substituted benzoic acid-amine systems (Ishida et al., 2001a,b,c,d,e). In the crystal, the cations and anions are held together by $N-H \cdots O$ hydrogen bonds (Table 2) and a weak C-H··· π interaction [H10···Cg 2.93 (3) Å, $C13 \cdots Cg \ 3.679 \ (3) \ \text{\AA}, \ C13 - H10 \cdots Cg \ 143.5 \ (19)^{\circ}, \text{ where } Cg$ denotes the centroid of the benzene ring C1-C6] (Fig. 1) to afford a 2_1 helical chain running along the *b* axis (Fig. 2). A similar helical chain is observed in imidazolium 2-chloro-4nitrobenzoate, giving a chiral crystal (Ishida et al., 2001e). The present salt, however, crystallizes in the centrosymmetric space group C2/c. Neighboring helical chains related by an inversion center are connected through C-H···O hydrogen bond (Fig. 2, Table 2), and those related by a twofold rotation axis are linked by a $\pi - \pi$ stacking interaction between the aromatic rings C9-C14 of the benzimidazolium ion. The dihedral angle between the aromatic rings is $1.35 (11)^{\circ}$, and their interplanar separation and the centroid offset are 3.480 (2) and 0.769 (2) Å, respectively. The carboxyl group is twisted considerably out of the plane of the benzene ring, probably because of the strong N-H...O hydrogen bonds; the dihedral angle between them is 67.15 (13)°. This may cause a close contact of O1···N1ⁱⁱⁱ, 2.832 (3) Å [symmetry code: (iii) 1 - x, 1 - y, -z], between two chains running in antiparallel directions. The N···O distance [2.610 (3) Å] in the N2– H4...O2 hydrogen bond is significantly shorter than the average $N \cdots O$ distance of 2.878 (3) Å for an $N-H \cdots O = C$ hydrogen bond (Taylor et al., 1984).



Experimental

Crystals of (I) were obtained by slow evaporation from an acetonitrile solution of benzimidazole and 2-chloro-4-nitrobenzoic acid in a molar ratio of 1:1.

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Figure 1

ORTEP-3 (Farrugia, 1997) drawing of (I), with the atom-labeling. Displacement ellipsoids of non-H atoms are drawn at the 50% probability level. The N-H···O hydrogen bond and C-H··· π interaction are indicated by a dashed and a dotted line, respectively.

intensity decay: 0.8%

 $w = 1/[\sigma^2(F_o^2) + (0.0502P)^2]$

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

+ 1.4779P]

 $\Delta \rho_{\rm max} = 0.20 \ {\rm e} \ {\rm \AA}^{-3}$

 $\Delta \rho_{\rm min} = -0.26 \text{ e} \text{ Å}^{-3}$

 $(\Delta/\sigma)_{\rm max} = 0.01$

Crystal data

$C_7H_7N_2^+ \cdot C_7H_3CINO_4^-$	$D_x = 1.493 \text{ Mg m}^{-3}$
$M_r = 319.70$	Mo K α radiation
Monoclinic, $C2/c$	Cell parameters from 25
a = 11.120 (3) Å	reflections
b = 14.771 (3) Å	$\theta = 11.3 - 12.1^{\circ}$
c = 17.414(3) Å	$\mu = 0.29 \text{ mm}^{-1}$
$\beta = 95.963 \ (18)^{\circ}$	$T = 300 { m K}$
$V = 2844.8 (11) \text{ Å}^3$	Prismatic, colorless
Z = 8	0.50 \times 0.35 \times 0.25 mm
Data collection	
Rigaku AFC-5R diffractometer	$R_{\rm int} = 0.025$
ω –2 θ scans	$\theta_{\rm max} = 29.0^{\circ}$
Absorption correction: ψ scan	$h = -4 \rightarrow 14$
(North et al., 1968)	$k = -4 \rightarrow 19$
$T_{\min} = 0.876, \ T_{\max} = 0.930$	$l = -23 \rightarrow 23$
6914 measured reflections	3 standard reflections
3776 independent reflections	every 97 reflections
2040 reflections with $I > 2\sigma(I)$	intensity decay: 0.8%

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.048$ wR(F²) = 0.134 S = 1.003776 reflections 239 parameters All H-atom parameters refined

Table 1

Selected geometric parameters (Å).

Cl-C2	1.731 (2)	N2-C8	1.320 (3)
O1-C7	1.228 (3)	N2-C14	1.388 (3)
O2-C7	1.242 (3)	N3-C8	1.313 (4)
O3-N1	1.223 (3)	N3-C9	1.395 (3)
O4-N1	1.222 (3)	C1-C7	1.520 (3)
N1-C4	1.473 (3)		



Figure 2

Packing diagram showing two 21 helical chains running in antiparallel directions along the b axis. $N-H\cdots O$ and $C-H\cdots O$ hydrogen bonds indicated by dashed and dotted lines, respectively [symmetry codes are as in Table 2].

Table 2

Hydrogen-bonding geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$\begin{array}{c} N2 - H4 \cdots O2 \\ N3 - H6 \cdots O1^{i} \\ C8 - H5 \cdots O3^{ii} \end{array}$	1.09 (3) 0.85 (3) 0.98 (3)	1.54 (3) 1.84 (3) 2.54 (2)	2.610 (3) 2.654 (3) 3.395 (4)	165 (3) 160 (3) 146 (2)

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

H atoms were found in difference Fourier maps and refined isotropically. Refined distances: C-H = 0.88 (3) - 0.98 (3) Å and N-H = 0.86(3) - 1.09(3) Å.

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: SIR92 (Altomare et al. 1993); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP-3 (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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