

Piperidinium *p*-nitrobenzoateYoshimitsu Moritani,^{a†} Akihiko Takeda^b and Setsuo Kashino^{b*}^aGraduate School of Natural Science and Technology, Okayama University, Tsushima, Okayama 700-8530, Japan, and ^bDepartment of Chemistry, Faculty of Science, Okayama University, Tsushima, Okayama 700-8530, Japan

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Key indicators

Single-crystal X-ray study

T = 298 K

Mean $\sigma(\text{C}-\text{C}) = 0.005 \text{ \AA}$

R factor = 0.057

wR factor = 0.205

Data-to-parameter ratio = 14.6

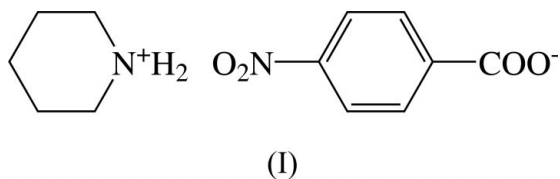
For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

A new type of hydrogen-bond pattern for piperidinium *p*-substituted benzoates is reported; this is found in piperidinium *p*-nitrobenzoate, $\text{C}_5\text{H}_{12}\text{N}^+\cdot\text{C}_7\text{H}_4\text{NO}_4^-$, (I). In the crystal of (I), the cations and anions are linked by $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds around a center of symmetry to form a cyclic dimer of the formula unit.

Comment

We report here a new type of hydrogen-bond pattern in the cyclic secondary amine-*p*-substituted benzoic acid (1/1) system. In piperidinium *p*-nitrobenzoate, (I), a centrosymmetric dimer of the formula unit is formed through $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds (Fig. 1 and Table 1).

In most crystalline salts formed between cyclic secondary amines and *p*-substituted benzoic acids, the cations and anions are arranged around a twofold screw axis to form $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds (Kashino *et al.*, 1972, 1978, 1981; Kashino, 1973). The same type of hydrogen bonding is also found in both pyrrolidinium *p*-nitrobenzoate and hexamethyleineiminium *p*-nitrobenzoate (Takeda, 1992). In a few cases, the cations and anions are arranged along a glide plane to form $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds (Kashino *et al.*, 1973, 1981).



The possibility of the formation of a cyclic dimer through the $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds was deduced based on the result of a molecular weight measurement in benzene solution, combined with a geometrical consideration of the hydrogen-bonded system (Kashino, 1973). However, the formation of a cyclic dimer has been found only in one of the dimorphs of hexamethyleineiminium *p*-bromobenzoate, in which the dimer is formed around a twofold axis (Kashino *et al.*, 1981). The present study established a centrosymmetric type as the fourth pattern of hydrogen bonding possible in salts of cyclic secondary amines and *p*-substituted benzoic acids.

Experimental

Stoichiometric amounts of piperidine and *p*-nitrobenzoic acid were dissolved in benzene at 340 K. After cooling the solution, crystals of (I) were grown by slow evaporation at room temperature.

Received 22 June 2001

Accepted 28 June 2001

Online 13 July 2001

Crystal data

$C_5H_{12}N^+ \cdot C_7H_4NO_4^-$
 $M_r = 252.27$
 Orthorhombic, *Pccn*
 $a = 8.949$ (2) Å
 $b = 22.451$ (5) Å
 $c = 12.712$ (4) Å
 $V = 2554.0$ (11) Å³
 $Z = 8$
 $D_x = 1.312$ Mg m⁻³

Mo $K\alpha$ radiation
 Cell parameters from 25 reflections
 $\theta = 10.6$ – 11.4°
 $\mu = 0.10$ mm⁻¹
 $T = 298$ K
 Prismatic, colorless
 $0.50 \times 0.30 \times 0.20$ mm

Data collection

Rigaku AFC-5R diffractometer
 ω - 2θ scans
 Absorption correction: ψ scan
 (North *et al.*, 1968)
 $T_{\min} = 0.952$, $T_{\max} = 0.980$
 2875 measured reflections
 2509 independent reflections
 1142 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.022$
 $\theta_{\text{max}} = 26.0^\circ$
 $h = 0 \rightarrow 11$
 $k = 0 \rightarrow 27$
 $l = 0 \rightarrow 15$
 3 standard reflections
 every 97 reflections
 intensity decay: 0.6%

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.057$
 $wR(F^2) = 0.205$
 $S = 0.91$
 2509 reflections
 172 parameters
 H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.1P)^2 + 1.1042P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.002$
 $\Delta\rho_{\text{max}} = 0.13$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.18$ e Å⁻³
 Extinction correction: *SHELXL97*
 Extinction coefficient: 0.0059 (14)

Table 1

Selected geometric parameters (Å, °).

O1–C7	1.253 (4)	C3–C4	1.377 (5)
O2–C7	1.250 (4)	C4–C5	1.379 (5)
O3–N1	1.219 (4)	C5–C6	1.378 (5)
O4–N1	1.216 (4)	N2–C8	1.478 (5)
N1–C4	1.476 (5)	N2–C12	1.489 (5)
C1–C6	1.386 (5)	C8–C9	1.510 (6)
C1–C2	1.387 (5)	C10–C11	1.502 (6)
C1–C7	1.513 (5)	C11–C12	1.500 (6)
C2–C3	1.378 (5)		
O4–N1–O3	123.6 (4)	C6–C5–C4	118.5 (3)
O4–N1–C4	118.5 (4)	C5–C6–C1	120.9 (3)
O3–N1–C4	117.9 (4)	O2–C7–O1	125.0 (4)
C6–C1–C2	119.4 (3)	O2–C7–C1	117.7 (3)
C6–C1–C7	120.5 (3)	O1–C7–C1	117.2 (3)
C2–C1–C7	120.1 (3)	C8–N2–C12	111.1 (3)
C3–C2–C1	120.3 (3)	N2–C8–C9	109.6 (4)
C4–C3–C2	119.1 (3)	C10–C9–C8	110.0 (4)
C3–C4–C5	121.8 (4)	C11–C10–C9	111.0 (3)
C3–C4–N1	119.2 (3)	C12–C11–C10	111.4 (4)
C5–C4–N1	119.0 (4)	N2–C12–C11	110.3 (3)

Table 2

Hydrogen-bonding geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
N2–H5 \cdots O1	0.94 (4)	1.78 (4)	2.697 (5)	165 (4)
N2–H6 \cdots O2 ⁱ	1.02 (5)	1.71 (5)	2.731 (4)	171 (4)

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

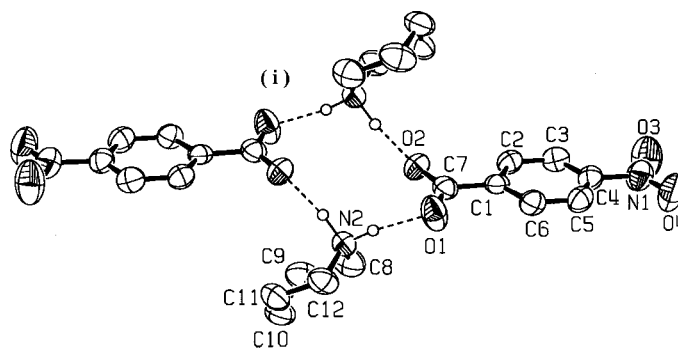


Figure 1

A molecular view of (I) showing the hydrogen-bonding pattern and the atomic numbering scheme. Displacement ellipsoids are drawn at the 50% probability level and H atoms involved in hydrogen bonds are shown as small spheres of arbitrary radii. Hydrogen bonds are indicated by dashed lines. [Symmetry code: (i) $1 - x, 1 - y, 1 - z$.]

All H atoms were found from a difference Fourier map. At the final stage of the least-squares refinement, all H atoms except those involved in hydrogen bonds were fixed at the ideal positions, and their isotropic displacement parameters were fixed to $1.2U_{\text{eq}}$ of the parent atoms.

Data collection: *MSC/AFC Diffractometer Control Software* (Molecular Structure Corporation, 1994); 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*.

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