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Mohd Mustaqim Rosli,^a Hoong-Kun Fun,^a* Ibrahim Abdul Razak,^a Pinanong Rodwatcharapiban^b and Suchada Chantrapromma^b*

^aX-ray Crystallography Unit, School of Physics, Universiti Sains Malaysia, 11800 USM, Penang, Malaysia, and ^bDepartment of Chemistry, Faculty of Science, Prince of Songkla University, Hat-Yai, Songkhla 90112, Thailand

Correspondence e-mail: hkfun@usm.my, suchada.c@psu.ac.th

Key indicators

Single-crystal X-ray study T = 100 KMean σ (C–C) = 0.002 Å R factor = 0.036 wR factor = 0.108 Data-to-parameter ratio = 11.5

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. In the title salt, $C_{10}H_{10}N^+ \cdot C_7H_3N_2O_6^-$, in addition to electrostatic interactions, the quinaldinium and 2,4-dinitrobenzoate ions are linked by N-H···O and C-H···O hydrogen bonds into a three-dimensional molecular network. The quinaldinium cations are stacked along the *a* axis with a centroid–centroid distance of 3.4662 (7) Å, indicating π - π interactions.

2-Methylquinolinium 2,4-dinitrobenzoate

Comment

Hydrogen bonding is a very important structural phenomenon and is still not fully understood, especially in phenol-benzoic acid systems. Interactions of organic acids with amine bases in the solid phase have been of interest due to their hydrogenbonding interactions (Coupar et al., 1997; Sobczyk et al., 2000) and their important role in biological systems (Blow, 1976). We have been studying hydrogen-bonding interactions in the adducts of phenols and carboxylic acids with amines (Usman et al., 2001; Chantrapromma et al., 2002a,b,c, 2004, 2006; Chantrapromma & Fun 2004; Fun et al., 2003; How et al., 2005; Rosli et al., 2006). 2,4-Dinitrobenzoic acid is an interesting acid since it readily forms the 2,4-dinitrobenzoate anion when cocrystallized with amine bases, by transferring a proton. This proton transfer leads to the formation of salt-like adducts consisting of 2,4-dinitrobenzoate anions and amine-based cations, in which the 2,4-dinitrobenzoate anion is expected to act as a multiple donor of hydrogen bonds via the nitro and carboxylate groups.



Our studies using 2,4-dinitrophenol and 3,5-dinitrobenzoic acid with some amines gave a number of proton-transfer compounds (Usman *et al.*, 2001; Chantrapromma *et al.*, 2002*a,b*, 2004; Fun *et al.*, 2003) and some of these undergo a phase transition (Fun *et al.*, 2003; How *et al.*, 2005). Owing to our interest in studying hydrogen bonding and in searching for further examples of phase transitions due to hydrogen bonding in the adducts of acids and amines (Fun *et al.*, 2003; How *et al.*, 2003; How *et al.*, 2003; How *et al.*, 2005), the title compound, (I), was prepared and we report its crystal structure here.

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Figure 1

The asymmetric unit of (I), showing 80% probability displacement ellipsoids and the atomic numbering scheme. The dashed line indicates a hydrogen bond.



Figure 2

A view of the hydrogen-bonded (dashed lines) chain in the crystal structure of (I).

The asymmetric unit of compound (I) (Fig. 1) contains a $C_{10}H_{10}N^+$ cation and a $C_7H_3N_2O_6^-$ anion. The 2,4-dinitrocarboxylic acid molecule has transferred a proton to the hetero-N atom of quinaldine. All bond distances and angles in (I) show normal values (Allen *et al.*, 1987). Selected bond distances and angles are given in Table 1.

The proton transfer from the carboxyl group results in a slight distortion of the functional groups of the 2,4-dinitrobenzoate ion. In this anion, the carboxylate and the two nitro groups are twisted away from the plane of the benzene ring, with torsion angles $O1-N1-C1-C2 = 167.17 (12)^{\circ}$, $O3-N2-C3-C4 = 63.09 (17)^{\circ}$ and C3-C4-C7-O5 =

32.66 (18)°. The dihedral angle between the planes of the adjacent carboxylate and nitro groups is 63.4 (2)°. The C4–C7 bond distance [1.5280 (18) Å] is relatively long compared with the corresponding distance in 3,5-dinitrobenzoic acid [1.5019 (10) Å; Rosli *et al.*, 2006]. The maximum deviation from planarity in the essentially planar quinaldinium cation (C8–C17/N3) is 0.029 (1) Å for atom C15. The dihedral angle between the plane of the benzene ring of the anion and the plane of the quinaldinium cation is 18.83 (5)°.

In the crystal structure of (I), in addition to electrostatic interactions, the quinaldinium and 2,4-dinitrobenzoate ions are linked by N3-H1N3···O5, C9-H9···O6 and C17-H17C···O5 hydrogen bonds, with the quinaldinium cation acting as a hydrogen-bond donor and the 2,4-dinitrobenzoate anion acting as a multiple hydrogen-bond acceptor. The C6-H6···O4ⁱ hydrogen bonds link the cation-anion pairs into a chain along the *b* axis (Fig. 2). These chains are interconnected through C12-H12···O1ⁱⁱ and C12-H12···O3ⁱⁱⁱ hydrogen bonds (symmetry codes are given in Table 2) into a threedimensional molecular network. The quinaldinium cations are stacked along the *a* axis with a centroid···centroid distance of 3.4662 (7) Å, indicating π - π interactions.

Experimental

Compound (I) was prepared by mixing equimolar amounts of quinaldine (0.07 ml, 0.5 mmol) and 2,4-dinitrobenzoic acid (0.11 g, 0.5 mmol) in ethanol (40 ml) with a few drops of water. The mixture was warmed until a clear solution was obtained. The solution was then filtered and left to evaporate slowly at ambient temperature. Pale-yellow block single crystals of (I) were obtained from the solution after a few days (m.p. 455–457 K).

Crystal data

$C_{10}H_{10}N^+ \cdot C_7H_3N_2O_6^-$	Z = 2
$M_r = 355.30$	$D_x = 1.554 \text{ Mg m}^{-3}$
Friclinic, $P\overline{1}$	Mo $K\alpha$ radiation
a = 7.3823 (2) Å	Cell parameters from 3305
p = 7.8262 (2) Å	reflections
z = 13.9272 (3) Å	$\theta = 1.5 - 27.0^{\circ}$
$\alpha = 79.062 \ (1)^{\circ}$	$\mu = 0.12 \text{ mm}^{-1}$
$\beta = 76.801 \ (1)^{\circ}$	T = 100.0 (1) K
$\nu = 78.810 \ (1)^{\circ}$	Block, pale yellow
$V = 759.54 (3) \text{ Å}^3$	$0.53 \times 0.23 \times 0.12 \ \mathrm{mm}$

Data collection

Bruker SMART APEX2 CCD areadetector diffractometer3295 independent reflections ω scans2850 reflections with $I > 2\sigma(I)$ ω scans $R_{int} = 0.027$ Absorption correction: multi-scan $\theta_{max} = 27.0^{\circ}$ (SADABS; Bruker, 2005) $h = -9 \rightarrow 9$ $T_{min} = 0.938, T_{max} = 0.985$ $k = -9 \rightarrow 6$ 10878 measured reflections $l = -17 \rightarrow 17$

Refinement

 Refinement on F^2 $w = 1/[\sigma^2(F_o^2) + (0.0581P)^2]$
 $R[F^2 > 2\sigma(F^2)] = 0.036$ + 0.2328P]

 $wR(F^2) = 0.108$ where $P = (F_o^2 + 2F_c^2)/3$

 S = 1.09 $(\Delta/\sigma)_{max} = 0.001$

 3295 reflections
 $\Delta\rho_{max} = 0.37$ e Å⁻³

 287 parameters
 $\Delta\rho_{min} = -0.26$ e Å⁻³

 All H-atom parameters refined
 $\Delta \rho_{min} = -0.26$ e Å⁻³

Table 1	
Selected geometric parameters (Å	, °).

O5-C7	1.2675 (17)	O6-C7	1.2354 (17)
C16-N3-C8	123.35 (12)	O6-C7-O5	128.64 (13)
O2-N1-C1-C2 O1-N1-C1-C6 O2-N1-C1-C6 O4-N2-C3-C2 O3-N2-C3-C2	-13.52 (18) -13.13 (18) 166.18 (12) 61.47 (16) -115.40 (13)	04-N2-C3-C4 N2-C3-C4-C7 C5-C4-C7-O6 C3-C4-C7-O6 C5-C4-C7-O5	-120.05 (14) 9.15 (19) 35.48 (18) -150.15 (13) -141 71 (13)

Table 2

Hydrogen-bond geometry (Å, °).

$D-\mathrm{H}\cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdot \cdot \cdot A$	
N3_H1N305	0.90(2)	1 74 (2)	2 6363 (15)	171 (2)	
$C6-H6\cdots O4^{i}$	0.972 (16)	2.438 (16)	3.1623 (18)	131 (1)	
$C9-H9\cdots O6$ $C12-H12\cdots O1^{ii}$	0.982(17) 0.953(17)	2.570 (17)	3.4986 (17) 3.2418 (18)	158(1) 144(1)	
$C12 - H12 \cdots O3^{iii}$	0.953(17) 0.953(17)	2.552 (17)	3.2086 (17)	144(1) 126(1)	
$C17 - H17C \cdots O5$	0.95 (2)	2.55 (2)	3.2987 (19)	137 (2)	

Symmetry codes: (i) x, y - 1, z; (ii) x + 1, y + 1, z - 1; (iii) -x + 1, -y + 2, -z.

All H atoms were located in a difference Fourier map and refined isotropically.

Data collection: *APEX2* (Bruker, 2005); cell refinement: *APEX2*; data reduction: *SAINT* (Bruker, 2005); program(s) used to solve structure: *SHELXTL* (Sheldrick, 1998); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*, *PARST* (Nardelli, 1995) and *PLATON* (Spek, 2003).

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