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Structure of complex of *N*-methylpiperidine betaine with *p*-hydroxybenzoic acid studied by X-ray, FT-IR and DFT methods

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Abstract

Crystal structure of the complex of *N*-methylpiperidine betaine (*N*-carboxymethyl-*N*-methylpiperidinium inner salt, MPB) with p-hydroxybenzoic acid (HBA) has been determined by X-ray diffraction. The crystals are triclinic, space group $P\bar{I}$, with a=6.1156(5), b=10.6869(10), c=12.0320(10) Å, $\alpha=109.55(1)^\circ$, $\beta=95.25(1)^\circ$, $\gamma=99.22(1)^\circ$, Z=2, R=0.034. Two molecules of p-hydroxybenzoic acid and two molecules of *N*-methylpiperidine betaine are linked together forming a centrosymmetric dimer, (MPB·HBA)₂, by four O–H···O hydrogen bonds of lengths 2.622(1) and 2.617(1) Å, between the carboxylic and hydroxy groups of HBA and both oxygen atoms of the carboxylate group of MPB, respectively. The piperidine ring has a chair conformation with the CH₂COO⁻ substituent in the axial position and the CH₃ group in the equatorial one. Two parallel aromatic rings in (MPB·HBA)₂ are distanced by 3.457 Å. In the crystals the complexes form "islands" related to the neighboring complexes by the inversion centers, weak C–H···O bonds and van der Waals forces. A broad band in the 3100–2400 cm⁻¹ region and two bands attributed to the ν C=O (1689 cm⁻¹) and ν _{as}COO (1607 cm⁻¹) vibrations in the FT-IR spectrum confirm the structure of the title complex. The two structures of MPB·HBA, denoted as **A** and **B**, have been optimized by the B3LYP/6-31G(d,p) method. In **A**, MPB forms a O–H···O hydrogen bond (2.562 Å) with the carboxylic group of HBA shorter than in the crystals, while in **B** it interacts with the phenolic group of HBA by a longer O–H···O hydrogen bond (2.661 Å) than in the crystals. Complex **A** is slightly more stable than **B** (0.15 kcal/mol).

Keywords: N-Methylpiperidine betaine; p-Hydroxybenzoic acid; Hydrogen bond; X-ray diffraction; FT-IR; DFT calculations

1. Introduction

Carboxybetaines are inner salts (zwitterionic compounds) consisting of an anionic carboxylate group and the oppositely charged quaternary ammonium moiety. The charged groups can be either coupled via π -electrons or are separated by a neutral skeleton and they are not conjugated [1–3].

In previous studies, we have determined the crystal structures of complexes of p-hydroxybenzoic acid with N-methylmorpholine betaine, N,N'-dimethylpiperazine mono- and di-betaines and nipecotic acid [4–7]. To get more details on the role of the hydroxyl group in the pro-

ton-donor molecules and the pattern of their hydrogen bonds, in this paper we present the molecular structure and spectroscopic properties of the complex of *N*-methylpiperidine betaine, MPB (p $K_a = 2.01$ [8]) with *p*-hydroxybenzoic acid, HBA, which has two different proton-donor groups, COOH (p $K_{a1} = 4.67$) and OH (p $K_{a2} = 9.37$) [9].

2. Experimental

N-Methylpiperidine betaine (N-carboxymethyl-N-methylpiperidinium inner salt, MPB) was obtained as described in Ref. [10]. Complex with p-hydroxybenzoic acid (HBA) was prepared by mixing one equivalent of MPB with one equivalent of HBA in methanol. The crude product was recrystallized from methanol, m.p. 178 °C. Analysis for $C_{15}H_{18}NO_5$: calcd: %C, 61.64; %H, 6.21; %N, 4.79; found:

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%C, 61.41; %H, 6.30; %N, 4.77. ¹H NMR (DMSO- d_6 , ppm): 1.51 (m, 2H, H-4), 1.79 (m, 4H, H-3, H-5), 3.20 (s, 3H, CH₃), 3.40 (m, 2H, H-2 ax, H-6 ax), 3.63 (m, 2H, H-2 eq, H-6 eq), 3.72 (s, 2H, N⁺CH₂), 6.86 (d, 2H, H-13, H-15), 8.26 (d, 2H, H-12, H-16). ¹³C NMR (DMSO- d_6 , ppm): 19.49 (C-3, C-5), 21.07 (C-4), 47.28 (CH₃), 59.81 (C-2, C-6), 63.28 (N⁺CH₂), 115.12 (C-13, C-15), 121.26 (C-11), 131.36 (C-12, C-16), 161.77 (C-14), 164.71 (COO), 167.17 (C=O). The deuterated analogue was prepared by threefold crystallization form CH₃OD.

Crystals of (MPB·HBA)₂ for diffraction measurements were grown from methanol. Diffraction measurements were carried out at ambient temperature on a KUMA-KM4 diffractometer equipped with CCD detector, using CuKα radiation [11]. Data reduction was carried out with the KM4 CCD software [12]. The structure was solved with direct methods SHELXS-97 program [13]. All atom positions were found from E-map, non-hydrogen atoms were refined with anisotropic displacement parameters and the parameters of hydrogen atoms were refined by isotropic methods, using the SHELXL-97 program [14]. The final *R* factor equals 0.0344 for 3777 reflections and 275 refined parameters. The crystal data and details of data processing are given in Table 1, and the final fractional atomic coordinates are in Table 2. The supplementary information in the

Table 1 Crystal data and structure refinement for the complex of *N*-methylpiperidine betaine with *p*-hydroxybenzoic acid, (MPB·HBA)₂

1 7 7	72		
Empirical formula	C ₁₅ H ₂₁ NO ₅		
Formula weight	295.33		
Temperature	293(2) K		
Wavelength	0.71073 Å		
Crystal system	Triclinic		
Space group	P $\bar{1}$		
Unit cell dimensions	a = 6.1156(5) Å		
	b = 10.6869(10) Å		
	c = 12.0320(10) Å		
	$\alpha = 109.545(8)^{\circ}$		
	$\beta = 95.249(7)^{\circ}$		
	$\gamma = 99.223(7)^{\circ}$		
Volume	$722.53(11) \text{ Å}^3$		
Z	2		
Calculated density	1.357 g/cm^3		
Absorption coefficient	$0.102~{\rm mm}^{-1}$		
F(000)	316		
Crystal size	$0.2 \times 0.2 \times 0.55 \text{ mm}$		
θ range for data collection	2.07–29.76°		
Limiting indices	$-8 \leqslant h \leqslant 8$		
	$-14 \leqslant k \leqslant 12$		
	$-14 \leqslant l \leqslant 16$		
Reflections collected/unique	$10274/3777 [R_{\text{int}} = 0.0126]$		
Completeness to $\theta = 25.00$	99.3%		
Refinement method	Full-matrix least-squares on F^2		
Data/restraints/parameters	3777/0/275		
Goodness-of-fit on F^2	1.091		
Final R indices $[I > 2\sigma(I)]$	R1 = 0.0344, wR2 = 0.1035		
R indices (all data)	R1 = 0.0414, wR2 = 0.1102		
Extinction coefficient	0.075(8)		
Largest diff. peak and hole	$0.315 \text{ and } -0.179 \text{ e Å}^{-3}$		

Table 2 Atomic coordinates and equivalent isotropic displacement parameters $(\mathring{A}^2 \times 10^3)$ for the complex of *N*-methylpiperidine betaine with *p*-hydroxybenzoic acid

Atom	X	у	Z	U(eq)
N(1)	0.9324(1)	0.2604(1)	0.5942(1)	28(1)
C(2)	1.1092(2)	0.2264(1)	0.6702(1)	32(1)
C(3)	1.2401(2)	0.3511(1)	0.7713(1)	38(1)
C(4)	1.0859(2)	0.4212(1)	0.8523(1)	46(1)
C(5)	0.9032(2)	0.4554(1)	0.7798(1)	40(1)
C(6)	0.7790(2)	0.3339(1)	0.6741(1)	36(1)
C(7)	0.7885(2)	0.1319(1)	0.5056(1)	45(1)
C(8)	1.0334(2)	0.3480(1)	0.5288(1)	29(1)
C(9)	1.1934(2)	0.2923(1)	0.4426(1)	32(1)
O(1)	1.2376(2)	0.1782(1)	0.4278(1)	46(1)
O(2)	1.2649(1)	0.3715(1)	0.3920(1)	45(1)
C(10)	0.6113(2)	0.0567(1)	0.8157(1)	33(1)
O(3)	0.7746(1)	-0.0112(1)	0.7905(1)	46(1)
O(4)	0.4717(2)	0.0605(1)	0.7395(1)	51(1)
C(11)	0.6203(2)	0.1291(1)	0.9452(1)	29(1)
C(12)	0.8024(2)	0.1397(1)	1.0284(1)	33(1)
C(13)	0.8046(2)	0.2048(1)	1.1490(1)	35(1)
C(14)	0.6230(2)	0.2610(1)	1.1885(1)	31(1)
C(15)	0.4389(2)	0.2497(1)	1.1062(1)	33(1)
C(16)	0.4382(2)	0.1841(1)	0.9857(1)	32(1)
O(5)	0.6355(1)	0.3265(1)	1.3075(1)	44(1)
H(2A)	1.0270(20)	0.1612(13)	0.6995(11)	40(3)
H(2B)	1.2020(20)	0.1809(12)	0.6163(11)	35(3)
H(3B)	1.3250(20)	0.4172(13)	0.7407(11)	38(3)
H(3A)	1.3490(20)	0.3221(14)	0.8158(12)	44(3)
H(4A)	1.0190(30)	0.3590(15)	0.8895(13)	55(4)
H(4B)	1.1750(30)	0.5085(17)	0.9188(15)	63(4)
H(5A)	0.7880(20)	0.4894(14)	0.8280(13)	51(4)
H(5B)	0.9680(20)	0.5295(14)	0.7522(12)	44(3)
H(6A)	0.6990(20)	0.2669(12)	0.7013(11)	37(3)
H(6B)	0.6760(20)	0.3586(13)	0.6193(11)	43(3)
H(7A)	0.7220(30)	0.0765(16)	0.5509(14)	61(4)
H(7B)	0.8860(30)	0.0848(15)	0.4520(14)	55(4)
H(7C)	0.6730(30)	0.1554(15)	0.4645(14)	55(4)
H(8A)	0.9120(20)	0.3630(13)	0.4835(11)	38(3)
H(8B)	1.1140(20)	0.4344(12)	0.5893(11)	35(3)
H(12)	0.9260(20)	0.1005(14)	1.0016(13)	48(4)
H(13)	0.9290(20)	0.2101(14)	1.2069(12)	45(3)
H(15)	0.3120(20)	0.2872(14)	1.1362(12)	45(3)
H(16)	0.3070(20)	0.1720(14)	0.9275(13)	48(3)
H(3)	0.7560(30)	-0.0699(18)	0.7107(16)	71(5)
H(5)	0.5010(30)	0.3374(18)	1.3259(16)	69(5)

U(eq) is defined as one third of the trace of the orthogonalized Uij tensor.

CIF form is available from Cambridge Crystallographic Data Base Center, No CCDC 642346.

FT-IR spectra were measured on a Bruker IFS 66v/S instrument, evacuated to avoid water and CO₂ absorptions. Solid state spectra were recorded in Nujol and Fluorolube suspension using KBr plates. Each spectrum consisted of 64 scans. The Raman spectrum was measured on a Bruker IFS 66 instrument.

The NMR spectra were recorded on a Varian Gemini 300 VT spectrometer operating at 300.07 and 75.46 MHz for 1 H and 13 C, respectively. The spectra were measured in DMSO- d_6 relative to TMS as internal reference. The 2D (COSY, HETCOR) spectra were obtained with standard Varian software.

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