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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.002 Å R factor = 0.053 wR factor = 0.157 Data-to-parameter ratio = 16.9

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

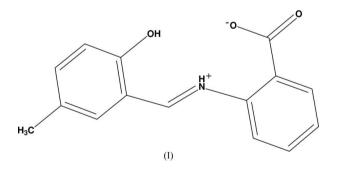
2-(2-Hydroxy-5-methylbenzylideneammonio)benzoate

The N atom in the title compound, $C_{15}H_{13}NO_3$, is protonated by the -COOH group. The molecular conformation is stabilized by strong N-H···O hydrogen bonds. The crystal packing is stabilized by O-H···O and C-H···O intermolecular interactions.

Comment

The recognition (Schmidtchen & Berger, 1997) and sensing (Czanik, 1994) of anionic analytes have been considered as a key research theme within the generalized area of supramolecular chemistry. A significant amount of work has been devoted to obtain specific chemosensors that are able to change one or several macroscopic properties, upon complexation with the target guests. In response to the molecular recognition event, changes in colour or optical properties [fluorescence (Bargossi *et al.*, 2000) and absorbance (Löhr & Vögtle, 1985)] are the output signals used in the development of optical chemosensors.

The bond lengths and bond angles of the title compound, (I), are comparable to the literature values (Allen *et al.*, 1987). The N atom is protonated by the –COOH group. The dihedral angle between the two benzene rings in the structure is 5.6 (1)°.



The molecular structure is stabilized by strong N-H···O interactions in which N1 acts as a bifurcated donor to O1 and O2, generating two S(6) motifs. The crystal packing is stabilized by O-H···O and C-H···O intermolecular interactions. Atom O1 acts as donor to atom O2 at $(-x, y, \frac{3}{2} - z)$, generating a dimer of the type $R_2^2(20)$. The packing of molecules in the unit cell viewed down the *a* axis resembles the shape of a wave.

Experimental

© 2007 International Union of Crystallography All rights reserved 2-(2-Hydroxy-5-methylbenzylideneammonio)benzoate was synthesized by Schiff base condensation between 2-aminobenzoic acid and

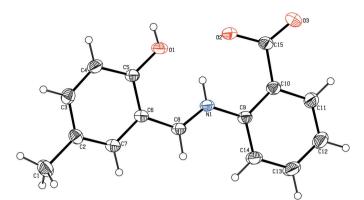


Figure 1

The molecular structure of the title compound, showing 30% probability displacement ellipsoids.

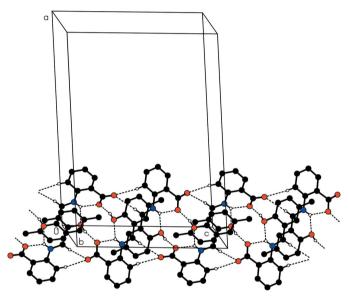


Figure 2

A view of the N-H···O intramolecular and O-H···O and C-H···O intermolecular interactions. For clarity, H atoms that are not involved in hydrogen-bonding have been omitted. Hydrogen bonds are shown as dashed lines.

2-hydroxy-5-methylbenzaldehyde. 2-Aminobenzoic acid (1 g, 7 mmol) was dissolved in methanol (25 ml), and a solution of 2hydroxy-5-methylbenzaldehyde (1 g, 7 mmol) in methanol was added with stirring. The resulting mixture was heated at reflux for 1 h and cooled to room temperature. The solid product was collected by filtration and washed with cold methanol. The microcrystalline compound was recrystallized from hot methanol; crystals suitable for X-ray diffraction studies were obtained on slow evaporation. Yield 92%, m.p. 476 K.

Crystal data

-	
C ₁₅ H ₁₃ NO ₃	Z = 8
$M_r = 255.26$	$D_x = 1.367 \text{ Mg m}^{-3}$
Monoclinic, $C2/c$	Mo $K\alpha$ radiation
a = 22.5716 (15) Å	$\mu = 0.10 \text{ mm}^{-1}$
b = 7.1742 (4) Å	T = 293 (2) K
c = 15.4336 (9) Å	Block, colourless
$\beta = 97.093 \ (2)^{\circ}$	$0.24 \times 0.22 \times 0.21 \text{ mm}$
V = 2480.1 (3) Å ³	

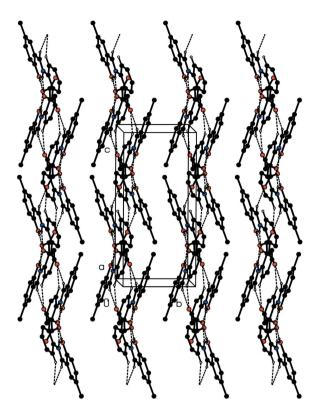


Figure 3

The molecular packing of (I), viewed down the a axis. For clarity, H atoms that are not involved in hydrogen bonding have been omitted. Hydrogen bonds shown as dashed lines.

2941 independent reflections

 $R_{\rm int} = 0.022$ $\theta_{\rm max} = 28.0^{\circ}$

2471 reflections with $I > 2\sigma(I)$

Data collection

Bruker SMART CCD diffractometer (i) scans Absorption correction: none 13716 measured reflections

Refinement

$w = 1/[\sigma^2(F_o^2) + (0.0927P)^2]$
+ 0.9763P]
where $P = (F_0^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{\rm max} = 0.001$
$\Delta \rho_{\rm max} = 0.34 \text{ e } \text{\AA}^{-3}$
$\Delta \rho_{\rm min} = -0.17 \text{ e } \text{\AA}^{-3}$

Table 1 Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
N1-H1···O1	0.86	2.01	2.663 (2)	132
$N1 - H1 \cdots O2$	0.86	1.91	2.592 (2)	135
$O1 - H1D \cdots O2^{i}$	0.82	1.66	2.464 (2)	166
C8-H8···O3 ⁱⁱ	0.93	2.56	3.462 (2)	165
C11-H11O3	0.93	2.47	2.784 (2)	100
$C14-H14\cdots O3^{ii}$	0.93	2.41	3.282 (2)	156

Symmetry codes: (i) $-x, y, -z + \frac{3}{2}$; (ii) $x, -y, z + \frac{1}{2}$.

All H atoms were refined using a riding model with C-H = 0.93 Å, $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}({\rm C})$ for aromatic, ${\rm C-H} = 0.96$ Å, $U_{\rm iso}({\rm H}) = 1.5U_{\rm eq}$ (C) for CH₃, N–H = 0.86 Å, $U_{iso}(H) = 1.2U_{eq}$ (N) for the NH group and O–H = 0.83 Å, $U_{iso}(H) = 1.5U_{eq}$ (O) for the OH group. In addition, the torsion angles for the rotation of the OH and methyl groups were refined.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT* (Bruker, 2001); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXL97* and *PARST* (Nardelli, 1995).

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