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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.002 Å R factor = 0.038 wR factor = 0.116 Data-to-parameter ratio = 12.0

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

2-Hydroxy-3-methoxybenzoic acid

The title compound, $C_8H_8O_4$, has an intramolecular O-H···O hydrogen bond. Intermolecular O-H···O hydrogen bonding gives rise to a dimeric structure, which is further extended into infinite stacks parallel to the *a* axis via π - π interactions between the aromatic rings of neigboring molecules.

Comment

Hydrogen-bonding interactions between ligands are specific and directional. When present in metal complexes they are usually not dependent on the properties of the metal ions, but they play a critical role in the overall structures and functions of the complexes and the way in which they pack in the solid state. In this sense, 2-hydroxy-3-methoxybenzoic acid is an excellent candidate for the construction of supramolecular complexes, since it not only has multiple coordination modes but can also form regular hydrogen bonds, functioning as both a hydrogen-bond donor and acceptor (Moncol *et al.*, 2006; Kozlevcar *et al.*, 2006). In this context we report here the crystal structure of the title compound, (I).



The molecular structure of (I) is depicted in Fig. 1. The C– O and C–C distances show no remarkable features, with C– O distances in the range 1.237 (2)–1.426 (2) Å. The title molecule acts as both a hydrogen-bond donor and acceptor, forming dimers with neighboring molecules through O– H···O hydrogen bonding (Table 1), while π - π interactions between the dimers lead to the formation of infinite stacks of the molecules along the *a* axis (Fig. 2). The centroid-tocentroid and interplanar distances between adjacent aromatic rings (symmetry code: x - 1, y, z) are 3.867 (4) and 3.528 (3) Å, respectively.

Experimental

© 2007 International Union of Crystallography All rights reserved 2-Hydroxy-3-methoxybenzoic acid was dissolved in hot methanol with stirring. Yellow single crystals suitable for X-ray diffraction were

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Figure 1



obtained at room temperature by slow evaporation of the solvent over a period of several days.

V = 762.81 (6) Å³

Mo $K\alpha$ radiation

 $0.20 \times 0.18 \times 0.15~\text{mm}$

1346 independent reflections

1051 reflections with $I > 2\sigma(I)$

 $\mu = 0.12 \text{ mm}^{-1}$

T = 293 (2) K

 $R_{\rm int} = 0.024$

Z = 4

Crystal data

Data collection

Bruker APEXII area-detector diffractometer Absorption correction: none 4322 measured reflections

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.038$	112 parameters
$wR(F^2) = 0.116$	H-atom parameters constrained
S = 1.05	$\Delta \rho_{\rm max} = 0.12 \text{ e} \text{ \AA}^{-3}$
1346 reflections	$\Delta \rho_{\rm min} = -0.19 \ {\rm e} \ {\rm \AA}^{-3}$

Table 1

Hydrogen-bond geometry (Å, $^{\circ}$).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D{\cdots}A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$O2-H2\cdots O1^i$	0.82	1.85	2.6703 (15)	176
$O3-H3\cdots O1$	0.82	1.90	2.6169 (14)	145

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

All H atoms were placed in calculated positions (C–H = 0.93 or 0.96 Å; O–H = 0.82 Å) and were refined using a riding model, with



Figure 2

The molecular packing of (I), showing the intra- and intermolecular hydrogen bonding and $\pi - \pi$ interactions. The view is along the *a* axis and hydrogen-bonding interactions are shown as dashed lines.

 $U_{iso}(H) = 1.2U_{eq}(C)$ for H atoms of the aromatic ring and $1.5U_{eq}(C,O)$ for methyl and hydroxyl H atoms.

Data collection: *APEXII* (Bruker, 2004); cell refinement: *SAINT* (Bruker, 2004); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 2004); software used to prepare material for publication: *SHELXTL*.

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