Received 25 November 2005 Accepted 9 December 2005

Online 14 December 2005

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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#### **Key indicators**

Single-crystal X-ray study T = 296 KMean  $\sigma$ (C–C) = 0.003 Å R factor = 0.046 wR factor = 0.109 Data-to-parameter ratio = 13.4

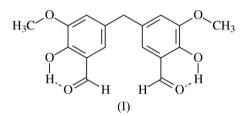
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

# Bis(3-formyl-4-hydroxy-5-methoxyphenyl)methane

In the title compound,  $C_{17}H_{16}O_3$ , the asymmetric unit contains one half-molecule; a twofold rotation axis bisects the molecule. The structure is stabilized by  $O-H\cdots N$  intramolecular hydrogen bonds and  $C-H\cdots \pi$  and  $\pi-\pi$  intermolecular interactions.

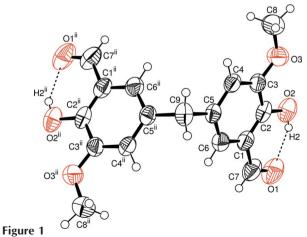
## Comment

Hydroxy-substituted benzaldehyde reagents used for condensation with primary amines, hydrazines, hydroxylamine and other primary amine derivatives afford imine derivatives which can function as ligands towards a number of metal cations (Loudon, 2002; Khandar & Nejati, 2000; Khandar & Rezvani, 1999).



The asymmetric unit of the title compound, (I), contains one half-molecule; a twofold rotation axis passes through C9 (Fig. 1). The bond lengths and angles (Table 1) are in normal ranges (Allen *et al.*, 1987).

In (I), molecules have strong intramolecular  $O-H\cdots O$ hydrogen bonds (Table 2) and they are linked through C8–  $H8a\cdots Cg1$  (Cg1 is the centroid of the C1–C6 ring) and Cg1 $\cdots$ Cg1 intermolecular interactions (Fig. 2). For the C8–



The molecular structure of (I), with the atomic numbering scheme. Displacement ellipsoids are drawn at the 50% probalitity level. The intramolecular hydrogen bonds are shown as dashed lines. [Symmetry code: (i) 1 - x, y,  $\frac{1}{2} - z$ .]

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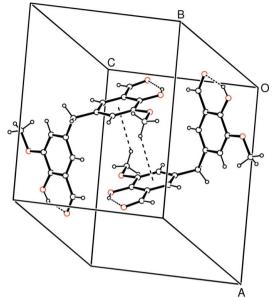


Figure 2

A packing diagram of (I) with the C–H··· $\pi$  intermolecular interactions shown as dashed lines.

H8*a*···*Cg*1 contact, the distance between atom H8*a* and the aromatic ring centroid is 3.01 (3) Å (symmetry code: 1 - x, 1 - y, -z). There is also  $\pi$ - $\pi$  stacking between adjacent molecules at (x, y, z) and  $(-x, y, \frac{1}{2} - z)$ , with distances of 3.608 (16) Å between the rings centroids and perpendicular distances of 2.481 (16) Å between the rings.

## **Experimental**

A mixture of *o*-vanillin (0.1 mol) and formaldehyde (0.1 mol) was stirred at 393 K for 2 h. The solution was added to boiling ethyl alcohol and stirred at 393 K for 20 min and cooled to room temperature. The precipitate was filtered off and recrystallized from ethyl alcohol by slow evaporation (yield 1.58 g, 10%, m.p. 424–426 K).

 $D_x = 1.411 \text{ Mg m}^{-3}$ Mo  $K\alpha$  radiation Cell parameters from 3016 reflections  $\theta = 2.9-27.5^{\circ}$ 

 $\mu = 0.11~\mathrm{mm}^{-1}$ 

T = 296 (2) K

 $0.40\,\times\,0.24\,\times\,0.07$  mm

Plate, yellow

### Crystal data

| $C_{17}H_{16}O_{6}$   |
|---|
| $M_r = 316.30$  |
| Monoclinic, $C2/c$  |
| a = 14.960 (3)  Å   |
| $b = 8.2889 (11) \text{\AA}$                                    |
| c = 13.249 (3) Å  |
| $\beta = 115.015 (15)^{\circ}$<br>V = 1488.8 (5) Å <sup>3</sup> |
| V = 1488.8 (5) Å <sup>3</sup>                                   |
| Z = 4   |
|   |
| Data collection   |

#### Data collection

| Stoe IPDS-II diffractometer            | 828 reflections with $I > 2\sigma(I)$ |
|--|---------------------------------------|
| $\omega$ scans                         | $R_{\rm int} = 0.065$                 |
| Absorption correction: integration     | $\theta_{\rm max} = 26.0^{\circ}$     |
| (X-RED32; Stoe & Cie, 2002)            | $h = -18 \rightarrow 18$              |
| $T_{\min} = 0.963, \ T_{\max} = 0.993$ | $k = -10 \rightarrow 10$              |
| 6042 measured reflections              | $l = -16 \rightarrow 16$              |
| 1470 independent reflections           |                                       |

Refinement

| Refinement on $F^2$<br>$R[F^2 > 2\sigma(F^2)] = 0.046$<br>$wR(F^2) = 0.109$<br>S = 1.00<br>1470 reflections<br>110 parameters | H atoms treated by a mixture of<br>independent and constrained<br>refinement<br>$w = 1/[\sigma^2(F_o^2) + (0.049P)^2]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{max} < 0.001$ $\Delta\rho_{max} = 0.12$ e Å <sup>-3</sup> $\Delta\rho_{min} = -0.11$ e Å <sup>-3</sup> |
|---|--|
| Table 1   |  |

| Selected | geometric | parameters | (Å | °) |  |
|----------|-----------|------------|----|----|--|
|          |           |            |    |    |  |

| C1-C2                    | 1.388 (3)   | C3-C4                    | 1.381 (3)   |
|--------------------------|-------------|--------------------------|-------------|
| C1-C6                    | 1.399 (3)   | C4-C5                    | 1.400 (3)   |
| C2-O2                    | 1.360 (2)   | C5-C6                    | 1.368 (3)   |
| C2-C3                    | 1.395 (3)   | C7-O1                    | 1.219 (3)   |
| C2-C1-C7                 | 120.9 (2)   | O3-C3-C4                 | 124.8 (2)   |
| O2-C2-C3                 | 118.0 (2)   | C5-C9-C5 <sup>i</sup>    | 113.6 (3)   |
| C6-C1-C7-O1              | -176.1 (2)  | C4-C5-C9-C5 <sup>i</sup> | 108.23 (19) |
| C6-C5-C9-C5 <sup>i</sup> | -72.25 (18) | C4-C3-O3-C8              | -1.6 (3)    |
|                          |             |                          |             |

Symmetry code: (i)  $-x + 1, y, -z + \frac{1}{2}$ .

. .

| Table 2                        |  |
|--------------------------------|--|
| Hydrogen-bond geometry (Å, °). |  |

| $D - H \cdot \cdot \cdot A$ | D-H  | $H \cdot \cdot \cdot A$ | $D \cdots A$ | $D - \mathbf{H} \cdot \cdot \cdot A$ |
|-----------------------------|------|-------------------------|--------------|--------------------------------------|
| O2−H2···O1                  | 0.82 | 1.94                    | 2.648 (3)    | 145                                  |

Atom H9, attached to C9, was located in a difference map and refined isotropically [C-H = 0.98 (2) Å]. The remaining H atoms were positioned geometrically [0.82 (OH), 0.93 (CH) and 0.96 Å (CH<sub>3</sub>)] and constrained to ride on their parent atoms with  $U_{iso}(H)$  values of 1.5 (1.2 for methine) times  $U_{eq}(C,O)$ .

Data collection: X-AREA (Stoe & Cie, 2002); cell refinement: X-AREA; data reduction: X-RED32; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP3 for Windows (Farrugia, 1997); software used to prepare material for publication: WinGX (Farrugia, 1999).

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