Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 298 KMean $\sigma(\text{C}-\text{C}) = 0.003 \text{ Å}$ R factor = 0.067 wR factor = 0.226 Data-to-parameter ratio = 17.7

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

tert-Butyl 3-(3,4-dihydroxyphenyl)prop-2-enoate

The molecule of the title compound, $C_{13}H_{16}O_4$, adopts an *E* configuration, the carboxyl group and benzene ring being located on opposite sides of the C=C bond. Intermolecular $O-H\cdots O$ hydrogen bonding helps to stabilize the crystal structure.

Received 21 August 2006 Accepted 22 August 2006

Comment

Caffeic acid and its derivatives are widely distributed in the plant kingdom (Chen *et al.*, 1999). These compounds are known to have anti-atherosclerotic, antibacterial, anti-inflammatory, antiproliferative, immunostimulatory, anti-oxidative, antiviral and neuroprotective properties (Son & Lewis, 2002). As part of our ongoing research on caffeic acid derivatives (Xia & Hu, 2005), we present here the structure of the title compound, (I).



The molecular structure of (I) is illustrated in Fig. 1. The molecule displays an *E* configuration, with the benzene ring and carboxyl group located on opposite sides of the C7—C8 double bond. Bond distances agree with those found in similar caffeate structures (Xia *et al.*, 2004, 2006). The propenoate group is tilted with respect to the benzene ring with a dihedral angle of 22.71 (10)°.

Intermolecular hydrogen bonding between hydroxy groups and between carboxyl and hydroxy groups (Table 1) helps to stabilize the crystal structure.

Experimental

Meldrum's acid (3.6 g, 25 mmol) and *tert*-butanol (1.85 g, 25 mmol) were mixed in toluene (50 ml) and the mixture was refluxed for 5 h. After cooling to room temperature, 3,4-dihydroxybenzaldehyde (1.4 g, 10 mmol), pyridine (2.5 ml) and piperidine (0.25 ml) were added. The resulting mixture was stirred for 12 h at room temperature. The solvents were then distilled out under vacuum. The residue was dissolved in diethyl ether (30 ml), washed in turn with a saturated solution of sodium bicarbonate (20 ml), dilute hydrochloric acid (20 ml) and distilled water (20 ml). The ether phase was dried by anhydrous MgSO₄ overnight. After removal of the drier, the solvent was distilled out, giving a yellowish sheet-like solid (1.45 g, 61.4%). Recrystallization from a mixture of acetone, diethyl ether and water (8:1:1) gave single crystals of (I).

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Crystal data

 $C_{13}H_{16}O_4$ $M_r = 236.26$ Monoclinic, P_{21}^2/c a = 9.334(3) Å b = 11.000 (3) Å c = 12.701 (4) Å β = 98.005 (4)° V = 1291.4 (7) Å³

Data collection

Bruker APEX CCD area-detector diffractometer φ and ω scans Absorption correction: none 6285 measured reflections

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_0^2) + (0.1179P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.067$	+ 0.1691P]
$wR(F^2) = 0.226$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.11	$(\Delta/\sigma)_{\rm max} < 0.001$
2826 reflections	$\Delta \rho_{\rm max} = 0.26 \ {\rm e} \ {\rm \AA}^{-3}$
160 parameters	$\Delta \rho_{\rm min} = -0.42 \text{ e} \text{ Å}^{-3}$
H-atom parameters constrained	Extinction correction: SHELXL97
*	Extinction coefficient: 0.031 (6)

Z = 4

 $D_x = 1.215 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation

Prism, light brown

 $0.25 \times 0.20 \times 0.20$ mm

2826 independent reflections

1857 reflections with $I > 2\sigma(I)$

 $\mu = 0.09 \text{ mm}^{-1}$

T = 298 (2) K

 $R_{\rm int} = 0.078$

 $\theta_{\rm max} = 27.1^{\circ}$

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	$D-\mathrm{H}$	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdot \cdot \cdot A$
$O1-H1\cdots O3^i$	0.82	1.85	2.669 (2)	172
$O2-H2\cdots O1^{ii}$	0.82	2.12	2.826 (2)	144
Summatry and (i)	v v 1 m 1,	(ii) x y z	1	

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

H atoms bonded to O atoms were located in a difference Fourier map and refined with the constraint O-H = 0.82 Å $[U_{iso}(H) = 1.5U_{eq}(O)]$. Methyl H atoms were placed in calculated positions, with C-H = 0.96 Å and $U_{iso}(H) = 1.5U_{eq}(C)$. Other H atoms were placed



Figure 1

The molecular structure of (I), shown with 30% probability displacement ellipsoids (arbitrary spheres for H atoms).

in calculated positions, with C–H = 0.93 Å, and refined in riding mode $[U_{iso}(H) = 1.2U_{eq}(C)]$.

Data collection: *SMART* (Bruker, 2003); cell refinement: *SAINT* (Bruker, 2003); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Bruker, 2003); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

We are very grateful to the National Natural and Scientific Foundation (grant No. 20272053). We also acknowledge financial support from the Science and Technology Bureau of Zhejiang Province (grant No. 2005 C23022).

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