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## Structure Reports

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## Key indicators

Single-crystal X-ray study
$T=298 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.003 \AA$
$R$ factor $=0.067$
$w R$ 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.

[^0]
# tert-Butyl 3-(3,4-dihydroxyphenyl)prop-2-enoate 

The molecule of the title compound, $\mathrm{C}_{13} \mathrm{H}_{16} \mathrm{O}_{4}$, adopts an $E$ configuration, the carboxyl group and benzene ring being located on opposite sides of the $\mathrm{C}=\mathrm{C}$ bond. Intermolecular $\mathrm{O}-\mathrm{H} \cdots \mathrm{O}$ hydrogen bonding helps to stabilize the crystal structure.

## 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, antiinflammatory, antiproliferative, immunostimulatory, antioxidative, 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).

(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 $\mathrm{C} 7=\mathrm{C} 8$ 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) ${ }^{\circ}$.

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

## Experimental

Meldrum's acid ( $3.6 \mathrm{~g}, 25 \mathrm{mmol}$ ) and tert-butanol $(1.85 \mathrm{~g}, 25 \mathrm{mmol})$ were mixed in toluene ( 50 ml ) and the mixture was refluxed for 5 h . After cooling to room temperature, 3,4-dihydroxybenzaldehyde $(1.4 \mathrm{~g}, 10 \mathrm{mmol})$, pyridine $(2.5 \mathrm{ml})$ and piperidine $(0.25 \mathrm{ml})$ 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 $(20 \mathrm{ml})$ and distilled water $(20 \mathrm{ml})$. The ether phase was dried by anhydrous $\mathrm{MgSO}_{4}$ overnight. After removal of the drier, the solvent was distilled out, giving a yellowish sheet-like solid ( $1.45 \mathrm{~g}, 61.4 \%$ ). Recrystallization from a mixture of acetone, diethyl ether and water (8:1:1) gave single crystals of (I).

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## Crystal data

$\mathrm{C}_{13} \mathrm{H}_{16} \mathrm{O}_{4}$
$M_{r}=236.26$
Monoclinic, $P 2_{1} / c$
$a=9.334$ (3) $\AA$
$b=11.000$ (3) $\AA$
$c=12.701$ (4) $\AA$
$\beta=98.005(4)^{\circ}$
$V=1291.4(7) \AA^{3}$
Data collection
Bruker APEX CCD area-detector
diffractometer
$\varphi$ and $\omega$ scans
Absorption correction: none
6285 measured reflections

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.067$
$w R\left(F^{2}\right)=0.226$
$S=1.11$
2826 reflections
160 parameters
H -atom parameters constrained

## $Z=4$

$D_{x}=1.215 \mathrm{Mg} \mathrm{m}^{-3}$
Mo $K \alpha$ radiation
$\mu=0.09 \mathrm{~mm}^{-1}$
$T=298$ (2) K
Prism, light brown
$0.25 \times 0.20 \times 0.20 \mathrm{~mm}$

2826 independent reflections 1857 reflections with $I>2 \sigma(I)$
$R_{\text {int }}=0.078$
$\theta_{\text {max }}=27.1^{\circ}$

$$
\begin{aligned}
& \begin{array}{l}
w=1 /\left[\sigma^{2}\left(F_{\mathrm{o}}^{2}\right)+(0.1179 P)^{2}\right. \\
\quad \\
\quad+0.1691 P] \\
\quad \text { where } P=\left(F_{\mathrm{o}}^{2}+2 F_{\mathrm{c}}^{2}\right) / 3 \\
(\Delta / \sigma)_{\max }<0.001 \\
\Delta \rho_{\max }=0.26 \mathrm{e} \AA^{-3} \\
\Delta \rho_{\min }=-0.42 \mathrm{e} \AA^{-3} \\
\text { Extinction correction: } S H E L X L 97 \\
\text { Extinction coefficient: } 0.031(6)
\end{array}
\end{aligned}
$$

Table 1
Hydrogen-bond geometry ( $\mathrm{A},{ }^{\circ}$ ).

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| $\mathrm{O}_{1}-\mathrm{H} 1 \cdots \mathrm{O}^{\text {i }}$ | 0.82 | 1.85 | $2.669(2)$ | 172 |
| $\mathrm{O}^{\mathrm{i}}-\mathrm{H} 2 \cdots \mathrm{O}^{1 i}$ | 0.82 | 2.12 | $2.826(2)$ | 144 |

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 $\mathrm{O}-\mathrm{H}=0.82 \AA\left[U_{\text {iso }}(\mathrm{H})=\right.$ $\left.1.5 U_{\text {eq }}(\mathrm{O})\right]$. Methyl H atoms were placed in calculated positions, with $\mathrm{C}-\mathrm{H}=0.96 \AA$ and $U_{\text {iso }}(\mathrm{H})=1.5 U_{\text {eq }}(\mathrm{C})$. 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 $\mathrm{C}-\mathrm{H}=0.93 \AA$, and refined in riding $\operatorname{mode}\left[U_{\text {iso }}(\mathrm{H})=1.2 U_{\text {eq }}(\mathrm{C})\right]$.

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.

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