Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

Chun-Nian Xia, Wei-Xiao Hu* and Wei Zhou

College of Pharmaceutical Science, Zhejiang University of Technology, Hangzhou 310032, People's Republic of China

Correspondence e-mail: huyang@mail.hz.zj.cn

Key indicators

Single-crystal X-ray study T = 298 K Mean σ (C–C) = 0.006 Å Disorder in main residue R factor = 0.095 wR factor = 0.243 Data-to-parameter ratio = 12.2

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

3-(4-Hydroxy-3-methoxy-2-nitrophenyl)acrylic acid

The title compound, $C_{10}H_9NO_6$, has an *E* configuration, with the carboxyl group and the benzene ring located on opposite sides of the C=C bond. Intermolecular $O-H \cdots O$ hydrogen bonding helps to stabilize the crystal structure.

Received 19 July 2006 Accepted 27 July 2006

Comment

Caffeic acid and its derivatives have antiatherosclerotic and neuroprotective properties (Son & Lewis, 2002). Also these compounds can inhibit the peroxynitrite-dependent tyrosine nitration (Pannala *et al.*, 1998). 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 has an *E* configuration, with the carboxyl group and the benzene ring located on opposite sides of the C7=C8 bond. The nitro group is twisted by 60.66 (18)° with respect to the benzene ring. Intermolecular $O-H\cdots O$ hydrogen bonding occurs in the crystal structure (Table 1).



Figure 1

The structure of (I), shown with 30% probability displacement ellipsoids (arbitrary spheres for H atoms). The dashed line indicates one of the disordered components.

© 2006 International Union of Crystallography All rights reserved

Experimental

Compound (I) was synthesized according to the method reported by Grenier *et al.* (2000). Single crystals were obtained by slow evaporation of a methanol/water solution (1:1).

Z = 4

 $D_{\rm r} = 1.485 {\rm Mg} {\rm m}^{-3}$

Mo Ka radiation

Prism, light yellow

 $0.15 \times 0.10 \times 0.05 \text{ mm}$

 $w = 1/[\sigma^2(F_0^2) + (0.0647P)^2]$

+ 3.1253*P*] where $P = (F_0^2 + 2F_c^2)/3$

 $\Delta \rho_{\rm max} = 0.34 \text{ e} \text{ Å}^{-3}$

 $\Delta \rho_{\rm min} = -0.34$ e Å⁻³

 $(\Delta/\sigma)_{\rm max} = 0.002$

 $\mu = 0.13 \text{ mm}^{-1}$

T = 298 (2) K

Crystal data

 $\begin{array}{l} C_{10}H_9NO_6\\ M_r = 239.18\\ Monoclinic, P2_1/c\\ a = 3.8710 \ (14) \ \text{\AA}\\ b = 8.049 \ (5) \ \text{\AA}\\ c = 34.342 \ (5) \ \text{\AA}\\ \beta = 90.03 \ (2)^\circ\\ V = 1070.0 \ (8) \ \text{\AA}^3 \end{array}$

Data collection

Bruker SMART1000 CCD
diffractometer1885 independent reflections
1477 reflections with $I > 2\sigma(I)$
 φ and ω scans φ and ω scans $R_{int} = 0.127$
 $\theta_{max} = 25.0^{\circ}$ 4259 measured reflections $\theta_{max} = 25.0^{\circ}$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.095$ $wR(F^2) = 0.243$ S = 1.101885 reflections 155 parameters H-atom parameters constrained

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$\begin{array}{c} O1 - H1 \cdots O2^{i} \\ O2 - H2 \cdots O1^{i} \\ O6 - H6A \cdots O3^{ii} \end{array}$	0.89	1.80	2.670 (6)	164
	0.92	1.76	2.670 (6)	171
	0.84	2.01	2.847 (5)	171

Symmetry codes: (i) -x + 2, -y + 2, -z; (ii) x, y - 1, z.

The carboxyl group is disordered over two sites; in a difference Fourier map two peaks were found at positions near to both O1 and O2. They were set as H1 and H2 with each having 0.5 occupancy. The hydroxy H was also located in a difference Fourier map. H atoms bonded to O atoms were refined as riding in their as-found relative positions, with $U_{iso}(H) = 1.5U_{eq}(O)$. Methyl H atoms were placed in calculated positions, with C-H = 0.96 Å, and the torsion angle was refined to fit the electron density, with $U_{iso}(H) = 1.5U_{eq}(C)$. Other H atoms were placed in calculated positions, with C-H = 0.93 Å, and refined in riding mode, with $U_{iso}(H) = 1.2U_{eq}(C)$.

Data collection: *SMART* (Bruker, 1997); cell refinement: *SAINT* (Bruker, 1997); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1999); software used to prepare material for publication: *SHELXL97*.

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

References

Bruker (1997). SMART and SAINT. Bruker AXS Inc., Madison, Wisconsin, USA.

- Farrugia, L. J. (1999). J. Appl. Cryst. 32, 837-838.
- Grenier, J. L., Cotelle, N., Gatteau, J. P. & Cottelle, P. (2000). J. Phys. Org. Chem. 13, 511–517.
- Pannala, A., Razaq, R., Halliwell, B., Singh, S. & Rice-Evans, C. A. (1998). Free Rad. Biol. Med. 24, 594–606.
- Sheldrick, G. M. (1997). SHELXL97 and SHELXS97. University of Göttingen, Germany.
- Son, S. & Lewis, B. A. (2002). J. Agric. Food Chem. 50, 468-472.
- Xia, C.-N. & Hu, W.-X. (2005). J. Chem. Res. 5, 332-334.