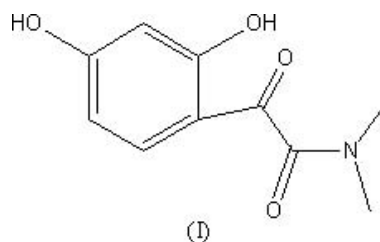


Wei Huang, Peiliang Zhao,
Zhongzhen Zhou, Qiong Chen,
Deyou Teng and Guangfu Yang*Key Laboratory of Pesticide and Chemical
Biology of the Ministry of Education, College
of Chemistry, Central China Normal University,
Wuhan 430079, ChinaCorrespondence e-mail:
gfyang@mail.ccnu.edu.cn**Key indicators**Single-crystal X-ray study
 $T = 283\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.002\text{ \AA}$
 R factor = 0.049
 wR factor = 0.148
Data-to-parameter ratio = 16.8For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.**2-(2,4-Dihydroxyphenyl)-*N,N*-dimethyl-
2-oxoacetamide**In the crystal structure of the title compound, $\text{C}_{10}\text{H}_{11}\text{NO}_4$, O—H \cdots O inter- and intramolecular hydrogen bonds are present, as well as intermolecular C—H \cdots O interactions.

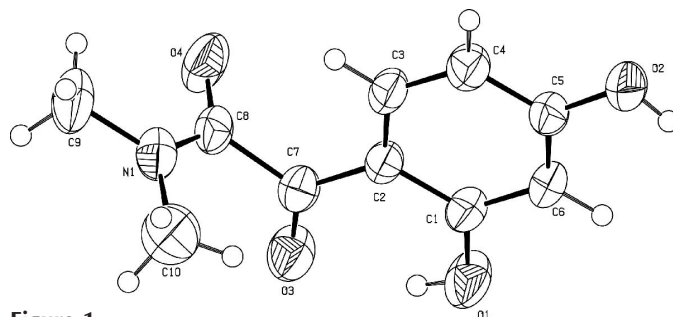
Received 23 March 2005

Accepted 21 April 2005

Online 27 April 2005

Comment α -Ketoamides have potential applications for the synthesis of various useful products, including α -amino acids, α -hydroxyacids and heterocyclic compounds (Zhou & Chen, 2001). Furthermore, α -ketoamide derivatives have been reported to exhibit a range of biological properties, including antimicrobial and antipsychotic activity (Yang *et al.*, 2002). For these reasons, we are interested in exploring and developing novel strategies for the synthesis of α -ketoamides.In this paper, we present the X-ray crystallographic analysis of the title compound, (I). The bond lengths and angles are unremarkable. The dihedral angle between the plane through C9/C10/N1/C8/O4/C7 and the plane of the benzene ring is $89.5(2)^\circ$ (Fig. 1). As shown in Fig. 2, the molecules are linked by both intra- and intermolecular hydrogen bonds. Geometric details of these are listed in Table 1. No π - π stacking interactions are observed in the crystal structure.**Experimental**

2,4-Dihydroxyacetophenone (10 mmol) and dimethyl oxalate (1.28 mmol) were dissolved in dimethylformamide (5 ml). This was

**Figure 1**

A view of the molecule of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level and H atoms are represented by circles of arbitrary size.

added to a solution of sodium (0.8 g) in methanol (20 ml) and the mixture was stirred at room temperature for 48 h. Acetic acid (36 ml) was then added and the mixture was heated to 373 K for 5 h. The resulting mixture was purified by column chromatography on silica gel using acetone/diethyl ether (9:1 v/v) as eluent to afford (I) (yield 40%, m.p. 464 K). ¹H NMR (CDCl₃, 400 MHz): 11.10 (s, 1H, 2'OH), 10.79 (s, 1H, 4'OH), 7.45 (d, 1H, 6'-H), 6.43 (d, 1H, 5'-H), 6.32 (d, 1H, 3'-H), 2.93 (s, 3H, N-CH₃), 2.85 (s, 3H, N-CH₃); MS (EI 70 eV) *m/z* (%): 209 (4), 191 (4), 136 (66), 107 (28), 70 (100), 52 (84). Crystals suitable for an X-ray diffraction study were grown from acetone at 277 K.

Crystal data

C₁₀H₁₁NO₄
M_r = 209.20
 Monoclinic, *P*2₁/*c*
a = 9.3158 (13) Å
b = 6.4393 (9) Å
c = 17.399 (2) Å
 β = 92.022 (2)°
V = 1043.1 (2) Å³
Z = 4
D_x = 1.332 Mg m⁻³
 Mo Kα radiation
 Cell parameters from 1822 reflections
 θ = 2.5–26.5°
 μ = 0.10 mm⁻¹
T = 283 (2) K
 Block, yellow
 0.40 × 0.20 × 0.10 mm

Data collection

Bruker SMART 4K CCD area-detector diffractometer
 φ and ω scans
 Absorption correction: none
 5991 measured reflections
 2373 independent reflections
 1740 reflections with *I* > 2σ(*I*)
R_{int} = 0.025
 θ_{max} = 27.5°
h = -12 → 12
k = -8 → 7
l = -22 → 17

Refinement

Refinement on *F*²
R[*F*² > 2σ(*F*²)] = 0.049
wR(*F*²) = 0.148
S = 1.05
 2373 reflections
 141 parameters
 H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0777P)^2 + 0.1291P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 (Δ/σ)_{max} = 0.001
 Δρ_{max} = 0.22 e Å⁻³
 Δρ_{min} = -0.17 e Å⁻³

Table 1

Hydrogen-bonding geometry (Å, °).

<i>D</i> –H... <i>A</i>	<i>D</i> –H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> –H... <i>A</i>
O1–H1...O3	0.82	1.90	2.6158 (18)	146
O2–H2...O4 ⁱ	0.82	1.82	2.6333 (17)	174
C3–H3...O2 ⁱⁱ	0.93	2.60	3.422 (2)	148

Symmetry codes: (i) *x*, ½ - *y*, *z* - ½; (ii) 1 - *x*, *y* - ½, ½ - *z*.

H atoms were positioned geometrically and treated as riding, with O–H = 0.82 Å and C–H = 0.93 or 0.96 Å. *U*_{iso}(H) values were set

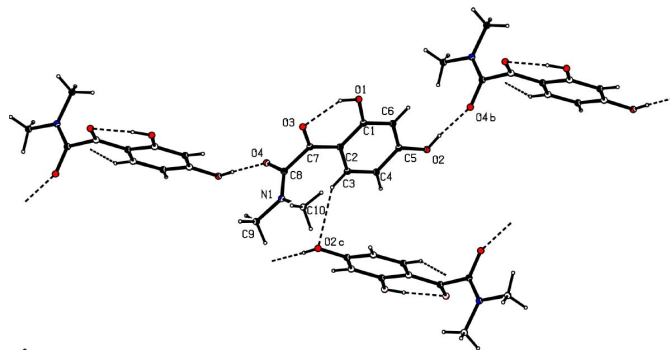


Figure 2

Hydrogen bonding in the crystal structure of (I). Hydrogen bonds are shown as dashed lines. [Symmetry codes: (b) *x*, -*y* + ½, *z* - ½; (c) -*x* + 1, *y* - ½, -*z* + ½]

equal to *xU*_{eq}(carrier atom), where *x* = 1.2 for *Csp*² and *x* = 1.5 for O and methyl C.

Data collection: SMART (Bruker, 1997); cell refinement: SAINT (Bruker, 1999); 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, 1997); software used to prepare material for publication: SHELXTL.

The authors acknowledge financial support from the National Key Project for Basic Research (No. 2002CCA00500), the National Natural Science Foundation of China (Nos. 20432010, 20476036 and 20172017), the Program for New Century Excellent Talents in Universities of China and the Program for Excellent Research Group of Hubei Province (No. 2004ABC002).

References

Bruker (1997). SMART. Version 5.054. Bruker AXS Inc., Madison, Wisconsin, USA.
 Bruker (1999). SAINT. Version 6.01. Bruker AXS Inc., Madison, Wisconsin, USA.
 Bruker (2001). SHELXTL. Version 6.12. Bruker AXS Inc., Madison, Wisconsin, USA.
 Sheldrick, G. M. (1997). SHELXS97 and SHELXL97. University of Göttingen, Germany.
 Yang, Z., Zhang, Z. X., Meanwell, N. A., Kadow, J. F. & Wang, T. (2002). Org. Lett. 7, 1103–1105.
 Zhou, T. & Chen, Z. C. (2001). J. Chem. Res. (S), pp. 116–117.