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#### **Key indicators**

Single-crystal X-ray study T = 298 KMean  $\sigma(\text{C-C}) = 0.006 \text{ Å}$  R factor = 0.060 wR factor = 0.156Data-to-parameter ratio = 7.0

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

# 7-Methoxy-4,6-dimethyl-3*H*-isobenzo-furan-1-one

The title compound,  $C_{11}H_{12}O_3$ , exists as a nearly planar molecule, the dihedral angle between the five- and six-membered rings being 1.9 (2)°.

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#### Comment

7-Methoxy-4,6-dimethyl-3*H*-isobenzofuran-1-one, (I), exists as a nearly planar molecule, the dihedral angle between the five- and six-membered rings being  $1.9 (2)^{\circ}$ . The compound exhibits moderate cytotoxicity towards the KB cell line (IC<sub>50</sub> 50 µg ml<sup>-1</sup>).

#### **Experimental**

The title compound was crystallized from the ethyl acetate extract of the liquid culture of an unidentified marine fungus isolated from the leaves of the sea lotus found in Fujian Province, China.

#### Crystal data

 $C_{11}H_{12}O_3$ Mo  $K\alpha$  radiation  $M_r = 192.21$ Cell parameters from 1595 Orthorhombic, Pna21 reflections a = 7.5508 (7) Å $\theta=2.6\text{--}21.8^\circ$  $\mu = 0.10 \text{ mm}^{-1}$ b = 12.616 (1) Åc = 10.1223 (9) ÅT = 298 (2) K $V = 964.27 (15) \text{ Å}^3$ Needle, colorless Z = 4 $0.45 \times 0.12 \times 0.09 \text{ mm}$  $D_x = 1.324 \text{ Mg m}^{-3}$ 

#### Data collection

Bruker APEX area-detector diffractometer  $R_{\rm int} = 0.070$   $\varphi$  and  $\omega$  scans  $\theta_{\rm max} = 25.0^{\circ}$  Absorption correction: none  $h = -8 \rightarrow 8$  12924 measured reflections  $k = -15 \rightarrow 14$  897 independent reflections  $l = -11 \rightarrow 12$ 

#### Refinement

 $\begin{array}{lll} \mbox{Refinement on } F^2 & w = 1/[\sigma^2(F_o^2) + (0.1P)^2 \\ R[F^2 > 2\sigma(F^2)] = 0.060 & + 0.1111P] \\ wR(F^2) = 0.156 & where $P = (F_o^2 + 2F_c^2)/3$ \\ S = 1.13 & (\Delta/\sigma)_{\rm max} = 0.001 \\ 897 \mbox{ reflections} & \Delta\rho_{\rm max} = 0.15 \mbox{ e Å}^{-3} \\ 129 \mbox{ parameters} & \Delta\rho_{\rm min} = -0.26 \mbox{ e Å}^{-3} \end{array}$ 

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 Table 1

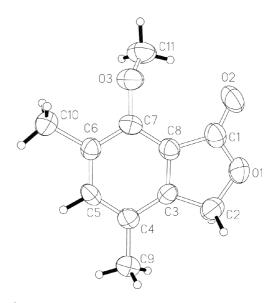
 Selected geometric parameters ( $\mathring{A}$ , °).

O1-C1	1.342 (6)	C3-C4	1.400 (7)
O1-C2	1.434 (7)	C4-C5	1.381 (7)
O2-C1	1.205 (6)	C4-C9	1.489 (7)
O3-C7	1.361 (6)	C5-C6	1.385 (6)
O3-C11	1.418 (7)	C6-C7	1.381 (6)
C1-C8	1.462 (6)	C6-C10	1.515 (6)
C2-C3	1.480 (7)	C7-C8	1.409 (6)
C3-C8	1.377 (7)		. ,
C1-O1-C2	110.3 (4)	C3-C4-C9	121.1 (4)
C7-O3-C11	117.7 (3)	C4-C5-C6	123.7 (4)
O2-C1-O1	120.4 (5)	C7-C6-C5	120.3 (4)
O2-C1-C8	130.7 (5)	C7-C6-C10	119.3 (4)
O1-C1-C8	108.8 (4)	C5-C6-C10	120.4 (4)
O1-C2-C3	105.2 (4)	O3-C7-C6	118.8 (4)
C8-C3-C4	122.4 (4)	O3-C7-C8	123.5 (4)
C8-C3-C2	108.0 (4)	C6 - C7 - C8	117.5 (4)
C4-C3-C2	129.6 (5)	C3-C8-C7	120.7 (4)
C5-C4-C3	115.3 (4)	C3-C8-C1	107.7 (4)
C5-C4-C9	123.6 (5)	C7-C8-C1	131.5 (4)

The H atoms were positioned geometrically and were included in the refinement in the riding-model approximation. C—H distances were set to 0.93–0.97 Å, with  $U_{\rm iso}$  values for H atoms of 1.2 or 1.5 (methyl H) times  $U_{\rm eq}$  of the parent atom. The methyl groups bonded to the aromatic ring were allowed to rotate but not to tip. In the absence of significant anomalous scattering effects, Friedel pairs were merged.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SMART*; data reduction: *SAINT* (Bruker, 2001); program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *ORTEP*II (Johnson, 1976); software used to prepare material for publication: *SHELXL*97.

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**Figure 1** *ORTEPII* (Johnson, 1976) plot of (I), with displacement ellipsoids drawn at the 50% probability level. H atoms are drawn as spheres of arbitrary radii.

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