

**6,7-Dimethoxy-2-methyl-3,4-dihydro-isoquinolin-1(2H)-one****Ikram M. Said, Norizan A. A. Hamid, Jalifah Latif, Laily B. Din and Bohari M. Yamin\***

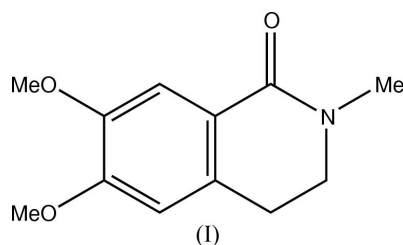
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Correspondence e-mail:  
bohari\_bob@yahoo.com**Key indicators**Single-crystal X-ray study  
 $T = 273$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.003$  Å  
 $R$  factor = 0.058  
 $wR$  factor = 0.145  
Data-to-parameter ratio = 15.8For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

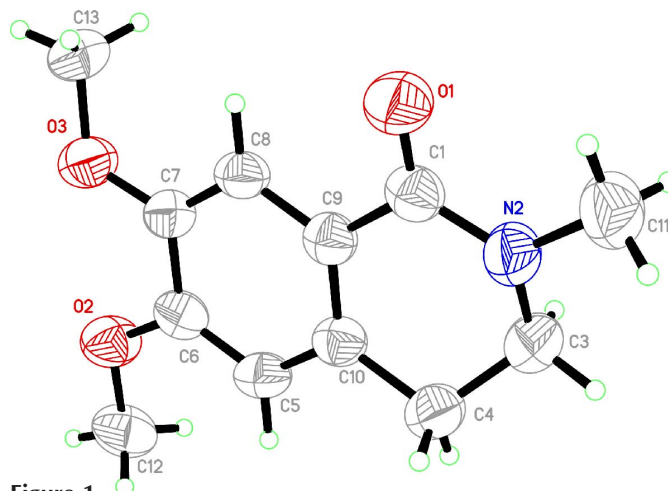
In the title compound,  $\text{C}_{12}\text{H}_{15}\text{NO}_3$ , the isoquinoline moiety is essentially planar, except for the N and adjacent  $\text{Csp}^3$  atoms, which deviate from the mean plane by 0.198 (2) and 0.686 (2) Å, respectively. The bond length between the N and carbonyl C atoms [1.353 (3) Å] is close to a double-bond value, indicating electron delocalization. No significant intermolecular interactions are observed.

**Comment**

Tetrahydroisoquinoline derivatives, obtained either from plants or by synthesis, are of great interest due to their biological and pharmacological properties (Brzezinska, 1994; Anderson *et al.*, 1998; Loesel *et al.*, 1987).



The title compound, (I) (Fig. 1), is an isoquinoline analogue of 6,7-dimethoxy-1-phenyl-1,2,3,4-tetrahydroisoquinoline, (II) (Olszak *et al.*, 1996), and 2-chlorobenzoxazolo[3,2-*b*]isoquinoline-6-one, (III) (Ravishankar *et al.*, 2001). Structural studies have shown that the degree of planarity of the isoquinoline moiety is affected by the presence of various substituents. For example, the molecule of (III) is essentially



**Figure 1**  
The molecular structure of (I), shown with 50% probability displacement ellipsoids and H atoms drawn as spheres of arbitrary radius.

Received 10 February 2005  
Accepted 22 February 2005  
Online 4 March 2005

planar (Ravishankar *et al.*, 2001), whereas in a dione system such as *N*-(2-chlorobenzyl)-1,2,3,4-tetrahydroisoquinoline-1,3-dione, the isoquinoline moiety is slightly folded about the fused C9–C10 bond (Pandi *et al.*, 2002). In contrast to (II), where the saturated fragment has a twist conformation, in (I) it has a half-chair or envelope conformation [ $Q = 0.430$  (2) Å,  $\theta = 116.8$  (3) and  $\varphi_2 = 100.1$  (3); Cremer & Pople, 1975], with C1–N2–C3–C4 and N2–C3–C4–C10 torsion angles of  $-42.8$  (3) and  $49.5$  (2)°, respectively. The C1/C4/C5/C6/C7/C8/C9/C10 fragment of the isoquinoline moiety is essentially planar, with a maximum deviation from the mean plane of 0.026 (2) Å for atom C6. Atoms N2 and C3 are displaced from that plane by 0.198 (2) and 0.686 (2) Å, respectively. As a result, the N2–C1 bond length [1.353 (3) Å] is shorter than those in (II) [1.461 (3) Å] and in (III) [1.408 (4) Å], indicating electron delocalization in (I) involving the O1–C1 and C1–N2 bonds. All other bond lengths and angles in (I) (Table 1) are in normal ranges (Allen *et al.*, 1987) and agree with those in other isoquinoline compounds. In contrast to the crystal structures of (II) and (III), no significant intermolecular interactions are observed in (I).

## Experimental

The roots of *Alseodaphne perakensis* were collected from Sungkai, Perak, and a voucher specimen (ALM 5735) was deposited at the UKMB herbarium of Universiti Kebangsaan Malaysia. The roots were air-dried, ground and soaked in MeOH for 48 h. The resulting extract was filtered and concentrated under reduced pressure to give 108.3 g of crude extract. The crude extract was acidified with 5% H<sub>2</sub>SO<sub>4</sub> and filtered. The acid extract was then basified with 5% Na<sub>2</sub>CO<sub>3</sub> until pH 8–9 was attained and extracted with CHCl<sub>3</sub> to obtain the crude alkaloid. The crude alkaloid (15.8 g) was chromatographed using liquid chromatography (VLC) and eluted with CHCl<sub>3</sub> and increasing amounts of MeOH. The VLC of the crude alkaloid yielded four fractions, denoted 01–04. Fraction 03 was further chromatographed using VLC with CHCl<sub>3</sub>–MeOH (7:3 v/v) as eluents to yield *N*-methylcorydaline. Colourless crystals were obtained from the chromatographed solution after 3 d upon evaporation at room temperature (m.p. 398–400 K). EIMS  $m/z$  (% int.): [ $M^+$ ] 221 (77), 178 (82), 150 (100), 135 (15), 107 (10), 92 (13). <sup>1</sup>H NMR:  $\delta$  2.88 (2H, *t*,  $J = 6.9$  Hz, H-4), 3.08 (3H, *s*, NCH<sub>3</sub>), 3.49 (2H, *t*,  $J = 6.9$  Hz, H-3), 3.86 (3H, *s*, OCH<sub>3</sub>-7), 3.87 (3H, *s*, OCH<sub>3</sub>-6), 6.58 (1H, *s*, H-5), 7.55 (1H, *s*, H-8), <sup>13</sup>C NMR:  $\delta$  27.5 (C4), 35.2 (C12), 48.4 (C3), \*56.07 (C12), \*56.14 (C13), 109.3 (C5), 110.5 (C8), 121.9 (C9), 131.6 (C10), 148.0 (C7), 152.0 (C6), 165.0 (C1) (\* values may be interchanged).

### Crystal data

C<sub>12</sub>H<sub>15</sub>NO<sub>3</sub>  
 $M_r = 221.25$   
 Monoclinic,  $P2_1/c$   
 $a = 9.7846$  (17) Å  
 $b = 16.865$  (3) Å  
 $c = 6.8914$  (12) Å  
 $\beta = 103.934$  (3)°  
 $V = 1103.7$  (3) Å<sup>3</sup>  
 $Z = 4$

$D_x = 1.331$  Mg m<sup>-3</sup>  
 Mo  $K\alpha$  radiation  
 Cell parameters from 912 reflections  
 $\theta = 2.1$ – $26.5^\circ$   
 $\mu = 0.10$  mm<sup>-1</sup>  
 $T = 273$  (2) K  
 Plate, colourless  
 0.31 × 0.11 × 0.06 mm

### Data collection

Bruker SMART APEX CCD area-detector diffractometer  
 $\omega$  scans  
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)  
 $T_{\min} = 0.971$ ,  $T_{\max} = 0.994$   
 6118 measured reflections

2285 independent reflections  
 1673 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.020$   
 $\theta_{\max} = 26.5^\circ$   
 $h = -10 \rightarrow 12$   
 $k = -21 \rightarrow 20$   
 $l = -6 \rightarrow 8$

### Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.058$   
 $wR(F^2) = 0.145$   
 $S = 1.09$   
 2285 reflections  
 145 parameters  
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0631P)^2 + 0.2375P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} < 0.001$   
 $\Delta\rho_{\max} = 0.24$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.15$  e Å<sup>-3</sup>

**Table 1**

Selected geometric parameters (Å, °).

O1–C1	1.227 (2)	N2–C11	1.451 (3)
O2–C6	1.359 (2)	N2–C3	1.458 (3)
O3–C7	1.364 (2)	C1–C9	1.486 (3)
N2–C1	1.353 (3)	C3–C4	1.497 (3)
C11–N2–C1–O1	–4.7 (3)	C3–N2–C1–C9	11.0 (3)
C3–N2–C1–O1	–170.6 (2)		

After the location of the H atoms in a difference map, their positions were idealized. The H atoms were allowed to ride on the parent C atoms, with C–H = 0.93–0.96 Å and  $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$  for CH<sub>3</sub> and  $1.2U_{\text{eq}}(\text{C})$  for CH groups.

Data collection: SMART (Siemens, 1996); cell refinement: SAINT (Siemens, 1996); data reduction: SAINT; program(s) used to solve structure: SHELXTL (Sheldrick, 1997); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL, PARST (Nardelli, 1995) and PLATON (Spek, 2003).

The authors thank the Malaysian Government for the research grant IRPA No. 09-02-02-0163.

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