

Methyl 1-*n*-butyl-2-(3,4-dichlorophenyl)-1*H*-benzimidazole-5-carboxylate

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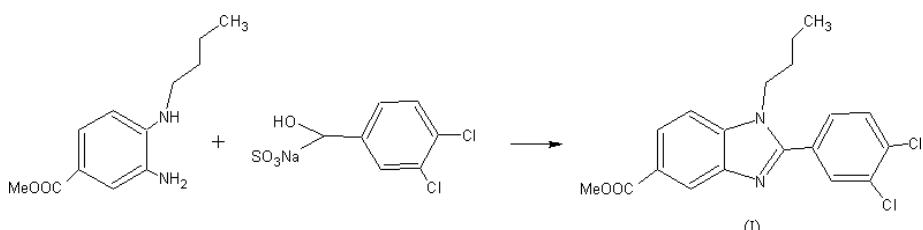
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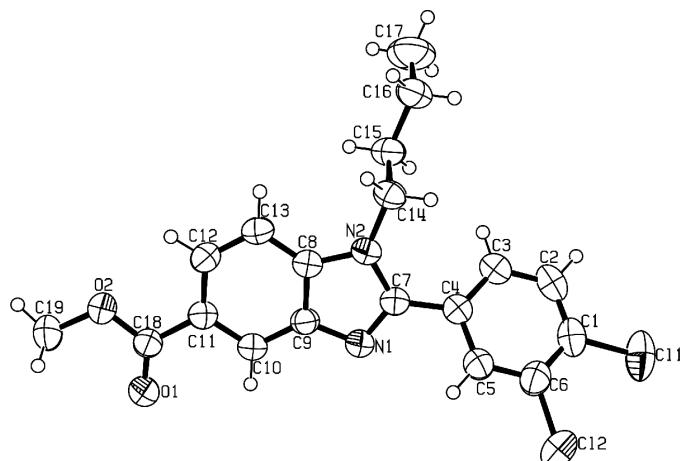
A new benzimidazole compound, methyl 1-*n*-butyl-2-(3,4-dichlorophenyl)-1*H*-benzimidazole-5-carboxylate, $C_{19}H_{18}Cl_2N_2O_2$, has been synthesized by the condensation of methyl 3-amino-4-(*n*-butylamino)benzoate with an $Na_2S_2O_5$ adduct of 3,4-dichlorobenzaldehyde. The molecule is twisted with a C—C—C—N torsion angle of $-39.7(3)^\circ$ between the phenyl and benzimidazole groups. In the crystal structure, symmetry-related molecules are linked by C—H···O interactions, forming a chain.

Comment

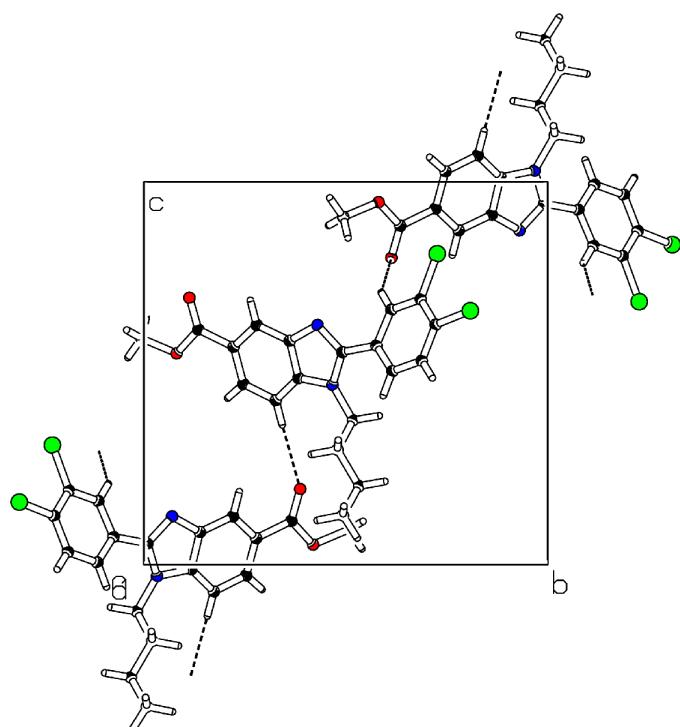
The benzimidazole ring system is of interest because of its diverse biological activities, including antifungal (Göker *et al.*, 2002), antibacterial (Weidner-Wells *et al.*, 2001), antiparasitic (Navarrete-Vazquez *et al.*, 2001), anticancer (Badawey & Kappe, 1999), anti-allergic (Nakano *et al.*, 2000), anti-ulcer (Göker & Düver, 1990) and antihypertensive (Matsumori, 2003). New drugs carrying a benzimidazole moiety, such as omeprazole (Göker & DüVer, 1990), candesartane (Matsumori, 2003) and mizolastine (Dubertret *et al.*, 1999), have been used clinically, and considerable effort has been invested recently to discover new potent agents (Mekapati & Hansch, 2001). From our laboratory, the synthesis and crystal structure analyses of several benzimidazoles have already been reported (Göker *et al.*, 1995, 1999; Özbeý *et al.*, 1998; Kendi *et al.*, 1999). The versatility of this ring system has prompted us to synthesize new analogs, including the title compound, (I).



The molecular structure of (I) is shown in Fig. 1 and selected bond distances and angles are given in Table 1. The dihedral angle between the plane of the ring defined by atoms N1/C7/N2/C8/C9 and the C1—C6 phenyl ring is $36.68(7)^\circ$, with a C3—C4—C7—N1 torsion angle of $-39.7(3)^\circ$. The molecule shows small deviations from planarity, the largest being $0.014(2)$ Å for atom C8 in the benzimidazole ring system and $0.015(4)$ Å for atom C1 in the C1—C6 phenyl ring. The C18=O1 bond length is $1.198(2)$ Å and the C19—O2—C18—C11 torsion angle is $178.78(17)^\circ$. In the molecule, the C—Cl bond lengths are very similar, Cl—Cl1 being $1.725(2)$ Å and C6—Cl2 being $1.729(2)$ Å.

**Figure 1**

An ORTEP-3 (Farrugia, 1997) view of (I), with the atomic numbering scheme and 50% probability displacement ellipsoids.

**Figure 2**

An ORTEP-3 (Farrugia, 1997) packing diagram of (I), viewed along the *a* axis. The C—H···O hydrogen bonds are shown as dashed lines.

In the crystal structure, symmetry-related molecules are connected by C—H···O hydrogen bonds, forming a polymer chain (see Table 2 and Fig. 2).

Experimental

To a suspension of methyl 3-amino-4-(*n*-butylamino)benzoate (0.22 g, 1 mmol) in dimethylformamide (1 ml), a sodium metabisulfite adduct of 3,4-dichlorobenzaldehyde (0.347 g, 1.25 mmol) was added and heated at 403 K for 4 h. The reaction mixture was cooled then poured into water. The solid product obtained was collected by filtration and washed with water. It was then chromatographed with EtOAc-*n*-hexane (1:3) (yield 0.2 g, 53%). Pale-green crystals of (I) were obtained (m.p. 353 K). IR (CO): 1706 cm⁻¹; ¹H NMR (DMSO-

*d*₆): δ 0.67 (*t*, 3H, CH₂CH₃), 1.03–1.09 (*m*, 2H, CH₂CH₃), 1.54–1.58 (*m*, 2H, CH₂CH₂), 3.79 (*s*, 3H, OCH₃), 4.26 (*t*, 2H, N—CH₂, *J* = 7.2 Hz), 7.70–7.8 (*m*, 3H, H-5,6,7), 7.84–7.87 (*dd*, 1H, H-6, *J*_o = 8.6, *J*_m = 1.4 Hz), 7.99 (*d*, 1H, H-2, *J*_m = 1.8 Hz), 8.20 (*d*, 1H, H-4, *J*_m = 1.2 Hz); MS (ES+): 377 (*M* + 1) (100%).

Crystal data

C ₁₉ H ₁₈ Cl ₂ N ₂ O ₂	<i>D</i> _x = 1.364 Mg m ⁻³
<i>M</i> _r = 377.25	Mo <i>K</i> α radiation
Monoclinic, <i>P</i> 2 ₁ / <i>n</i>	Cell parameters from 15013 reflections
<i>a</i> = 9.3359 (6) Å	<i>θ</i> = 1.5–29.0°
<i>b</i> = 14.4051 (8) Å	<i>μ</i> = 0.37 mm ⁻¹
<i>c</i> = 16.3707 (11) Å	<i>T</i> = 293 (2) K
<i>β</i> = 123.459 (4)°	Prismatic, pale green
<i>V</i> = 1836.8 (2) Å ³	<i>Z</i> = 4 0.50 × 0.30 × 0.10 mm

Data collection

Stoe IPDS-2 two-circle goniometer diffractometer	2727 reflections with <i>I</i> > 2σ(<i>I</i>)
<i>w</i> scans	<i>R</i> _{int} = 0.088
Absorption correction: none	<i>θ</i> _{max} = 27.5°
30007 measured reflections	<i>h</i> = -12 → 12
4133 independent reflections	<i>k</i> = -18 → 18
	<i>l</i> = -21 → 21

Refinement

Refinement on <i>F</i> ²	<i>w</i> = 1/[<i>σ</i> ² (<i>F</i> _o ²) + (0.0573 <i>P</i>) ² + 0.0182 <i>P</i>]
<i>R</i> [<i>F</i> ² > 2σ(<i>F</i> ²)] = 0.043	where <i>P</i> = (<i>F</i> _o ² + 2 <i>F</i> _c ²)/3
<i>wR</i> (<i>F</i> ²) = 0.117	(Δ/ <i>σ</i>) _{max} = 0.001
<i>S</i> = 1.04	Δρ _{max} = 0.29 e Å ⁻³
4133 reflections	Δρ _{min} = -0.29 e Å ⁻³
227 parameters	Extinction correction: <i>SHELXL97</i>
H-atom parameters constrained	Extinction coefficient: 0.0164 (18)

Table 1
Selected geometric parameters (Å, °).

Cl1—C1	1.725 (2)	O1—C18	1.198 (2)
Cl2—C6	1.729 (2)		
C3—C4—C7—N1	-39.7 (3)	C19—O2—C18—C11	178.78 (17)

Table 2
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>
C5—H5···O1 ⁱ	0.93	2.53	3.336 (3)	146
C13—H13···O1 ⁱⁱ	0.93	2.41	3.340 (2)	176

Symmetry codes: (i) $\frac{3}{2} - x, \frac{1}{2} + y, \frac{3}{2} - z$; (ii) $x - \frac{1}{2}, \frac{1}{2} - y, z - \frac{1}{2}$.

H atoms were included in calculated positions and treated as riding atoms; C—H = 0.93–0.97 Å and *U*_{iso}(H) = 1.2 or 1.5*U*_{eq}(C).

Data collection: *X*-AREA (Stoe & Cie, 1996); cell refinement: *X*-AREA; data reduction: *X*-RED32 (Stoe & Cie, 1996); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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