

Data collection

Enraf–Nonius CAD-4 diffractometer
 $\omega/2\theta$ scans
 Absorption correction: empirical ψ scan (North, Phillips & Mathews, 1968)
 $T_{\min} = 0.046$, $T_{\max} = 0.069$
 3202 measured reflections
 3187 independent reflections

2300 reflections with $I > 2.5\sigma(I)$
 $R_{\text{int}} = 0.028$
 $\theta_{\text{max}} = 24.97^\circ$
 $h = 0 \rightarrow 12$
 $k = 0 \rightarrow 17$
 $l = -13 \rightarrow 12$
 3 standard reflections every 200 reflections
 intensity decay: none

Refinement

Refinement on F^2
 $R = 0.053$
 $wR = 0.061$
 $S = 1.905$
 2300 reflections
 208 parameters
 H atoms not refined
 Weighting scheme based on measured s.u.'s; $w = 1/[\sigma_c^2(F_o) + (p^2/4F_o^2)]$, with $p = 0.03$

$(\Delta/\sigma)_{\text{max}} = 0.0034$
 $\Delta\rho_{\text{max}} = 1.41 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.69 \text{ e } \text{\AA}^{-3}$
 Extinction correction: none
 Scattering factors from *International Tables for X-ray Crystallography* (Vol. IV)

Table 1. Selected geometric parameters (\AA , $^\circ$)

Br(1)—C(8)	1.972 (5)	N(1)—C(1)	1.306 (7)
Br(2)—C(9)	2.007 (6)	N(1)—C(7)	1.385 (7)
Cl—C(11)	1.736 (5)	C(1)—C(8)	1.492 (7)
S—C(1)	1.729 (5)	C(8)—C(9)	1.476 (8)
S—C(2)	1.711 (5)	C(9)—C(10)	1.504 (7)
C(1)—S—C(2)	89.5 (3)	Br(1)—C(8)—C(1)	110.4 (4)
C(1)—N(1)—C(7)	109.9 (4)	Br(1)—C(8)—C(9)	106.2 (4)
S—C(1)—N(1)	116.2 (4)	C(1)—C(8)—C(9)	113.2 (5)
S—C(1)—C(8)	120.4 (4)	Br(2)—C(9)—C(8)	107.0 (4)
N(1)—C(1)—C(8)	123.3 (5)	Br(2)—C(9)—C(10)	106.7 (4)
S—C(2)—C(3)	129.2 (5)	C(8)—C(9)—C(10)	117.0 (5)
S—C(2)—C(7)	109.4 (4)	C(9)—C(10)—C(11)	122.2 (5)
N(1)—C(7)—C(2)	114.9 (5)	C(9)—C(10)—C(15)	120.5 (4)
N(1)—C(7)—C(6)	125.2 (5)		

The largest peak in the final difference Fourier map ($1.41 \text{ e } \text{\AA}^{-3}$) is 0.39 \AA from the Br(2) atom.

Data collection: *CAD-4-PC Software* (Enraf–Nonius, 1992). Cell refinement: *CAD-4-PC Software*. Data reduction: *TEXSAN* (Molecular Structure Corporation, 1985, 1992). Program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1993). Program(s) used to refine structure: *TEXSAN*. Software used to prepare material for publication: *TEXSAN*.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: SX1024). Services for accessing these data are described at the back of the journal.

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***N*-(*o*-Tolyl)-2-bromo-4,5-dimethoxybenzamide†**

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Abstract

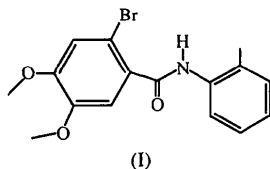
The molecular structure of the title compound, $\text{C}_{16}\text{H}_{16}\text{BrNO}_3$, consists of 2-bromo-4,5-dimethoxybenzene and *o*-methylphenyl rings linked by an amide group. The two methoxy groups are almost coplanar with the phenyl ring.

Comment

A number of antipsychotic agents of the *o*-methoxybenzamide type show interesting selectivity for dopamine–D2 receptors (Hogberg, 1991). Benzamides with an *N*-(*cis*-2-phenylcycloheptyl) or an *N*-ethyl-2-(pyrrolidine)methyl side chain and a series of

† Alternative name: 2-bromo-4,5-dimethoxy-2'-methylbenzamide.

5-substituted dimethoxybenzamides also display a high stereoselective affinity for dopamine-D2 receptors (Hogberg *et al.*, 1987; Hogberg, 1991). During our studies on the organic synthesis of new antipsychotic drugs, we obtained the title compound, (I), and in order to evaluate its bioactivity on the basis of structure-property relationships, we determined its X-ray crystal structure.



An ORTEPII (Johnson, 1976) representation of the title compound is presented in Fig. 1. The structure consists of 2-bromo-4,5-dimethoxybenzene and *o*-methylphenyl rings linked by an amide group. The dihedral angle between the two aromatic rings is 74.3 (8)°. The two methoxy groups are almost coplanar with the phenyl ring [torsion angles C(3)—C(4)—O(1)—C(7) -6.6 (6) and C(6)—C(5)—O(2)—C(8) 6.8 (5)°]. All the other bond distances and angles are in the normal ranges.

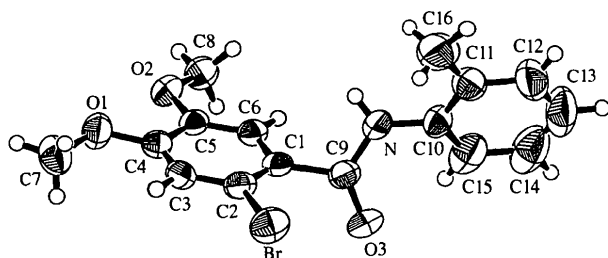


Fig. 1. An ORTEPII (Johnson, 1976) representation of the title compound showing 50% probability displacement ellipsoids.

Experimental

The title compound was synthesized by the reaction of bromo-veratric acid chloride with *o*-toluidine. The crude product was recrystallized from acetone to give white single crystals (m.p. 462–464 K) of X-ray quality.

Crystal data

C₁₆H₁₆BrNO₃

M_r = 350.21

Monoclinic

*P*2₁/*n*

a = 9.022 (3) Å

b = 11.615 (4) Å

c = 14.787 (6) Å

β = 99.44 (4)°

V = 1528.6 (9) Å³

Z = 4

D_x = 1.522 Mg m⁻³

D_m not measured

Mo Kα radiation

λ = 0.7107 Å

Cell parameters from 25 reflections

θ = 14–16°

μ = 2.706 mm⁻¹

T = 295 (1) K

Prism

0.40 × 0.26 × 0.22 mm

Colorless

Data collection

Enraf-Nonius CAD-4 diffractometer

ω/2θ scans

Absorption correction:

empirical ψ scan (North, Phillips & Mathews, 1968)

*T*_{min} = 0.379, *T*_{max} = 0.551

3048 measured reflections

3031 independent reflections

2146 reflections with

I > 2.5σ(*I*)

*R*_{int} = 0.021

θ_{max} = 24.97°

h = 0 → 10

k = 0 → 13

l = -17 → 17

3 standard reflections

every 200 reflections

intensity decay: none

Refinement

Refinement on *F*

R = 0.069

wR = 0.072

S = 2.425

2146 reflections

190 parameters

H atoms not refined

Weighting scheme based

on measured s.u.'s;

$$w = 1/[\sigma^2(F_{\text{obs}}) + 0.0026|F_{\text{obs}}|^2]$$

(Δ/σ)_{max} = 0.0212

Δρ_{max} = 1.28 e Å⁻³

Δρ_{min} = -0.54 e Å⁻³

Extinction correction: none

Scattering factors from International Tables for X-ray Crystallography (Vol. IV)

Table 1. Selected geometric parameters (Å, °)

Br—C(2)	1.900 (4)	O(2)—C(8)	1.410 (6)
O(1)—C(4)	1.358 (5)	O(3)—C(9)	1.209 (5)
O(1)—C(7)	1.413 (6)	N—C(9)	1.347 (5)
O(2)—C(5)	1.370 (5)	N—C(10)	1.410 (6)
C(4)—O(1)—C(7)	117.9 (4)	O(2)—C(5)—C(4)	115.4 (4)
C(5)—O(2)—C(8)	116.9 (3)	O(2)—C(5)—C(6)	124.8 (4)
C(9)—N—C(10)	125.2 (4)	O(3)—C(9)—N	124.9 (4)
Br—C(2)—C(1)	121.0 (3)	O(3)—C(9)—C(1)	122.2 (4)
Br—C(2)—C(3)	117.1 (3)	N—C(9)—C(1)	113.0 (3)
O(1)—C(4)—C(3)	125.8 (4)	N—C(10)—C(11)	117.9 (4)
O(1)—C(4)—C(5)	114.6 (4)	N—C(10)—C(15)	120.4 (5)
C(3)—C(4)—C(5)	119.6 (4)		

The largest peak in the final difference Fourier map (1.28 e Å⁻³) is 0.47 Å from the Br atom.

Data collection: CAD-4-PC Software (Enraf-Nonius, 1992). Cell refinement: CAD-4-PC Software. Data reduction: TEXSAN (Molecular Structure Corporation, 1985, 1992). Program(s) used to solve structure: SIR92 (Altomare *et al.*, 1993). Program(s) used to refine structure: TEXSAN. Software used to prepare material for publication: TEXSAN.

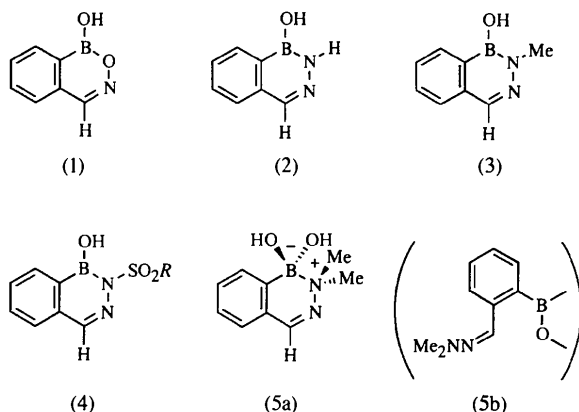
This work was supported by the National Science Foundation (OSR-9452893 and RII-8610677) and the National Institutes of Health (MBRS5S06-GM08224).

Supplementary data for this paper are available from the IUCr electronic archives (Reference: SX1023). Services for accessing these data are described at the back of the journal.

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A 2-Alkyl Substituted 2,3,1-Benzodiazaborine

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Abstract

The structure of the fourth member of a family of structurally related 2,3,1-benzodiheteroborines has been solved by crystallographic means, providing data for proper direct comparison with the others. The title compound, 1,2-dihydro-1-hydroxy-2-methyl-2,3,1-benzodiazaborine (C₈H₉BN₂O) is obtained *via* condensation of 2-formylbenzeneboronic acid and methylhydrazine, and is similar in most respects to its 2-unsubstituted parent, in internal geometry and intramolecular association topography, but has some subtle oxazaborine-like characteristics.

Comment

Although it has been known for quite some time that condensation of 2-formylbenzeneboronic acid with hydroxylamines or hydrazines affords 2,3,1-benzoxazaborine or benzodiazaborines, respectively (Tschampel & Snyder, 1964; Dewar & Dougherty, 1964), these and related boron heterocycles have until recently received little attention. We are interested in them as they are quite robust and potentially useful as platforms for the construction of new biologically active compounds (see, for example, Groziak *et al.*, 1994). To develop the 2,3,1-benzodiheteroborines further, we are undertaking a systematic physicochemical study of heterocycles like compounds (1)–(5) by crystallographic and other means.

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The parent oxaza- and diazaborines, (1) and (2), respectively, were the subjects of our most recent report (Groziak *et al.*, 1997), which among other determinations reported their crystal structures. The structure of the title compound (3), which is the 2-methyl-substituted version of (2), is presented here, while that of a representative 2-alkyl- or arylsulfonylated version, (4), is currently being sought. Although a crystal structure determination of a thiophene analog of (3), 7-hydroxy-6-methyl-7,6-borazarothieno[3,2-*c*]pyridine, has been reported (Aurivillius & Löefving, 1974), we solved the structure of (3) in order to compare it directly with (1) and (2) without the need to allow for differences in ring-fusion identity. Interestingly, the thiophene analog reportedly decomposed in air of normal humidity so rapidly that the crystal for analysis had to be kept over a saturated aqueous solution of the substance. No such unusual lability was noted for our crystalline (3).

The 2,3,1-benzodiazaborines in particular can exhibit a wide diversity of structures. Some of the known biocidal (Grassberger *et al.*, 1984) heterocycles, (4), have been found by crystallography to form a tetrahedral borate complex in the active site of enoyl reductase (Baldock *et al.*, 1996) and the 2-dimethyl-substituted 2,3,1-benzodiazaborine, (5a), exists as a zwitterion, but only in water. An unusual double intramolecular chelate form of triarylboroxine (5b) was found to be its structure in the solid state (Robinson *et al.*, 1996).

An *ORTEP* (Johnson, 1965) view of the molecule of crystalline (3), together with its atom numbering, is given in Fig. 1. This molecule is a true B—N for C=C replacement analog of the enol form of 3-methyl-3*H*-isoquinolin-4-one, but to our knowledge no simple 3-alkyl-4-hydroxyisoquinoline has yet been studied crystallographically. Instead, the closest nitrogen heterocycle reference compounds for (3) appear to be several 3,4-dihydro-4-oxo-3-(aryl-methyl)-1-phthalazineacetic acids (Mylari *et al.*, 1991) and a 4-(*p*-chlorobenzyl)-2-(alkyl)-1(2*H*)-phthalazinone monohydrate (Scheffler *et al.*, 1988). Features associated with the benzene ring and methyl substituent of (3) are