

C. R. Girija and
Noor Shahina Begum*Department of Studies in Chemistry, Central
College Campus, Bangalore University,
Bangalore 560 001, Karnataka, India

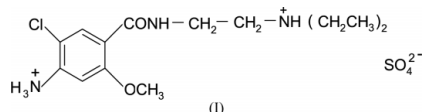
Correspondence e-mail: noorsb@rediffmail.com

Key indicators

Single-crystal X-ray study
 $T = 293\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.006\text{ \AA}$
 R factor = 0.057
 wR factor = 0.178
Data-to-parameter ratio = 12.2For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.4-Ammonio-5-chloro-*N*-[2-(*N,N*-diethyl-
ammonio)ethyl]-2-methoxybenzamide
sulfateThe title compound, $\text{C}_{14}\text{H}_{24}\text{ClN}_3\text{O}_2^{2+}\cdot\text{SO}_4^{2-}$, also known as metoclopramide sulfate, is non-planar, both the molecular and crystal structures being stabilized by $\text{N}-\text{H}\cdots\text{O}$ and $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds.Received 9 December 2003
Accepted 22 December 2003
Online 23 January 2004

Comment

The title compound, (I), a substituted benzamide, is a dopamine antagonist clinically used as neuroleptic drug (Pitre & Stradi, 1987). It is also a potential ligand, owing to its significant tendency to coordinate to metals. Its crystal structure is reported here.



In the title molecule (Fig. 1), the $\text{C}=\text{O}$ group is almost *cis* with respect to the $\text{C}1-\text{C}6$ bond, the torsion angle $\text{C}6-\text{C}1-\text{C}11-\text{O}12$ being $-5.5(5)^\circ$. The methoxy group is coplanar with the attached ring. A *gauche* conformation is observed in an amino ethyl side chain. The geometrical parameters of the molecule are comparable with those in a similar structure (Blaton & Peeters, 1980). An intramolecular hydrogen bond is formed between methoxy O atom O7 and amide H atom H13 (Table 2). This hydrogen bond creates a virtual ring which may be a key feature for the binding of neuroleptic benzamides to the dopamine receptor (van de Waterbeemd & Testa, 1981). The molecular structure is further stabilized by $\text{N}16-\text{H}16\cdots\text{O}12$ and $\text{C}6-\text{H}6\cdots\text{O}12$ intramolecular hydrogen bonds. In the crystal structure, the cation and anions

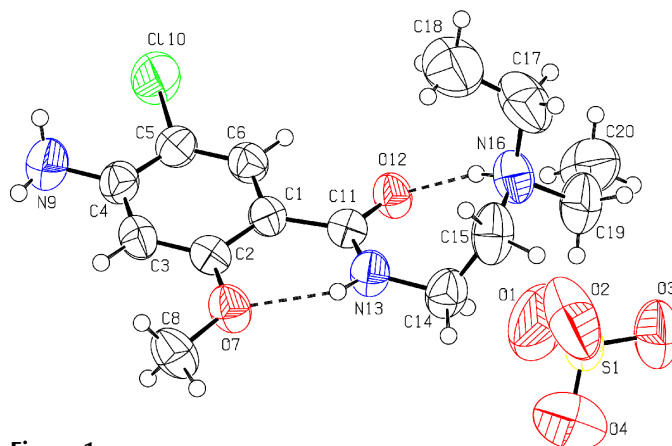


Figure 1

Molecular structure of the title compound, showing 50% probability displacement ellipsoids and the atom-numbering scheme.

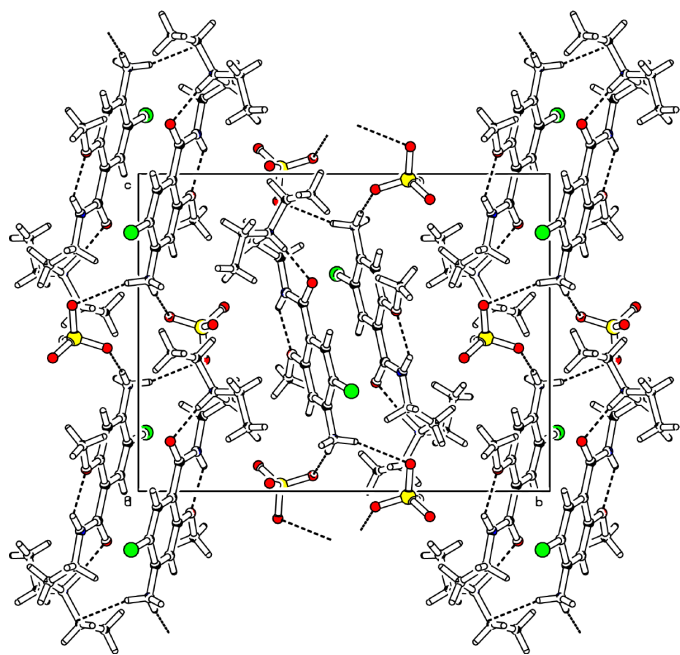


Figure 2
Packing of the title molecules, viewed down the *a* axis.

are involved in N—H···O and C—H···O hydrogen bond (Table 2). Further, Cl10···O12(1 - *x*, 1 - *y*, 1 - *z*) short contacts [3.262 (3) Å] link inversion-related molecules into a dimer (Fig. 2). A short H9C···Cl10 (2.90 Å) interaction is also observed.

Experimental

The title compound (98% pure) was bought from IPCA laboratories. Single crystals were grown from a methanol solution by slow evaporation at room temperature.

Crystal data

$C_{14}H_{24}ClN_3O_2^{2+} \cdot SO_4^{2-}$
 $M_r = 397.88$
 Monoclinic, $P2_1/c$
 $a = 9.1514$ (2) Å
 $b = 16.5218$ (3) Å
 $c = 12.8524$ (3) Å
 $\beta = 97.186$ (1)°
 $V = 1927.99$ (7) Å³
 $Z = 4$

$D_x = 1.371$ Mg m⁻³
 Mo $K\alpha$ radiation
 Cell parameters from 7814 reflections
 $\theta = 2.0$ – 23.3 °
 $\mu = 0.34$ mm⁻¹
 $T = 293$ (2) K
 Prism, colourless
 $0.30 \times 0.25 \times 0.10$ mm

Data collection

Bruker SMART CCD area-detector diffractometer
 ω scans
 Absorption correction: none
 7814 measured reflections
 2768 independent reflections

1932 reflections with $I > 2\sigma(I)$
 $R_{int} = 0.034$
 $\theta_{max} = 23.3$ °
 $h = -10 \rightarrow 10$
 $k = -14 \rightarrow 18$
 $l = -13 \rightarrow 14$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.057$
 $wR(F^2) = 0.179$
 $S = 1.03$
 2768 reflections
 227 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0961P)^2 + 1.029P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{max} = 0.004$
 $\Delta\rho_{max} = 0.56$ e Å⁻³
 $\Delta\rho_{min} = -0.28$ e Å⁻³
 Extinction correction: *SHELXL97*
 Extinction coefficient: 0.0074 (19)

Table 1

Selected geometric parameters (Å, °).

S1—O1	1.360 (5)	O7—C8	1.428 (5)
S1—O2	1.372 (4)	C11—O12	1.251 (4)
S1—O4	1.389 (5)	C11—N13	1.325 (4)
S1—O3	1.390 (3)	N13—C14	1.456 (5)
C1—C11	1.485 (5)	C15—N16	1.504 (6)
C2—O7	1.350 (4)	N16—C17	1.486 (6)
C4—N9	1.376 (5)	N16—C19	1.532 (6)
C5—Cl10	1.748 (4)		
C2—O7—C8	119.5 (3)	C17—N16—C19	111.6 (4)
C11—N13—C14	123.4 (3)	C15—N16—C19	109.5 (4)
C17—N16—C15	114.4 (4)		
C3—C2—O7—C8	5.4 (5)	C14—C15—N16—C17	150.6 (4)
C6—C1—C11—O12	-5.5 (5)	C14—C15—N16—C19	-83.3 (4)
C1—C11—N13—C14	174.7 (4)	C15—N16—C17—C18	-73.7 (6)
C11—N13—C14—C15	73.5 (5)	C15—N16—C19—C20	159.5 (5)
N13—C14—C15—N16	-78.0 (5)		

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>
N13—H13···O7	0.86	2.00	2.658 (4)	132
N16—H16···O12	0.91	1.79	2.665 (4)	160
C6—H6···O12	0.93	2.39	2.735 (4)	102
N9—H9A···O2 ⁱ	0.89	2.45	3.325 (7)	169
N9—H9B···O3 ⁱⁱ	0.89	2.51	3.372 (6)	164

Symmetry codes: (i) 2 - *x*, 1 - *y*, 1 - *z*; (ii) *x*, *y*, 1 + *z*.

H atoms were placed at calculated positions (N—H = 0.86–0.91 Å and C—H = 0.93–0.97 Å) and allowed to ride on the atoms to which they are bonded; $U_{iso}(H)$ values were set at $1.5U_{eq}(\text{parent})$ for CH₃ and NH₃ H atoms and $1.2U_{eq}(\text{parent})$ for the remaining H atoms. The highest peak and deepest hole in the final difference map were located at 1.39 and 1.09 Å, respectively, from atom O1. The large U_{eq} values for atoms O1, O2 and O4 indicate a possible disorder in the sulfate anion.

Data collection: *SMART* (Siemens, 1993); cell refinement: *SMART*; data reduction: *SAINT* (Siemens, 1995); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 1990); software used to prepare material for publication: *SHELXL97*.

CRG thanks UGC–FIP for a Teacher fellowship. The authors thank Dr G. U. Kulkarni, JNCASR, for data collection.

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