

Twinned 1-[2-[bis(4-fluorophenyl)methoxy]ethyl]-4-(3-phenyl)propyl)piperazinium chloride (GBR12909)

Judith L. Flippen-Anderson,^{a*}
Jeffrey R Deschamps,^a Clifford
George,^a John E. Folk,^b Arthur E.
Jacobson^b and Kenner C. Rice^b^aLaboratory for the Structure of Matter, Code
6030, Naval Research Laboratory, Washington,
DC 20375, USA, and ^bLaboratory of Medicinal
Chemistry, Building 8, Room B1-23, National
Institute of Diabetes and Digestive and Kidney
Diseases, National Institutes of Health, 8 Center
Drive MSC 0815, Bethesda, MD 20892, USACorrespondence e-mail:
flippen@harker.nrl.navy.mil

Key indicators

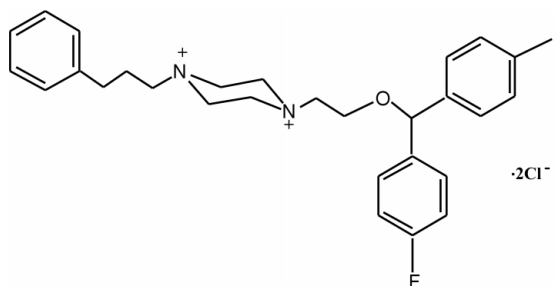
Single-crystal X-ray study
T = 293 K
Mean $\sigma(\text{C}-\text{C})$ = 0.008 Å
R factor = 0.065
wR factor = 0.190
Data-to-parameter ratio = 14.2For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.GBR12909, C₂₈H₃₄F₂N₂O, is one of the earliest agents found
to have high affinity and selectivity for the dopamine
transporter (DAT). The triclinic crystal, of the hydrochloride
salt of GBR12909, C₂₈H₃₄F₂N₂O²⁺·2Cl⁻, exhibited non-
merohedral twinning.

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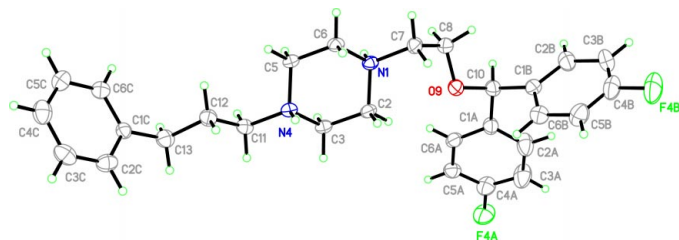
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Comment

Cocaine is one of the most widely abused drugs in the world.
The reinforcing and locomotor stimulating properties of
cocaine are said to be mainly mediated by the binding of
cocaine to the dopamine transporter (DAT) and subsequent
blockade of dopamine re-uptake into presynaptic terminals,
resulting in increased neurotransmission in the mesolimbic
dopaminergic system (Kuhar *et al.*, 1991; Rothman *et al.*,
1989). Studies have been aimed at the discovery and devel-
opment of potent and selective DAT ligands, (Zhang *et al.*,
2000; Newman *et al.*, 1994) that may reduce the effects of
cocaine and, therefore, could be used for the treatment of
cocaine abuse (Glowa *et al.*, 1995*a,b*). GBR12909, (I), a potent
if not overly selective DAT ligand, is now being examined in
man for its toxicity and efficacy prior to exploration of its
potential as a cocaine-treatment agent. The X-ray structure of
GBR12909 will provide some of the necessary information for
structure–activity studies that might lead to a new generation
of more potent and selective cocaine-treatment agents.

(I)

GBR12909 hydrochloride crystallizes with two molecules
per asymmetric unit. The two independent molecules are
almost identical, the only difference being the relative twist of
the two fluorophenyl groups (Fig. 1). The difference between
the C1B–C10–C1A–C2A torsion angle and the corre-
sponding torsion in the second molecule in the asymmetric
unit is approximately 16°. The central six-membered ring has a
normal chair conformation. Both N atoms in the ring are
positively charged and hydrogen bond to a Cl ion (see Table 1).
There are no other intermolecular interactions. Both aromatic
moieties are extended away from the central ring and from


Figure 1

View of GBR12909 with 20% probability ellipsoids, showing one of the two independent molecules in the asymmetric unit.

each other. The 4-fluorophenyl rings are approximately perpendicular to one another (angle between the ring planes of 88.5 (1) and 88.6 (2)° for the two independent molecules). The third aromatic ring is approximately perpendicular to the plane formed by N4, C11, C12 and C13 [82.3 (3) and 82.2 (2)° for the two independent molecules].

Experimental

The sample of GBR12909 hydrochloride was obtained from the National Institutes of Health. Crystals were grown at NRL by slow evaporation from a mixture of 2-propanol and ethyl acetate.

Crystal data

$C_{28}H_{34}F_2N_2O^{2+} \cdot 2Cl^-$
 $M_r = 523.47$
 Triclinic, $P\bar{1}$
 $a = 8.5449$ (4) Å
 $b = 14.3779$ (7) Å
 $c = 23.2700$ (12) Å
 $\alpha = 83.131$ (3)°
 $\beta = 80.554$ (3)°
 $\gamma = 89.963$ (3)°
 $V = 2799.4$ (2) Å³

$Z = 4$
 $D_x = 1.242$ Mg m⁻³
 Cu $K\alpha$ radiation
 Cell parameters from 5787 reflections
 $\theta = 3.1$ – 61.4 °
 $\mu = 2.38$ mm⁻¹
 $T = 293$ (2) K
 Plate, colorless
 $0.40 \times 0.35 \times 0.02$ mm

Data collection

Bruker SMART 6000 diffractometer
 ω scans
 Absorption correction: multi-scan (SADABS; Bruker, 2001)
 $T_{\min} = 0.386$, $T_{\max} = 0.953$
 9044 measured reflections
 9044 independent reflections

6598 reflections with $I > 2\sigma(I)$
 $\theta_{\max} = 67.2$ °
 $h = -10 \rightarrow 10$
 $k = -16 \rightarrow 14$
 $l = -26 \rightarrow 25$
 56 standard reflections
 intensity decay: none

Refinement

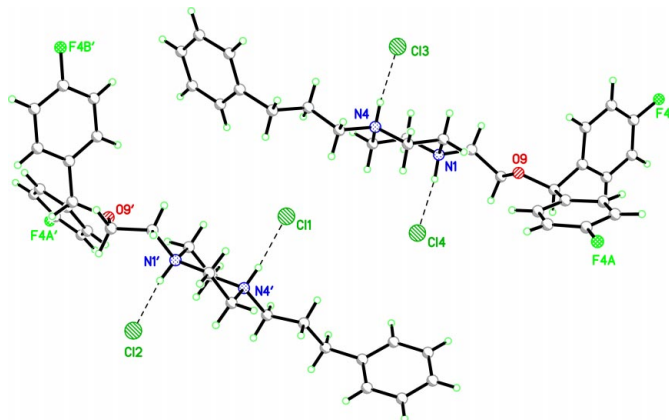
Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.065$
 $wR(F^2) = 0.190$
 $S = 0.97$
 9044 reflections
 635 parameters
 H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.128P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.047$
 $\Delta\rho_{\max} = 0.42$ e Å⁻³
 $\Delta\rho_{\min} = -0.26$ e Å⁻³
 Extinction correction: SHELXL97
 Extinction coefficient: 0.0015 (3)

Table 1

Hydrogen-bonding geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$N1-H1A \cdots Cl4$	0.91	2.14	3.034 (3)	166
$N4-H4A \cdots Cl3$	0.91	2.11	3.012 (3)	174
$N1'-H1'A \cdots Cl2$	0.91	2.13	3.027 (3)	167
$N4'-H4'A \cdots Cl1$	0.91	2.11	3.018 (3)	173


Figure 2

The asymmetric unit, showing each independent molecule hydrogen bonded to two Cl atoms.

The GBR12909 hydrochloride crystals exhibited non-merohedral twinning. Program *ROTAX*, found in the *CRYSTALS* (Watkin, 2001) suite of programs was used to determine the twin law $(-1, 0, 0/0, 1, 0/0, 0, 4, -1)$. Program *ROTWIN* (Young, 2001) was used to generate a new reflection file for refinement with *SHELXTL* (Bruker, 2001). Without twinning the refinement converged to an R factor of 0.094. Adding the single twin reduced the R factor to 0.065.

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

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