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Key indicators

Single-crystal X-ray study
 T = 291 K
 Mean $\sigma(C-C)$ = 0.005 Å
 R factor = 0.042
 wR factor = 0.113
 Data-to-parameter ratio = 14.6

For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

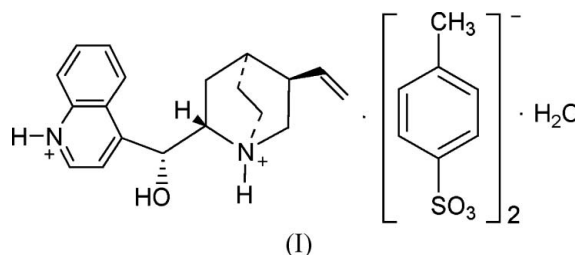
Cinchonidinium bis(4-methylbenzenesulfonate) monohydrate

In the title salt, $C_{19}H_{24}N_2O^{2+} \cdot 2C_7H_7SO_3^- \cdot H_2O$, the ions and the water molecule are held together by O—H...O, N—H...O and C—H...O hydrogen bonds, forming hydrogen-bonded layers; there are no hydrogen bonds between layers.

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Comment

Various salts of cinchonidine have been reported previously, e.g. cinchonidinium pyruvate oxime (Borszky *et al.*, 1996), cinchonidinium (*S*)-mandelate (Gjerløv & Larsen, 1997), bis(cinchonidinium) *L*-tartrate dihydrate (Zhang *et al.*, 2003), cinchonidinium (*R,R*)-tartrate monohydrate (Ryttersgaard & Larsen, 2003), cinchonidinium tris(tetrachlorobenzene-diolato)phosphate (Favarger *et al.*, 2004) and cinchonidinium bis(perchlorate) (Zhang *et al.*, 2006). As a continuation of our research, we report here the title salt, (I).



As shown in Fig. 1, the 4-methylbenzenesulfonate anion containing atom S1, the cinchonidinium cation and the water molecule are held together by N—H...O, O—H...O and C—H...O hydrogen bonds (Table 2) in an $R_3^3(9)$ ring (Etter, 1990).

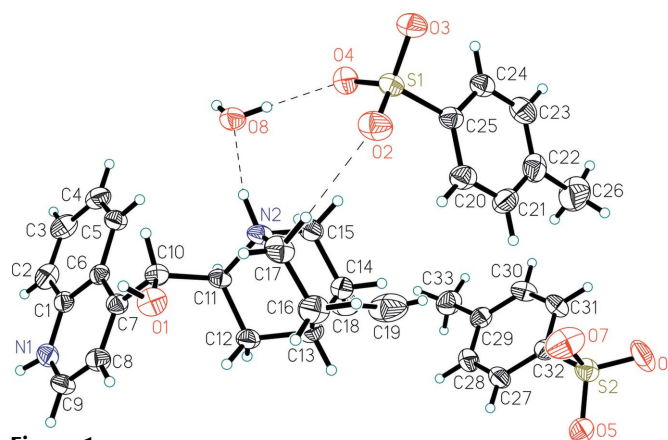


Figure 1
 The asymmetric unit of (I), showing the atom labelling and 40% probability displacement ellipsoids. Dashed lines indicate hydrogen bonds.

Further N—H···O, O—H···O and some weak C—H···O hydrogen bonds (Table 2) form a layer parallel to the (001) plane (Fig. 2). The whole structure is generated through translation of the layer; there are no hydrogen bonds between layers.

Experimental

Cinchonidine and 4-methylbenzenesulfonic acid in a 1:3 molar ratio were mixed and dissolved with stirring in sufficient water by heating to 373 K, at which point a clear solution resulted. The solution was cooled to room temperature and the colourless crystals were collected after 30 d.

Crystal data

$C_{19}H_{24}N_2O^{2+} \cdot 2C_7H_7SO_3^- \cdot H_2O$	$V = 790.6 (6) \text{ \AA}^3$
$M_r = 656.81$	$Z = 1$
Triclinic, $P1$	$D_x = 1.380 \text{ Mg m}^{-3}$
$a = 6.565 (3) \text{ \AA}$	Mo $K\alpha$ radiation
$b = 9.599 (4) \text{ \AA}$	$\mu = 0.22 \text{ mm}^{-1}$
$c = 12.748 (5) \text{ \AA}$	$T = 291 (2) \text{ K}$
$\alpha = 85.25 (3)^\circ$	Block, colourless
$\beta = 84.08 (3)^\circ$	$0.54 \times 0.42 \times 0.26 \text{ mm}$
$\gamma = 82.71 (3)^\circ$	

Data collection

Siemens P4 diffractometer	6198 independent reflections
ω scans	5316 reflections with $I > 2\sigma(I)$
Absorption correction: ψ scan	$R_{int} = 0.000$
(<i>XSCANS</i> ; Siemens, 1994)	$\theta_{max} = 26.0^\circ$
$T_{min} = 0.932$, $T_{max} = 0.994$	3 standard reflections
(expected range = 0.884–0.943)	every 97 reflections
6402 measured reflections	intensity decay: 4.9%

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.066P)^2 + 0.0459P]$
$R[F^2 > 2\sigma(F^2)] = 0.042$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.113$	$(\Delta/\sigma)_{max} = 0.001$
$S = 1.05$	$\Delta\rho_{max} = 0.26 \text{ e \AA}^{-3}$
6198 reflections	$\Delta\rho_{min} = -0.39 \text{ e \AA}^{-3}$
425 parameters	Extinction correction: <i>SHELXL97</i>
H atoms treated by a mixture of independent and constrained refinement	Extinction coefficient: 0.016 (3)
	Absolute structure: Flack (1983),
	2176 Friedel pairs
	Flack parameter: $-0.03 (6)$

Table 1

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

S1—O2	1.438 (3)	C1—C2	1.395 (4)
S1—O3	1.439 (2)	C5—C6	1.413 (4)
S1—O4	1.460 (3)	C10—C11	1.540 (4)
S1—C25	1.775 (3)	C14—C18	1.502 (5)
S2—O5	1.462 (2)	C18—C19	1.293 (6)
O1—C10	1.413 (4)	C20—C25	1.382 (4)
N2—C11	1.509 (4)	C21—C22	1.379 (6)
O2—S1—O3	114.45 (15)	O1—C10—C7	111.1 (2)
O2—S1—C25	106.39 (15)	C7—C10—C11	109.0 (2)
O7—S2—O6	116.4 (2)	C13—C12—C11	109.0 (2)
C9—N1—C1	123.2 (3)	C19—C18—C14	124.7 (5)
C15—N2—C11	107.6 (2)	C21—C22—C23	117.8 (3)
C2—C3—C4	120.5 (3)	C24—C25—C20	119.5 (3)

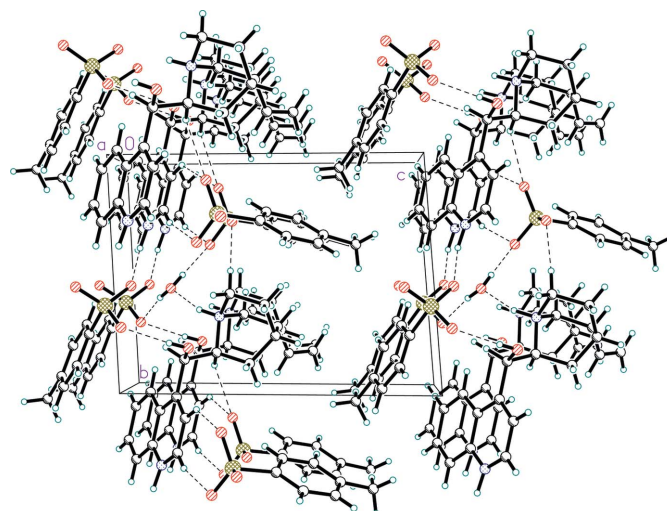


Figure 2

A packing diagram of (I), viewed along the a axis. Hydrogen bonds are indicated by dashed lines.

Table 2

Hydrogen-bond geometry (\AA , $^\circ$).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
N1—H1N···O6 ⁱ	0.98 (6)	1.86 (6)	2.749 (4)	150 (5)
N2—H2N···O8	0.99 (4)	1.75 (4)	2.718 (3)	164 (4)
O1—H1O···O5 ⁱⁱ	0.82	1.96	2.754 (3)	163
O8—H0A···O4	0.82 (1)	1.91 (1)	2.731 (4)	179 (5)
O8—H0B···O5 ⁱⁱⁱ	0.83 (1)	1.98 (1)	2.806 (3)	172 (4)
C8—H8···O3 ^{iv}	0.93	2.46	3.205 (4)	138
C9—H9···O4 ^{iv}	0.93	2.34	3.247 (4)	164
C11—H11···O3 ^v	0.98	2.51	3.320 (4)	140
C17—H17A···O1	0.97	2.39	3.026 (4)	123
C17—H17B···O2	0.97	2.50	3.441 (3)	164

Symmetry codes: (i) $x+1, y+1, z-1$; (ii) $x+1, y, z-1$; (iii) $x, y, z-1$; (iv) $x+1, y+1, z$; (v) $x, y+1, z$.

H atoms of the water molecule and those attached to N atoms were located in difference Fourier maps. Water H atoms were refined with O—H and H···H distances restrained to 0.82 (1) and 1.34 (1) \AA , respectively, while the H atoms of the NH group were refined freely. All other H atoms were placed in calculated positions and allowed to ride on their parent atoms at distances of 0.93 (aromatic and alkene), 0.96 (methyl), 0.97 (methylene), 0.98 (methine) and 0.82 \AA (hydroxyl), with $U_{iso}(H) = 1.2U_{eq}(C, O)$.

Data collection: *XSCANS* (Siemens, 1994); cell refinement: *XSCANS*; data reduction: *SHELXTL* (Bruker, 1998); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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