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## Structure Reports

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## 2-[(E)-2-(4-Methoxyphenyl)ethenyl]-1-methylquinolinium 4-bromobenzenesulfonate

In the title compound $\mathrm{C}_{19} \mathrm{H}_{18} \mathrm{NO}^{+} \cdot \mathrm{C}_{6} \mathrm{H}_{4} \mathrm{BrSO}_{3}{ }^{-}$, the cation is almost planar. The benzene ring of the anion makes dihedral angles of $84.23(12)$ and $84.59(15)^{\circ}$, respectively, with the quinolinium group and the benzene ring of the cation. The cations and anions are arranged in an alternating onedimensional chain along the $b$ axis, and these chains are interconnected through $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ and $\mathrm{C}-\mathrm{H} \cdots \mathrm{Br}$ interactions to form a three-dimensional network. $\mathrm{C}-\mathrm{H} \cdots \pi$ interactions are also observed in the crystal structure.

## Comment

Our ongoing interest in new nonlinear optical (NLO) materials (Rahman et al., 2003; Jindawong, Chantrapromma, Fun, Yu \& Karalai, 2005; Jindawong, Chantrapromma, Fun \& Karalai, 2005; Chantrapromma et al., 2005, 2006) led us to synthesize the title compound, (I), in order to explore its molecular behaviour and crystal structure. However, since second-order nonlinear effects require a non-centrosymmetric structure for the bulk material (Williams, 1984) and (I) crystallizes in a centrosymmetric space group, it cannot exhibit second-order NLO properties.


The asymmetric unit of (I) consists of a $\mathrm{C}_{19} \mathrm{H}_{18} \mathrm{NO}^{+}$cation and a $\mathrm{C}_{6} \mathrm{H}_{4} \mathrm{BrSO}_{3}{ }^{-}$anion (Fig. 1). Bond lengths and angles are in normal ranges (Allen et al., 1987) and similar to those in some closely related structures, such as 2-[(E)-2-(3-hydroxy-4-methoxyphenyl)ethenyl]-1-methylquinolinium 4-methylbenzenesulfonate, (II) (Jindawong, Chantrapromma, Fun \& Karalai, 2005) and 2-[(E)-2-(3-hydroxy-4-methoxyphenyl)-ethenyl]-1-methylquinolinium 4-chlorobenzenesulfonate, (III) (Chantrapromma et al., 2006). In the cation, the quinolinium group (C9-C17/N1) is planar, with a maximum deviation from the plane of 0.054 (3) $\AA$ for atom N1. Overall, the cation is almost planar, as indicated by the dihedral angle of $1.12(14)^{\circ}$ between the quinolinium group and the $\mathrm{C} 1-\mathrm{C} 6$ benzene ring. The H atoms attached to atoms C7 and C8 are mutually trans, so that the cation adopts an $E$ configuration with a $\mathrm{C} 6-\mathrm{C} 7-$

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## Key indicators

Single-crystal X-ray study
$T=100 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.004 \AA$
$R$ factor $=0.048$
$w R$ factor $=0.112$
Data-to-parameter ratio $=21.4$

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


Figure 1
The asymmetric unit of (I), showing $80 \%$ probability displacement ellipsoids and the atomic numbering scheme.


Figure 2
The crystal packing of (I), viewed down the $a$ axis. Weak $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ and $\mathrm{C}-\mathrm{H} \cdots \mathrm{Br}$ interactions are shown as dashed lines.

C8-C9 torsion angle of 177.4 (4) ${ }^{\circ}$. The methoxy group deviates slightly from the plane of the benzene ring of the cation $\left[\mathrm{C} 19-\mathrm{O} 1-\mathrm{C} 3-\mathrm{C} 4=-5.6(5)^{\circ}\right]$. In the asymmetric unit, the molecular planes of the cation and anion are almost perpendicular to each other, with a dihedral angle of $84.23(12)^{\circ}$ between the benzene ring of the anion and the quinolinium plane. This compares with 59.17 (9) ${ }^{\circ}$ in (II) and 48.97 (6) ${ }^{\circ}$ in (III). The angle between the benzene rings of the anion and cation is $84.59(15)^{\circ}\left[60.59(10)^{\circ}\right.$ in (II) and 51.63 (7) ${ }^{\circ}$ in (III)].

In the crystal structure, all the O atoms of the 4-bromobenzenesulfonate anion are involved in weak $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ interactions (Table 1 and Fig. 2). The cations and anions form alternating one-dimensional chains along the $b$ axis. These chains are interconnected by weak $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ and $\mathrm{C}-\mathrm{H} \cdots \mathrm{Br}$ interactions to form a three-dimensional molecular network.
$\mathrm{C}-\mathrm{H} \cdots \pi$ interactions are also observed and details are given in Table 1.

## Experimental

Silver(I) 4-bromobenzenesulfonate (compound $A$ ) was synthesized as reported previously (Jindawong, Chantrapromma, Fun, Yu \& Karalai, 2005). 2-(4'-Methoxystyryl)-1-methylquinolinium iodide (compound $B$ ) was synthesized by refluxing a solution of $1,2-$ dimethylquinolinium iodide $(1.00 \mathrm{~g}, \quad 3.51 \mathrm{mmol})$, 4 -methoxybenzaldehyde $(0.43 \mathrm{ml}, \quad 3.51 \mathrm{mmol})$ and piperidine $(0.35 \mathrm{ml}$, $3.54 \mathrm{mmol})$, in a $1: 1: 1$ molar ratio, in methanol $(20 \mathrm{ml})$ for 7 h . The resulting solid was filtered off, washed with $\mathrm{CHCl}_{3}$ and recrystallized from $\mathrm{CH}_{3} \mathrm{OH}$, to give $B$ as a green solid ( $0.95 \mathrm{~g}, 67 \%$, mp. 498-499 K). The title compound, (I), was synthesized by mixing solutions of $A$ $(0.09 \mathrm{~g}, 0.25 \mathrm{mmol})$ and $B(0.10 \mathrm{~g}, 0.25 \mathrm{mmol})$, each in hot MeOH ( 70 ml ), which immediately gave a yellow mass of silver iodide. After stirring the mixture for 30 min , the silver iodide was removed and the resulting green-yellow solution was evaporated to yield a greenyellow solid. Single orange blocks of (I) were obtained by recrystallization from $\mathrm{MeOH}-\mathrm{EtOH}(2: 1 \mathrm{v} / \mathrm{v})$ at ambient temperature over several days (m.p. 539-540 K).

## Crystal data

$\mathrm{C}_{19} \mathrm{H}_{18} \mathrm{NO}^{+} . \mathrm{C}_{6} \mathrm{H}_{4} \mathrm{BrO}_{3} \mathrm{~S}^{-}$
$M_{r}=512.41$
Monoclinic, $P 2_{\mathrm{o}_{1}} / c$
$a=7.1945$ (1) А
$b=15.1542(2) \AA$
$c=19.6438$ (3) $\AA$
$\beta=90.960(1)^{\circ}$
$V=2141.40(5) \AA^{3}$

## Data collection

Bruker SMART APEX2 CCD areadetector diffractometer
$\omega$ scans
Absorption correction: multi-scan (SADABS; Bruker, 2005)
$T_{\text {min }}=0.415, T_{\max }=0.676$

$$
\begin{aligned}
& Z=4 \\
& D_{x}=1.589 \mathrm{Mg} \mathrm{~m}^{-3} \\
& \text { Mo } K \alpha \text { radiation } \\
& \mu=2.05 \mathrm{~mm}^{-1} \\
& T=100.0(1) \mathrm{K} \\
& \text { Block, orange } \\
& 0.52 \times 0.37 \times 0.21 \mathrm{~mm}
\end{aligned}
$$

## Refinement

Refinement on $F^{2}$

$$
\begin{aligned}
& w=1 /\left[\sigma^{2}\left(F_{\mathrm{o}}{ }^{2}\right)+(0.0369 P)^{2}\right. \\
& +4.0824 P] \\
& \text { where } P=\left(F_{\mathrm{o}}{ }^{2}+2 F_{\mathrm{c}}{ }^{2}\right) / 3 \\
& (\Delta / \sigma)_{\max }=0.001 \text { 。 } \\
& \Delta \rho_{\text {max }}=1.38 \mathrm{e}_{\mathrm{m}} \AA^{-3} \\
& \Delta \rho_{\text {min }}=-0.57 \mathrm{e}^{-3}
\end{aligned}
$$

## Table 1

Hydrogen-bond geometry ( $\AA,{ }^{\circ}$ ).
$C g 1$ is the centroid of the C20-C25 ring.

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :---: | :---: | :---: | :---: | :---: |
| $\mathrm{C} 5-\mathrm{H} 5 A \cdots \mathrm{O} 2$ | 0.93 | 2.54 | 3.436 (4) | 162 |
| C11-H11A $\cdots$ O $4^{\text {i }}$ | 0.93 | 2.58 | 3.481 (4) | 162 |
| C13-H13A $\cdots \mathrm{O}^{\text {i }}$ | 0.93 | 2.52 | 3.382 (4) | 155 |
| $\mathrm{C} 14-\mathrm{H} 14 A \cdots \mathrm{O} 3^{\text {ii }}$ | 0.93 | 2.58 | 3.219 (4) | 127 |
| $\mathrm{C} 18-\mathrm{H} 18 \mathrm{C} \cdots \mathrm{O} 2{ }^{\text {iii }}$ | 0.96 | 2.32 | 3.252 (4) | 165 |
| $\mathrm{C} 22-\mathrm{H} 22 \mathrm{~A} \cdots \mathrm{O} 4$ | 0.93 | 2.54 | 2.919 (4) | 105 |
| $\mathrm{C} 25-\mathrm{H} 25 A \cdots \mathrm{O} 4^{\text {iv }}$ | 0.93 | 2.29 | 3.083 (4) | 142 |
| $\mathrm{C} 4-\mathrm{H} 4 A \cdots \mathrm{Cg} 1$ | 0.93 | 2.87 | 3.655 (4) | 142 |

All H atoms bound to C atoms were refined using a riding model, with $\mathrm{C}-\mathrm{H}=0.93 \AA$ and $U_{\text {iso }}(\mathrm{H})=1.2 U_{\text {eq }}(\mathrm{C})$ for aromatic H , and $\mathrm{C}-$ $\mathrm{H}=0.96 \AA$ and $U_{\text {iso }}(\mathrm{H})=1.5 U_{\text {eq }}(\mathrm{C})$ for $\mathrm{CH}_{3}$ atoms. The highest peak is located $1.15 \AA$ from H19C.

Data collection: APEX2 (Bruker, 2005); cell refinement: APEX2; data reduction: SAINT (Bruker, 2005); program(s) used to solve structure: SHELXTL (Sheldrick, 1998); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL and PLATON (Spek, 2003).

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