

Vadivelu Manivannan,^a
Nagarajan Vembu,^{b,*} Maruthai
Nallu,^b Kandasamy Sivakumar^c
and Frank R. Fronczek^d^aDepartment of Physics, Presidency College,
Chennai 600 005, India, ^bDepartment of
Chemistry, Bharathidasan University, Tiruchir-
appalli 620 024, India, ^cDepartment of Physics,
Anna University, Chennai 600 025, India, and
^dDepartment of Chemistry, Louisiana State
University, Baton Rouge, LA 70803-1804, USA* Correspondence address: Department of
Chemistry, Urumu Dhanalakshmi College,
Tiruchirappalli 620 019, Tamil Nadu, India.

Correspondence e-mail: vembu57@yahoo.com

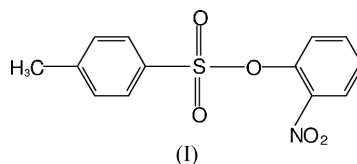
Key indicators

Single-crystal X-ray study
 $T = 150$ K
Mean $\sigma(\text{C}-\text{C}) = 0.006$ Å
 R factor = 0.061
 wR factor = 0.187
Data-to-parameter ratio = 15.3For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.2-Nitrophenyl 4-toluenesulfonate: supramolecular
aggregation through weak C—H···O, C—H··· π and
 π — π interactions

In the crystal structure of the title compound, $\text{C}_{13}\text{H}_{11}\text{NO}_5\text{S}$, the dihedral angle between the mean planes of the 4-tolyl and 2-nitrophenyl rings is $42.2(2)^\circ$. There are weak C—H···O hydrogen bonds, which generate rings of graph-set motifs $S(5)$, $R_1^2(4)$, $R_2^1(5)$, $R_2^2(6)$, $R_1^2(6)$, $R_2^2(7)$ and $R_2^2(10)$. The supramolecular aggregation is completed by the presence of intermolecular van der Waals short contacts, and C—H··· π and π — π interactions.

Comment

1-Anilino-8-naphthalenesulfonate, an aromatic sulfonate, is used in monitoring the merging of lipids in the binding of Rose bengal, a model organic anion, to sinusoidal and bile canalicular membrane fractions isolated from rat liver (Yachi *et al.*, 1989) and in many other fields (Spungin *et al.*, 1992; Tharakan *et al.*, 1992; Alford *et al.*, 1991; Jiang *et al.*, 1990; Narayanan & Krakow, 1983). The crystal structure of the title compound, (I), was determined because of the biological importance of its analogues. A search of Version 5.23 (July 2002 updates) of the Cambridge Structural Database (Allen, 2002) revealed 16 structures (with the following refcodes: KAWDAN, FIXCAQ, NEDXUP, NEDYAW, NEDYIE, NUNCHI, RASSOT, RELVUZ, SIMVUF, TCPTOS, TEBFOV, TMPDTS, TSMIPH, WOHCUR, ZZZBDA10 and MIWHIJ) that are closely related to the title compound in that they all contain the *p*-toluenesulfonyl group. The S—C, S—O and S=O bond lengths (Table 1) are comparable to those found in related structures (Vembu, Nallu, Garrison & Youngs, 2003*a,b,c,d,e,f*; Vembu, Nallu, Spencer & Howard, 2003*a,b,c,d,e,f,g*; Vembu, Nallu, Garrison, Hindi & Youngs, 2003; Vembu, Nallu, Durmus *et al.*, 2004*a,b,c*).



The molecular structure of (I) is shown in Fig. 1. The dihedral angle between the mean planes of the C1—C6 and C8—C13 rings is $42.2(2)^\circ$. This shows their non-coplanar orientation, which is similar to that reported for other aromatic sulfonates (Vembu, Nallu, Garrison & Youngs, 2003*b,c,d,e*; Vembu, Nallu, Spencer & Howard, 2003*a,b,c,d,f,g*; Vembu, Nallu, Durmus *et al.*, 2004*a,b,c*) and in contrast the near coplanar orientation observed in 2,4-dinitrophenyl 4-toluenesulfonate (Vembu, Nallu, Garrison & Youngs, 2003*a*), 4-methoxyphenyl 4-toluenesulfonate (Vembu, Nallu, Garrison, Hindi & Youngs, 2003) and 8-quinolyl 3-nitrobenzenesulfon-

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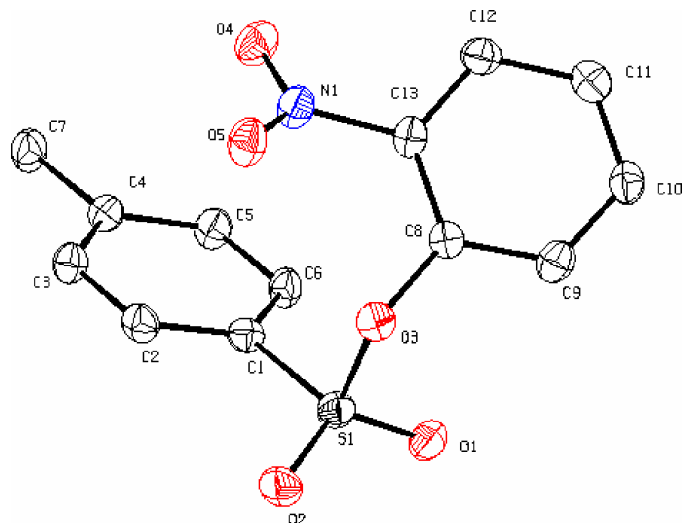


Figure 1

The molecular structure of the title molecule, showing 50% probability displacement ellipsoids. H atoms have been omitted.

ate (Vembu, Nallu, Spencer & Howard, 2003e). The aromatic carbon, C8, attached to an O atom is taken as fiducial and the molecule is viewed along the S—O bond (Fig. 2). The orientations of the sulfonyl atoms O1 and O2 and the tolyl atom C1 attached to the S atom have been deduced from the corresponding torsion angles (C8—O3—S1—O1, C8—O3—S1—O2 and C8—O3—S1—C1) and are depicted in Fig. 2. Since the C8—O3—S1—C1 torsion angle is 72.7 (3)°, which corresponds to a synclinal conformation, the two aromatic planes are, as expected, not coplanar [dihedral angle 42.2 (2)°].

The crystal structure of (I) is stabilized by weak C—H...O interactions (Table 2). The range for the H...O distances in (I) agrees with those found for weak C—H...O bonds (Desiraju & Steiner, 1999). In (I), each of the C2—H2...O2 and C6—H6...O1 interactions generates an *S*(5) graph-set motif (Bernstein *et al.*, 1995; Etter, 1990). The *S*(5) rings were found to be non-planar. The C2—H2...O4ⁱ and C2—H2...O5ⁱ interactions together constitute a pair of bifurcated donor bonds generating a chelate ring of graph set *R*₁²(4) (symmetry codes are as in Table 2). The C2—H2...O4ⁱ and C3—H3...O4ⁱ interactions together form a pair of bifurcated acceptor bonds generating an *R*₂¹(5) motif. The above *R*₁²(4) and *R*₂¹(5) motifs together form an *R*₂²(7) motif, which is formed by the interactions C2—H2...O5ⁱ and C3—H3...O4ⁱ. The C7—H7B...O3ⁱⁱ and C5—H5...O3ⁱⁱ interactions constitute a pair of bifurcated acceptor bonds generating an *R*₂¹(6) motif. The C7—H7B...O5ⁱⁱ and C7—H7B...O3ⁱⁱ interactions form a pair of bifurcated donor bonds generating an *R*₁²(6) motif. The above *R*₂¹(6) and *R*₁²(6) motifs together form an *R*₂²(10) motif, which is formed by the interactions, C5—H5...O3ⁱⁱ and C7—H7B...O5ⁱⁱ. The two symmetry-related 2-nitrophenyl rings (3 - *x*, 1 - *y*, 1 - *z*) are stacked over each other, with a centroid-centroid distance of 3.621 Å. There are several other C—H...O, C—H...π and van der Waals interactions, which contribute to the supramolecular aggregation (Fig. 3 and Table 2) of the title compound.

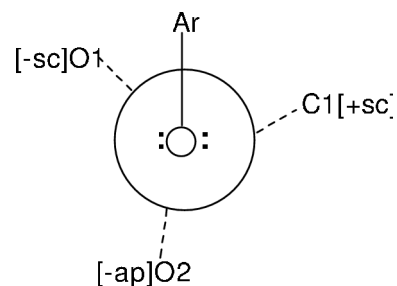


Figure 2

Newman projection of the title molecule along the S—O bond.

Experimental

4-Toluenesulfonyl chloride (4.7 mmol), dissolved in acetone, was added dropwise to 2-nitrophenol (4.3 mmol) in aqueous NaOH (2.5 ml, 10%) with constant shaking. The precipitated title compound (3.1 mmol, yield 72%) was filtered off and recrystallized from aqueous ethanol.

Crystal data

C₁₃H₁₁NO₅S
M_r = 293.29
 Monoclinic, *P*₂₁/*a*
a = 7.953 (1) Å
b = 20.468 (3) Å
c = 7.9450 (9) Å
 β = 99.516 (8)°
V = 1275.5 (3) Å³
Z = 4

D_x = 1.527 Mg m⁻³
 Mo *K*α radiation
 Cell parameters from 23 615 reflections
 θ = 2.5–27.4°
 μ = 0.27 mm⁻¹
T = 150 (2) K
 Irregular fragment, colorless
 0.35 × 0.25 × 0.10 mm

Data collection

Nonius KappaCCD diffractometer
 ω scans with κ offsets
 Absorption correction: multi-scan
 (*HKL SCALEPACK*;
 Otwinowski & Minor, 1997)
*T*_{min} = 0.902, *T*_{max} = 0.973
 3173 measured reflections

2839 independent reflections
 1446 reflections with *I* > 2σ(*I*)
*R*_{int} = 0.055
 θ _{max} = 27.4°
h = -10 → 10
k = -26 → 25
l = -10 → 10

Refinement

Refinement on *F*²
R [*F*² > 2σ(*F*²)] = 0.061
wR (*F*²) = 0.187
S = 0.98
 2839 reflections
 185 parameters

H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0948P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 (Δ/σ)_{max} < 0.001
 $\Delta\rho$ _{max} = 0.31 e Å⁻³
 $\Delta\rho$ _{min} = -0.41 e Å⁻³

Table 1

Selected geometric parameters (Å, °).

S1—O2	1.423 (3)	O3—C8	1.401 (4)
S1—O1	1.427 (3)	O4—N1	1.224 (4)
S1—O3	1.626 (3)	O5—N1	1.236 (4)
S1—C1	1.754 (3)	N1—C13	1.475 (5)
O2—S1—O1	121.17 (17)	O3—S1—C1	102.62 (16)
O2—S1—O3	102.07 (16)	C8—O3—S1	117.7 (2)
O1—S1—O3	107.75 (16)	O4—N1—O5	124.3 (3)
O2—S1—C1	111.42 (18)	O4—N1—C13	117.3 (3)
O1—S1—C1	109.82 (17)	O5—N1—C13	118.2 (3)
O2—S1—O3—C8	-171.8 (3)	C1—S1—O3—C8	72.7 (3)
O1—S1—O3—C8	-43.1 (3)		

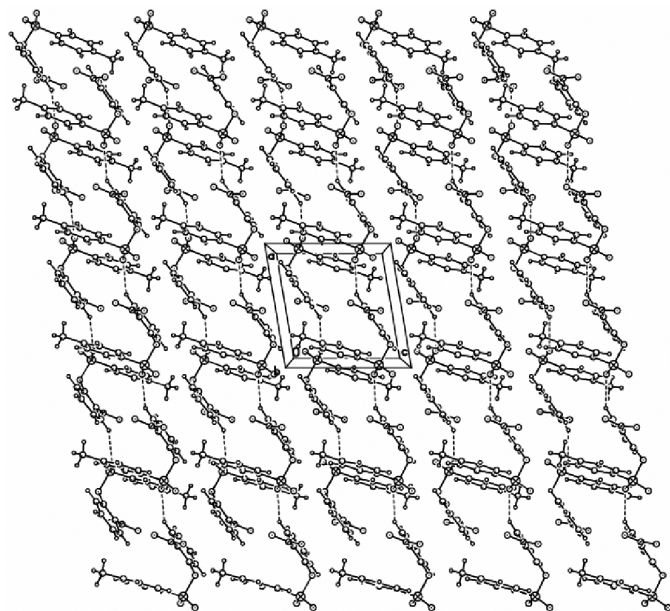


Figure 3
Packing of the molecules, viewed down the *b* axis. Dashed lines indicate hydrogen bonds.

Table 2
Hydrogen-bond geometry (Å, °).

<i>D</i> — <i>H</i> ··· <i>A</i>	<i>D</i> — <i>H</i>	<i>H</i> ··· <i>A</i>	<i>D</i> ··· <i>A</i>	<i>D</i> — <i>H</i> ··· <i>A</i>
C2—H2···O2	0.93	2.68	3.009 (5)	102
C6—H6···O1	0.93	2.57	2.929 (4)	103
C2—H2···O4 ⁱ	0.93	2.95	3.430 (5)	114
C2—H2···O5 ⁱ	0.93	2.84	3.688 (5)	153
C3—H3···O4 ⁱ	0.93	2.70	3.313 (5)	124
C7—H7B···O5 ⁱⁱ	0.96	2.72	3.570 (5)	149
C5—H5···O3 ⁱⁱⁱ	0.93	2.64	3.504 (4)	156
C7—H7B···O3 ⁱⁱⁱ	0.96	2.74	3.590 (5)	148
C6—H6···O1 ⁱⁱⁱ	0.93	2.69	3.463 (5)	141
C7—H7C···O5 ^{iv}	0.96	2.87	3.793 (5)	163
C9—H9···O1 ^v	0.93	2.89	3.566 (4)	130
C12—H12···O1 ^{vi}	0.93	2.57	3.392 (5)	148
C7—H7A···Cg1 ^{vii}	0.96	3.08	3.99	158

Symmetry codes: (i) $x - \frac{1}{2}, -y + \frac{1}{2}, z$; (ii) $x, y, z - 1$; (iii) $-x + 2, -y + 1, -z$; (iv) $x - \frac{1}{2}, -y + \frac{1}{2}, z - 1$; (v) $-x + 2, -y + 1, -z + 1$; (vi) $x + 1, y, z$; (vii) $x - \frac{1}{2}, -y - \frac{1}{2}, z$. Cg1 is the centroid of the C1–C6 ring.

All H atoms were included in calculated position with aromatic C—H distances of 0.93 Å and methyl C—H distances of 0.96 Å and refined with riding model. A group displacement parameter was refined for each type of H atom.

Data collection: *COLLECT* (Nonius, 2000); cell refinement: *DENZO* and *SCALEPAK* (Otwinowski & Minor, 1997); data reduction: *DENZO* and *SCALEPAK*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine

structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXL97*.

References

- Alford, R. L., Honda, S., Lawrence, C. B. & Belmont, J. W. (1991). *Virology*, **183**, 611–619.
- Allen, F. H. (2002). *Acta Cryst.* **B58**, 380–388.
- Bernstein, J., Davis, R. E., Shimon, L. & Chang, N. (1995). *Angew. Chem. Int. Ed. Engl.* **34**, 1555–1573.
- Desiraju, G. R. & Steiner, T. (1999). *The Weak Hydrogen Bond in Structural Chemistry and Biology*, New York: Oxford University Press.
- Etter, M. C. (1990). *Acc. Chem. Res.* **23**, 120–126.
- Jiang, F. N., Jiang, S., Liu, D., Richter, A. & Levy, J. G. (1990). *J. Immunol. Methods*, **134**, 139–149.
- Narayanan, C. S. & Krakow, J. S. (1983). *Nucleic Acids Res.* **11**, 2701–2716.
- Nonius (2000). *COLLECT*. Nonius BV, Delft, The Netherlands.
- Otwinowski, Z. & Minor, W. (1997). *Methods in Enzymology*, Vol. 276, *Macromolecular Crystallography*, Part A, edited by C. W. Carter Jr & R. M. Sweet, pp. 307–326. New York: Academic Press.
- Sheldrick, G. M. (1997). *SHELXL97*. University of Göttingen, Germany.
- Spungin, B., Levinshal, T., Rubenstein, S. & Breitbart, H. (1992). *FEBS Lett.* **311**, 155–160.
- Tharakan, J., Highsmith, F., Clark, D. & Drohsn, W. (1992). *J. Chromatogr.* **595**, 103–111.
- Vembu, N., Nallu, M., Durmus, S., Panzner, M., Garrison, J. & Youngs, W. J. (2004a). *Acta Cryst.* **E60**, o1–o3.
- Vembu, N., Nallu, M., Durmus, S., Panzner, M., Garrison, J. & Youngs, W. J. (2004b). *Acta Cryst.* **C60**, o65–o68.
- Vembu, N., Nallu, M., Durmus, S., Panzner, M., Garrison, J. & Youngs, W. J. (2004c). *Acta Cryst.* **C60**, o248–o251.
- Vembu, N., Nallu, M., Garrison, J., Hindi, K. & Youngs, W. J. (2003). *Acta Cryst.* **E59**, o830–o832.
- Vembu, N., Nallu, M., Garrison, J. & Youngs, W. J. (2003a). *Acta Cryst.* **E59**, o378–o380.
- Vembu, N., Nallu, M., Garrison, J. & Youngs, W. J. (2003b). *Acta Cryst.* **E59**, o503–o505.
- Vembu, N., Nallu, M., Garrison, J. & Youngs, W. J. (2003c). *Acta Cryst.* **E59**, o776–o779.
- Vembu, N., Nallu, M., Garrison, J. & Youngs, W. J. (2003d). *Acta Cryst.* **E59**, o936–o938.
- Vembu, N., Nallu, M., Garrison, J. & Youngs, W. J. (2003e). *Acta Cryst.* **E59**, o939–o941.
- Vembu, N., Nallu, M., Garrison, J. & Youngs, W. J. (2003f). *Acta Cryst.* **E59**, o1019–o1021.
- Vembu, N., Nallu, M., Spencer, E. C. & Howard, J. A. K. (2003a). *Acta Cryst.* **E59**, o1009–o1011.
- Vembu, N., Nallu, M., Spencer, E. C. & Howard, J. A. K. (2003b). *Acta Cryst.* **E59**, o1033–o1035.
- Vembu, N., Nallu, M., Spencer, E. C. & Howard, J. A. K. (2003c). *Acta Cryst.* **E59**, o1213–o1215.
- Vembu, N., Nallu, M., Spencer, E. C. & Howard, J. A. K. (2003d). *Acta Cryst.* **E59**, o1216–o1219.
- Vembu, N., Nallu, M., Spencer, E. C. & Howard, J. A. K. (2003e). *Acta Cryst.* **E59**, o1379–o1382.
- Vembu, N., Nallu, M., Spencer, E. C. & Howard, J. A. K. (2003f). *Acta Cryst.* **E59**, o1387–o1389.
- Vembu, N., Nallu, M., Spencer, E. C. & Howard, J. A. K. (2003g). *Acta Cryst.* **E59**, o1390–o1392.
- Yachi, K., Sugiyama, Y., Sawada, Y., Iga, T., Ikeda, Y., Toda, G. & Hanano, M. (1989). *Biochim. Biophys. Acta*, **978**, 1–7.