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Key indicators

Single-crystal X-ray study T = 273 K Mean σ (C–C) = 0.003 Å R factor = 0.047 wR factor = 0.146 Data-to-parameter ratio = 18.7

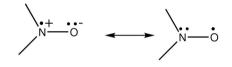
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

4-(Benzenesulfonyloxy)-2,2,6,6-tetramethylpiperidin-4-yl oxide

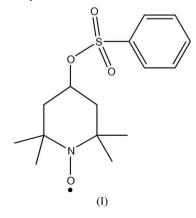
The title compound, $C_{15}H_{22}NO_4S$, is a nitroxyl radical, which is stable mainly because the N atom is surrounded by tertiary C atoms, preventing formation of the nitrone. More significantly, intermolecular C-H···O hydrogen bonding contributes to the stability of the structure with weak intramolecular C-H··· π (arene) interactions. The N-O distance is 1.2816 (18) Å.

Comment

Nitroxyl radicals are compounds containing an NO group that has one unpaired electron. The structure of this fragment can be conceived as a superposition of two resonance structures:



The contributions of the first and second structures to the ground state may be different, depending on the effects of conjugation and polarity of the medium. Piloty & Schwerin (1901) prepared the first organic nitroxyl. The structure of this nitroxyl free radical was later determined (Holden *et al.*, 1951). A widely used general procedure for the preparation of aromatic nitroxyls was developed (Wieland & Roth, 1920; Wieland & Kogl, 1922). A considerable amount of research, both theoretical and experimental, has been devoted to the electronic structure of nitroxyls (Stone *et al.*, 1965; Karimov *et al.*, 1969). It has been suggested that nitroxyl radicals are stable when the N atom is surrounded by tertiary C atoms, as formation of nitrones would not be possible in this case (Johnson *et al.*, 1956), the absence of α -H atoms ensuring the stability of nitroxyls.



The title compound, (I) (Fig. 1), is a stable radical whose structure cannot be identified through ¹H NMR. Selected

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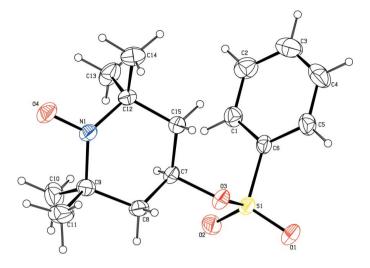


Figure 1

View of the molecule of (I), showing the atom-labelling scheme, with displacement ellipsoids drawn at the 50% probability level. H atoms are represented by circles of arbitrary size.

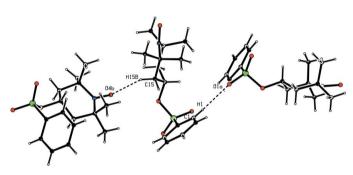


Figure 2

Intermolecular hydrogen bonding in the crystal structure of (I). Hydrogen-bonding interactions are indicated by dashed lines.

bond distances and angles describing the molecular conformation are listed in Table 1. The N1/C12/C15/C7–C9 ring exists as a chair conformation. The molecules are connected by intermolecular hydrogen bonding (Fig. 2 and Table 2). In addition, methyl atom C15 is involved in weak C–H··· π interactions (C15–H15A···Cg2 = 3.1452 Å).

Experimental

The title compound was synthesized according to the literature procedure (Rozantsev & Sholle, 1971). Crystals suitable for data collection were obtained by slow evaporation from an acetone–hexane solution at 293 K.

Crystal data

$C_{15}H_{22}NO_4S$
$M_r = 312.40$
Orthorhombic, Pbca
a = 11.4827 (9) Å
b = 15.6864 (13) Å
c = 18.4671 (14) Å
$V = 3326.3 (5) \text{ Å}^3$
Z = 8
$D_x = 1.248 \text{ Mg m}^{-3}$

Mo K α radiation Cell parameters from 3922 reflections $\theta = 2.5-22.4^{\circ}$ $\mu = 0.21 \text{ mm}^{-1}$ T = 273 (2) K Block, yellow $0.30 \times 0.30 \times 0.20 \text{ mm}$

Data collection

Bruker SMART CCD area-detector	2525 reflections with $I > 2\sigma(I)$
diffractometer	$R_{\rm int} = 0.034$
φ and ω scans	$\theta_{\rm max} = 27.0^{\circ}$
Absorption correction: none	$h = -12 \rightarrow 14$
18434 measured reflections	$k = -20 \rightarrow 19$
3629 independent reflections	$l = -23 \rightarrow 19$
Refinement	
Refinement on F^2	$w = 1/[\sigma^2(F_0^2) + (0.0745P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.047$	+ 0.3954P]
$wR(F^2) = 0.146$	where $P = (F_0^2 + 2F_c^2)/3$

 $(\Delta/\sigma)_{\text{max}} = 0.001$ $\Delta\rho_{\text{max}} = 0.29 \text{ e} \text{ Å}^{-3}$

 $\Delta \rho_{\rm min} = -0.28 \text{ e } \text{\AA}^{-3}$

Table 1

H-atom parameters constrained

S = 1.06

3629 reflections

194 parameters

Selected geometric parameters (Å, °).

C1-C2	1.366 (3)	C12-C15	1.526 (3)
C1-C6	1.374 (3)	C12-C14	1.527 (3)
C2-C3	1.372 (4)	C12-C13	1.530 (3)
C6-S1	1.755 (2)	N1-O4	1.2816 (18)
C7-O3	1.474 (2)	O1-S1	1.4224 (15)
C7-C15	1.499 (3)	O2-S1	1.4208 (15)
C12-N1	1.489 (3)	O3-S1	1.5649 (15)
O3-C7-C15	108.97 (14)	O4-N1-C9	115.77 (15)
N1-C9-C8	110.11 (15)	C12-N1-C9	125.03 (14)
N1-C12-C15	109.92 (14)	O3-S1-C6	104.09 (8)
O4-N1-C12	116.06 (15)		

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

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
C1-H1···O1 ⁱ	0.93	2.58	3.337 (3)	139
$C15-H15B\cdots O4^{ii}$	0.97	2.46	3.351 (2)	153
$C1 - H1 \cdots O2$	0.93	2.60	2.948 (3)	103
$C7 - H7 \cdot \cdot \cdot O2$	0.98	2.54	2.949 (2)	105

Symmetry codes: (i) $x - \frac{1}{2}, -y + \frac{1}{2}, -z + 1$; (ii) $x + \frac{1}{2}, y, -z + \frac{3}{2}$.

All H atoms were initially located in a difference Fourier map. The methyl H atoms were then constrained to an ideal geometry, with C–H distances of 0.96 Å and $U_{\rm iso}(\rm H) = 1.5 U_{eq}(\rm C)$. All other H atoms were placed in geometrically idealized positions, with C–H distances in the range 0.93–0.98 Å, and constrained to ride on their parent atoms with $U_{\rm iso}(\rm H) = 1.2 U_{eq}(\rm C)$.

Data collection: *SMART* (Bruker, 1997); cell refinement: *SAINT* (Bruker, 1999); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 2001); software used to prepare material for publication: *SHELXTL*.

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