

N-Cyclohexyl-N-ethyl-4-methylbenzene-sulfonamide

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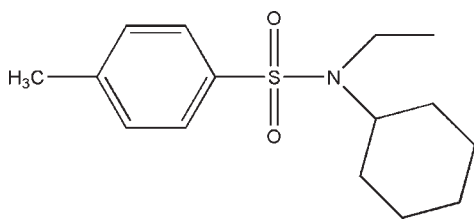
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Key indicators: single-crystal X-ray study; $T = 296$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.045; wR factor = 0.122; data-to-parameter ratio = 21.3.

The title compound, $\text{C}_{15}\text{H}_{23}\text{NO}_2\text{S}$, contains cyclohexyl and ethyl substituents on the sulfonamide N atom and the cyclohexyl ring adopts a classic chair conformation. The dihedral angle between the benzene ring plane and the mean plane through the six atoms of the cyclohexyl ring is $59.92(6)^\circ$. In the crystal structure, $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds link molecules into sheets extending in the bc plane.

Related literature

For ring conformations, see: Cremer & Pople (1975). For related structures, see: Arshad *et al.* (2008, 2009); Khan *et al.* (2009); Gowda *et al.* (2007a,b,c).



Experimental

Crystal data

$\text{C}_{15}\text{H}_{23}\text{NO}_2\text{S}$

$M_r = 281.40$

Monoclinic, $P2_1/c$

$a = 12.2269(5)$ Å

$b = 7.5818(3)$ Å

$c = 16.3045(6)$ Å

$\beta = 92.495(2)^\circ$

$V = 1510.03(10)$ Å³

$Z = 4$

Mo $K\alpha$ radiation

$\mu = 0.21$ mm⁻¹

$T = 296$ K

$0.43 \times 0.32 \times 0.15$ mm

Data collection

Bruker APEXII CCD area-detector diffractometer

Absorption correction: multi-scan (SADABS; Bruker, 2007)

$T_{\min} = 0.914$, $T_{\max} = 0.969$

16676 measured reflections

3714 independent reflections

2251 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.040$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.045$

$wR(F^2) = 0.122$

$S = 0.99$

3713 reflections

174 parameters

H-atom parameters constrained

$\Delta\rho_{\text{max}} = 0.23$ e Å⁻³

$\Delta\rho_{\text{min}} = -0.25$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{C10}-\text{H10B}\cdots\text{O2}^{\text{i}}$	0.97	2.66	3.530 (3)	150
$\text{C13}-\text{H13A}\cdots\text{O1}^{\text{ii}}$	0.96	2.60	3.512 (3)	159

Symmetry codes: (i) $x, y - 1, z$; (ii) $x, -y + \frac{5}{2}, z + \frac{1}{2}$.

Data collection: APEX2 (Bruker, 2007); cell refinement: SAINT (Bruker, 2007); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: ORTEP-3 (Farrugia, 1997) and Mercury (Macrae *et al.*, 2006); software used to prepare material for publication: SHELXL97, enCIFer (Allen *et al.*, 2004), PLATON (Spek, 2009) and publCIF (Westrip, 2009).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BT5132).

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supplementary materials

Acta Cryst. (2010). E66, o102 [doi:10.1107/S1600536809052593]

N-Cyclohexyl-*N*-ethyl-4-methylbenzenesulfonamide

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Comment

Our group has been involved in the synthesis and crystallographic studies of sulfonamide derivatives (Arshad *et al.*, 2009).

The title compound is a benzenesulfonamide with cyclohexyl and ethyl substituents on the sulfonamide N1 atom, Fig. 1. Bond distances in the molecule are comparable to those in similar structures (Arshad *et al.*, 2008; Khan *et al.*, 2009; Gowda *et al.*, 2007*a,b,c*). The cyclohexyl (C7···C12) ring adopts a classic chair conformation with puckering amplitude $Q = 0.567(2) \text{ \AA}$, $\theta = 178.8(2)^\circ$, $\varphi = 214(11)^\circ$ (Cremer & Pople, 1975) and the plane through this ring is inclined at $59.92(6)^\circ$ to that of the C1···C6 benzene ring. In the crystal structure C—H···O hydrogen bonds involving the C13—H13 bond of the methyl group and the C10—H10B bond of the cyclohexyl ring link each molecule to the O1 and O2 atoms of individual sulphonamide units forming an extended two dimensional network in the *bc* plane (Table 1, Fig. 2).

Experimental

A mixture of *N*-cyclohexyl-4-methyl benzene sulfonamide (1.089 g, 4.3 mmol), sodium hydride (0.21 g, 8.6 mmol) and *N,N*-dimethylformamide (10 ml) was stirred at room temperature for half an hour followed by addition of ethyl iodide (1.32 g, 8.6 mmol). Stirring was continued further for a period of three hours and the contents were poured over crushed ice. Precipitated product was isolated, washed and crystallized from a methanol solution.

Refinement

All H-atoms were positioned geometrically and refined using a riding model with $d(\text{C—H}) = 0.93 \text{ \AA}$, $U_{\text{iso}} = 1.2U_{\text{eq}}(\text{C})$ for aromatic 0.98 \AA , $U_{\text{iso}} = 1.2U_{\text{eq}}(\text{C})$ for CH, 0.97 \AA , $U_{\text{iso}} = 1.2U_{\text{eq}}(\text{C})$ for CH₂, 0.96 \AA , $U_{\text{iso}} = 1.5U_{\text{eq}}(\text{C})$ for CH₃ atoms.

The 1 0 0 reflection was identified as being obscured by the beamstop and was omitted.

Figures

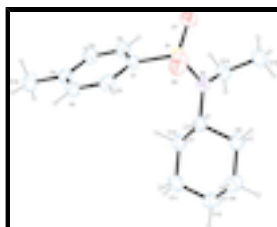


Fig. 1. The structure of (I) with displacement ellipsoids for the non-hydrogen atoms drawn at the 50% probability level.

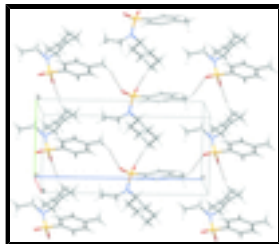


Fig. 2. Crystal packing of (I) viewed down the *a* axis with hydrogen bonds drawn as dashed lines.

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Monoclinic, $P2_1/c$

Hall symbol: -P 2ybc

$a = 12.2269$ (5) Å

$b = 7.5818$ (3) Å

$c = 16.3045$ (6) Å

$\beta = 92.495$ (2)°

$V = 1510.03$ (10) Å³

$Z = 4$

$F(000) = 608$

$D_x = 1.238$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 3495 reflections

$\theta = 2.5$ – 25.6 °

$\mu = 0.21$ mm⁻¹

$T = 296$ K

Needle, white

$0.43 \times 0.32 \times 0.15$ mm

Data collection

Bruker APEXII CCD area-detector diffractometer

Radiation source: fine-focus sealed tube graphite

φ and ω scans

Absorption correction: multi-scan (*SADABS*; Bruker, 2007)

$T_{\min} = 0.914$, $T_{\max} = 0.969$

16676 measured reflections

3714 independent reflections

2251 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.040$

$\theta_{\max} = 28.3$ °, $\theta_{\min} = 1.7$ °

$h = -16 \rightarrow 15$

$k = -10 \rightarrow 9$

$l = -19 \rightarrow 21$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.045$

$wR(F^2) = 0.122$

$S = 0.99$

3713 reflections

174 parameters

Primary atom site location: structure-invariant direct methods

Secondary atom site location: difference Fourier map

Hydrogen site location: inferred from neighbouring sites

H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.048P)^2 + 0.3506P]$

where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} = 0.001$

$\Delta\rho_{\max} = 0.23$ e Å⁻³

0 restraints

$$\Delta\rho_{\min} = -0.25 \text{ e } \text{\AA}^{-3}$$

Special details

Geometry. All s.u.'s (except the s.u. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell s.u.'s are taken into account individually in the estimation of s.u.'s in distances, angles and torsion angles; correlations between s.u.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell s.u.'s is used for estimating s.u.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > 2\sigma(F^2)$ is used only for calculating R -factors(gt) *etc.* and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
S1	0.22960 (5)	1.10442 (7)	0.56680 (3)	0.05930 (19)
O1	0.14930 (14)	1.1507 (2)	0.50392 (9)	0.0866 (5)
O2	0.32184 (13)	1.21640 (19)	0.58303 (10)	0.0784 (5)
N1	0.27372 (13)	0.9101 (2)	0.54440 (9)	0.0549 (4)
C1	0.16215 (14)	1.0870 (2)	0.66002 (11)	0.0451 (4)
C2	0.21516 (14)	1.1373 (2)	0.73271 (11)	0.0511 (5)
H2	0.2856	1.1835	0.7326	0.061*
C3	0.16309 (15)	1.1186 (3)	0.80559 (12)	0.0528 (5)
H3	0.1994	1.1519	0.8544	0.063*
C4	0.05847 (15)	1.0517 (2)	0.80760 (11)	0.0491 (5)
C5	0.00638 (16)	1.0042 (3)	0.73394 (13)	0.0571 (5)
H5	-0.0645	0.9598	0.7341	0.069*
C6	0.05660 (16)	1.0209 (3)	0.66051 (12)	0.0555 (5)
H6	0.0201	0.9881	0.6117	0.067*
C7	0.36213 (15)	0.8297 (2)	0.59696 (11)	0.0494 (5)
H7	0.3938	0.9237	0.6317	0.059*
C8	0.45338 (16)	0.7582 (3)	0.54568 (12)	0.0576 (5)
H8A	0.4245	0.6660	0.5096	0.069*
H8B	0.4808	0.8521	0.5118	0.069*
C9	0.54647 (16)	0.6847 (3)	0.59987 (14)	0.0671 (6)
H9A	0.6019	0.6358	0.5657	0.081*
H9B	0.5797	0.7793	0.6323	0.081*
C10	0.50692 (17)	0.5438 (3)	0.65621 (13)	0.0642 (6)
H10A	0.5673	0.5044	0.6922	0.077*
H10B	0.4811	0.4435	0.6239	0.077*
C11	0.4154 (2)	0.6108 (3)	0.70741 (13)	0.0697 (6)
H11A	0.4437	0.7014	0.7446	0.084*
H11B	0.3883	0.5148	0.7402	0.084*
C12	0.32174 (16)	0.6864 (3)	0.65394 (12)	0.0606 (5)
H12A	0.2670	0.7355	0.6887	0.073*
H12B	0.2876	0.5926	0.6214	0.073*

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C13	0.00216 (18)	1.0322 (3)	0.88731 (13)	0.0723 (6)
H13A	0.0434	1.0933	0.9300	0.108*
H13B	-0.0027	0.9095	0.9011	0.108*
H13C	-0.0701	1.0815	0.8817	0.108*
C14	0.21331 (17)	0.8024 (3)	0.48246 (12)	0.0608 (5)
H14A	0.1357	0.8263	0.4855	0.073*
H14B	0.2249	0.6787	0.4953	0.073*
C15	0.2469 (2)	0.8363 (3)	0.39659 (13)	0.0820 (7)
H15A	0.2338	0.9578	0.3828	0.123*
H15B	0.2050	0.7624	0.3592	0.123*
H15C	0.3233	0.8104	0.3927	0.123*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0778 (4)	0.0485 (3)	0.0529 (3)	0.0114 (3)	0.0175 (2)	0.0068 (2)
O1	0.1182 (13)	0.0855 (12)	0.0565 (9)	0.0423 (10)	0.0071 (9)	0.0196 (8)
O2	0.0959 (11)	0.0491 (9)	0.0935 (11)	-0.0132 (8)	0.0419 (9)	-0.0031 (8)
N1	0.0682 (10)	0.0513 (10)	0.0455 (9)	0.0081 (8)	0.0054 (7)	-0.0054 (8)
C1	0.0487 (10)	0.0398 (10)	0.0470 (10)	0.0075 (8)	0.0052 (8)	0.0001 (8)
C2	0.0410 (10)	0.0532 (12)	0.0591 (12)	-0.0008 (8)	0.0027 (9)	-0.0036 (9)
C3	0.0498 (11)	0.0590 (12)	0.0493 (11)	0.0003 (9)	-0.0015 (8)	-0.0077 (9)
C4	0.0537 (11)	0.0404 (10)	0.0537 (11)	0.0041 (8)	0.0078 (9)	-0.0022 (9)
C5	0.0451 (11)	0.0574 (13)	0.0692 (14)	-0.0064 (9)	0.0069 (9)	-0.0074 (10)
C6	0.0544 (12)	0.0569 (12)	0.0545 (12)	-0.0010 (10)	-0.0049 (9)	-0.0104 (10)
C7	0.0611 (11)	0.0441 (10)	0.0433 (10)	0.0015 (9)	0.0046 (8)	-0.0062 (8)
C8	0.0590 (12)	0.0559 (12)	0.0593 (12)	-0.0029 (10)	0.0164 (9)	0.0029 (10)
C9	0.0551 (12)	0.0620 (14)	0.0845 (16)	-0.0010 (11)	0.0058 (11)	-0.0003 (12)
C10	0.0640 (13)	0.0517 (12)	0.0760 (15)	0.0047 (10)	-0.0062 (11)	-0.0023 (11)
C11	0.0958 (16)	0.0593 (14)	0.0543 (12)	0.0046 (12)	0.0053 (11)	0.0072 (11)
C12	0.0674 (13)	0.0616 (13)	0.0543 (12)	0.0078 (11)	0.0207 (10)	0.0053 (10)
C13	0.0765 (15)	0.0772 (16)	0.0649 (14)	-0.0025 (12)	0.0229 (11)	-0.0008 (12)
C14	0.0638 (12)	0.0607 (13)	0.0576 (12)	-0.0026 (10)	-0.0001 (10)	-0.0015 (10)
C15	0.0992 (18)	0.0938 (19)	0.0525 (13)	-0.0011 (15)	-0.0013 (12)	-0.0099 (13)

Geometric parameters (\AA , $^\circ$)

S1—O2	1.4272 (16)	C8—H8B	0.9700
S1—O1	1.4320 (16)	C9—C10	1.502 (3)
S1—N1	1.6160 (17)	C9—H9A	0.9700
S1—C1	1.7654 (18)	C9—H9B	0.9700
N1—C14	1.472 (2)	C10—C11	1.513 (3)
N1—C7	1.481 (2)	C10—H10A	0.9700
C1—C2	1.380 (2)	C10—H10B	0.9700
C1—C6	1.385 (3)	C11—C12	1.521 (3)
C2—C3	1.380 (3)	C11—H11A	0.9700
C2—H2	0.9300	C11—H11B	0.9700
C3—C4	1.378 (3)	C12—H12A	0.9700
C3—H3	0.9300	C12—H12B	0.9700

C4—C5	1.383 (3)	C13—H13A	0.9600
C4—C13	1.504 (3)	C13—H13B	0.9600
C5—C6	1.375 (3)	C13—H13C	0.9600
C5—H5	0.9300	C14—C15	1.498 (3)
C6—H6	0.9300	C14—H14A	0.9700
C7—C8	1.523 (3)	C14—H14B	0.9700
C7—C12	1.525 (3)	C15—H15A	0.9600
C7—H7	0.9800	C15—H15B	0.9600
C8—C9	1.517 (3)	C15—H15C	0.9600
C8—H8A	0.9700		
O2—S1—O1	119.90 (11)	C8—C9—H9A	109.4
O2—S1—N1	108.38 (9)	C10—C9—H9B	109.4
O1—S1—N1	106.68 (9)	C8—C9—H9B	109.4
O2—S1—C1	106.25 (9)	H9A—C9—H9B	108.0
O1—S1—C1	107.65 (9)	C9—C10—C11	111.34 (18)
N1—S1—C1	107.42 (8)	C9—C10—H10A	109.4
C14—N1—C7	120.00 (16)	C11—C10—H10A	109.4
C14—N1—S1	119.89 (13)	C9—C10—H10B	109.4
C7—N1—S1	119.14 (12)	C11—C10—H10B	109.4
C2—C1—C6	119.88 (17)	H10A—C10—H10B	108.0
C2—C1—S1	119.95 (14)	C10—C11—C12	111.52 (17)
C6—C1—S1	120.16 (15)	C10—C11—H11A	109.3
C3—C2—C1	119.63 (17)	C12—C11—H11A	109.3
C3—C2—H2	120.2	C10—C11—H11B	109.3
C1—C2—H2	120.2	C12—C11—H11B	109.3
C2—C3—C4	121.47 (17)	H11A—C11—H11B	108.0
C2—C3—H3	119.3	C11—C12—C7	111.20 (17)
C4—C3—H3	119.3	C11—C12—H12A	109.4
C3—C4—C5	117.93 (17)	C7—C12—H12A	109.4
C3—C4—C13	121.15 (18)	C11—C12—H12B	109.4
C5—C4—C13	120.93 (18)	C7—C12—H12B	109.4
C6—C5—C4	121.74 (18)	H12A—C12—H12B	108.0
C6—C5—H5	119.1	C4—C13—H13A	109.5
C4—C5—H5	119.1	C4—C13—H13B	109.5
C5—C6—C1	119.34 (18)	H13A—C13—H13B	109.5
C5—C6—H6	120.3	C4—C13—H13C	109.5
C1—C6—H6	120.3	H13A—C13—H13C	109.5
N1—C7—C8	111.26 (15)	H13B—C13—H13C	109.5
N1—C7—C12	113.46 (16)	N1—C14—C15	113.33 (17)
C8—C7—C12	110.21 (16)	N1—C14—H14A	108.9
N1—C7—H7	107.2	C15—C14—H14A	108.9
C8—C7—H7	107.2	N1—C14—H14B	108.9
C12—C7—H7	107.2	C15—C14—H14B	108.9
C9—C8—C7	111.12 (16)	H14A—C14—H14B	107.7
C9—C8—H8A	109.4	C14—C15—H15A	109.5
C7—C8—H8A	109.4	C14—C15—H15B	109.5
C9—C8—H8B	109.4	H15A—C15—H15B	109.5
C7—C8—H8B	109.4	C14—C15—H15C	109.5
H8A—C8—H8B	108.0	H15A—C15—H15C	109.5

supplementary materials

C10—C9—C8	111.37 (17)	H15B—C15—H15C	109.5
C10—C9—H9A	109.4		
O2—S1—N1—C14	-144.40 (15)	C13—C4—C5—C6	-179.92 (19)
O1—S1—N1—C14	-14.01 (17)	C4—C5—C6—C1	0.0 (3)
C1—S1—N1—C14	101.19 (15)	C2—C1—C6—C5	-0.8 (3)
O2—S1—N1—C7	46.86 (16)	S1—C1—C6—C5	178.40 (15)
O1—S1—N1—C7	177.24 (14)	C14—N1—C7—C8	60.5 (2)
C1—S1—N1—C7	-67.56 (15)	S1—N1—C7—C8	-130.77 (15)
O2—S1—C1—C2	-15.54 (18)	C14—N1—C7—C12	-64.5 (2)
O1—S1—C1—C2	-145.16 (16)	S1—N1—C7—C12	104.27 (17)
N1—S1—C1—C2	100.29 (16)	N1—C7—C8—C9	177.02 (16)
O2—S1—C1—C6	165.26 (15)	C12—C7—C8—C9	-56.2 (2)
O1—S1—C1—C6	35.64 (18)	C7—C8—C9—C10	56.7 (2)
N1—S1—C1—C6	-78.91 (16)	C8—C9—C10—C11	-55.7 (2)
C6—C1—C2—C3	1.0 (3)	C9—C10—C11—C12	55.0 (2)
S1—C1—C2—C3	-178.17 (14)	C10—C11—C12—C7	-55.1 (2)
C1—C2—C3—C4	-0.5 (3)	N1—C7—C12—C11	-179.00 (15)
C2—C3—C4—C5	-0.3 (3)	C8—C7—C12—C11	55.5 (2)
C2—C3—C4—C13	-179.85 (19)	C7—N1—C14—C15	-103.9 (2)
C3—C4—C5—C6	0.6 (3)	S1—N1—C14—C15	87.5 (2)

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

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
C10—H10B \cdots O2 ⁱ	0.97	2.66	3.530 (3)	150.
C13—H13A \cdots O1 ⁱⁱ	0.96	2.60	3.512 (3)	159.

Symmetry codes: (i) $x, y-1, z$; (ii) $x, -y+5/2, z+1/2$.

Fig. 1

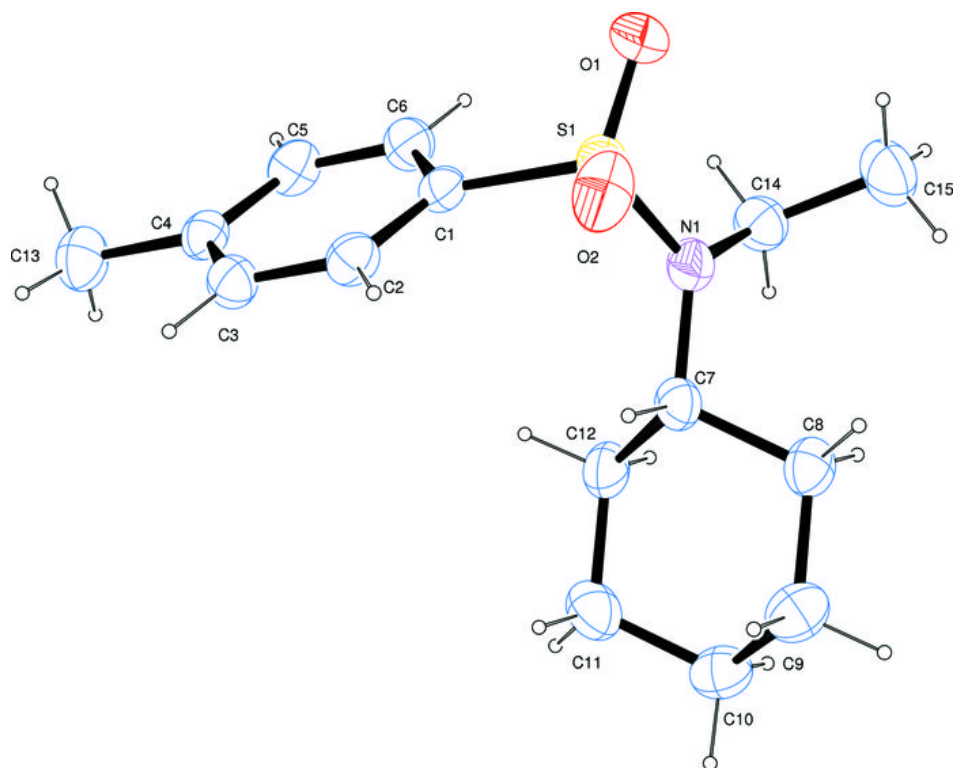


Fig. 2

