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[(3*R*,4*S*)-4-(4-Fluorophenyl)-1-methylpiperidin-3-yl]methyl 4-methylbenzenesulfonate

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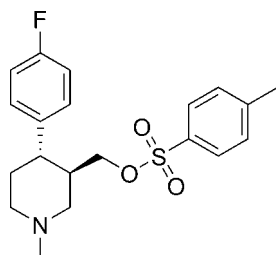
Received 19 May 2010; accepted 24 September 2010

Key indicators: single-crystal X-ray study; $T = 296$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å;
 R factor = 0.031; wR factor = 0.084; data-to-parameter ratio = 18.7.

In the title compound, $\text{C}_{20}\text{H}_{24}\text{FNO}_3\text{S}$, the piperidine ring adopts a chair conformation. The dihedral angle between the aromatic rings is 47.01 (17°).

Related literature

For general background to the design and synthesis of vinyl sulfonate derivatives, see: Curzons (2003), Segura *et al.* (2003). For related structures, see: Wang & Kanagawa (1997).



Experimental

Crystal data

$\text{C}_{20}\text{H}_{24}\text{FNO}_3\text{S}$
 $M_r = 377.46$

Monoclinic, $P2_1$
 $a = 9.1590$ (4) Å

$b = 10.0764$ (5) Å
 $c = 10.7644$ (6) Å
 $\beta = 95.718$ (1°)
 $V = 988.50$ (9) Å³
 $Z = 2$

Mo $K\alpha$ radiation $\mu = 0.19$ mm⁻¹ $T = 296$ K $0.32 \times 0.26 \times 0.20$ mm

Data collection

Rigaku R-Axis RAPID
diffractometer
Absorption correction: multi-scan
(*ABSCOR*; Higashi, 1995)
 $T_{\min} = 0.931$, $T_{\max} = 0.963$

9742 measured reflections
4457 independent reflections
3114 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.021$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.031$
 $wR(F^2) = 0.084$
 $S = 1.00$
4457 reflections
238 parameters
1 restraint

H-atom parameters constrained
 $\Delta\rho_{\max} = 0.17$ e Å⁻³
 $\Delta\rho_{\min} = -0.14$ e Å⁻³
Absolute structure: Flack (1983),
2086 Friedel pairs
Flack parameter: 0.05 (6)

Data collection: *PROCESS-AUTO* (Rigaku, 2006); cell refinement: *PROCESS-AUTO*; data reduction: *CrystalStructure* (Rigaku/MSK, 2007); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

Mr Jian-ming Gu of the X-ray crystallography facility of Zhejiang University is acknowledged for his assistance with the crystal structure analysis.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: KJ2148).

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supporting information

Acta Cryst. (2010). E66, o2660 [doi:10.1107/S1600536810038249]

[(3*R*,4*S*)-4-(4-Fluorophenyl)-1-methylpiperidin-3-yl]methyl 4-methylbenzenesulfonate**Jianfeng Qi, Hanjing Chen and Chen Zhang****S1. Comment**

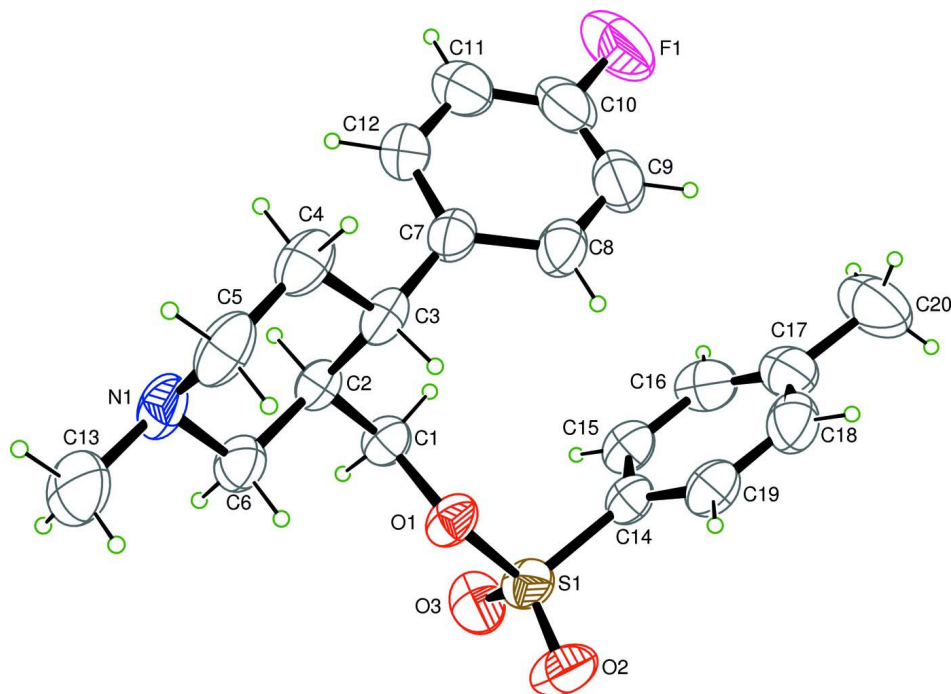
The title compound is a useful intermediate in preparing paroxetine [(3*S*,4*R*)-4-(4-fluorophenyl)-3-(3,4-methylenedioxyphenoxy-methyl)-piperidine]. Paroxetine is a well-known selective serotonin reuptake inhibitor (SSRI) antidepressant, used world wide in therapeutics (Segura *et al.*, 2003). In view of the above, ((3*R*,4*S*)-4-(4-fluorophenyl)-1-methylpiperidin-3-yl)methyl 4-methylbenzenesulfonate was synthesized and its crystal structure is reported here. A perspective view of the structure with the atomic numbering scheme is shown in Fig. 1. The dihedral angle between the two benzene rings is 47.01 (17)°. The piperidine ring adopts a chair conformation. The piperidine ring contains three planes (C2/C4/C5/C6, C3/C4/N1/C6, C2/C3/C5/N1), the first one of which is more planar than the other two.

S2. Experimental

To a stirred solution of trans-(-)-paroxo (10 g) in dichloromethane (50 ml) triethylamine (7 ml) was added. The mixture was cooled to 268 K. Toluenesulfonyl chloride (12 g) was slowly added and stirred for 1 h at 268 K. Methanesulfonic acid (4 ml) was then added gradually and the mixture was concentrated at about 323 K at atmospheric pressure. The residue was taken up in toluene, water was added and this was stirred for 30 minutes. The top toluene layer was separated. The pH of the aqueous layer was then adjusted to 9.0 with a saturated NaHCO₃ solution. The product was filtered, washed with water and dried to yield the title compound (13 g) as a white to off-white solid (m.p. 380-381 K).

S3. Refinement

All H atoms were placed in calculated positions, with C—H distances in the range 0.93-0.98 and included in the final cycles of refinement in the riding-model approximation, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ or $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$.

**Figure 1**

The molecular structure of the title compound showing atom labels and 40% probability displacement ellipsoids.

[(3*R*,4*S*)-4-(4-Fluorophenyl)-1-methylpiperidin-3-yl]methyl 4-methylbenzenesulfonate

Crystal data

$C_{20}H_{24}FNO_3S$

$M_r = 377.46$

Monoclinic, $P2_1$

Hall symbol: $P\ 2_1yb$

$a = 9.1590(4)\ \text{\AA}$

$b = 10.0764(5)\ \text{\AA}$

$c = 10.7644(6)\ \text{\AA}$

$\beta = 95.718(1)^\circ$

$V = 988.50(9)\ \text{\AA}^3$

$Z = 2$

$F(000) = 400$

$D_x = 1.268\ \text{Mg m}^{-3}$

Melting point: 380 K

Mo $K\alpha$ radiation, $\lambda = 0.71073\ \text{\AA}$

Cell parameters from 7787 reflections

$\theta = 3.0\text{--}27.4^\circ$

$\mu = 0.19\ \text{mm}^{-1}$

$T = 296\ \text{K}$

Chunk, yellow

$0.32 \times 0.26 \times 0.20\ \text{mm}$

Data collection

Rigaku R-AXIS RAPID
diffractometer

Radiation source: rolling anode

Graphite monochromator

Detector resolution: $10.00\ \text{pixels mm}^{-1}$

ω scans

Absorption correction: multi-scan

(*ABSCOR*; Higashi, 1995)

$T_{\min} = 0.931$, $T_{\max} = 0.963$

9742 measured reflections

4457 independent reflections

3114 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.021$

$\theta_{\max} = 27.4^\circ$, $\theta_{\min} = 3.0^\circ$

$h = -11 \rightarrow 11$

$k = -13 \rightarrow 13$

$l = -13 \rightarrow 13$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.031$

$wR(F^2) = 0.084$

$S = 1.00$

4457 reflections

238 parameters

1 restraint

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

$w = 1/[\sigma^2(F_o^2) + (0.0395P)^2 + 0.063P]$

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

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

$\Delta\rho_{\max} = 0.17 \text{ e } \text{\AA}^{-3}$

$\Delta\rho_{\min} = -0.14 \text{ e } \text{\AA}^{-3}$

Extinction correction: *SHELXL97* (Sheldrick,
2008), $F_c^* = kF_c[1 + 0.001xF_c^2\lambda^3/\sin(2\theta)]^{-1/4}$

Extinction coefficient: 0.0052 (13)

Absolute structure: Flack (1983), 2086 Friedel
pairs

Absolute structure parameter: 0.05 (6)

Special details

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds 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 $F^2 > \sigma(F^2)$ 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)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
S1	0.34071 (5)	0.65560 (6)	0.19254 (4)	0.05870 (14)
O1	0.37947 (12)	0.66249 (15)	0.33871 (9)	0.0550 (3)
C14	0.51158 (17)	0.6533 (2)	0.13267 (13)	0.0496 (4)
O2	0.2721 (2)	0.52973 (18)	0.17277 (14)	0.0832 (5)
C2	0.4586 (2)	0.77398 (17)	0.53214 (15)	0.0481 (4)
H2	0.4844	0.8619	0.5665	0.058*
O3	0.26443 (18)	0.77282 (18)	0.15090 (14)	0.0788 (5)
N1	0.3547 (2)	0.71908 (16)	0.72847 (16)	0.0659 (5)
C3	0.5921 (2)	0.68346 (16)	0.56744 (16)	0.0533 (5)
H3	0.5703	0.5965	0.5292	0.064*
C7	0.7277 (2)	0.73680 (18)	0.51511 (18)	0.0568 (5)
C4	0.6116 (2)	0.6639 (3)	0.70958 (16)	0.0684 (5)
H4A	0.6855	0.5963	0.7303	0.082*
H4B	0.6463	0.7460	0.7492	0.082*
C1	0.4245 (2)	0.79152 (17)	0.39340 (16)	0.0507 (4)
H1A	0.3462	0.8558	0.3764	0.061*
H1B	0.5105	0.8238	0.3571	0.061*
C15	0.5673 (2)	0.7700 (2)	0.08678 (18)	0.0595 (5)
H15	0.5152	0.8491	0.0888	0.071*
C6	0.3242 (2)	0.7278 (2)	0.59338 (18)	0.0615 (5)
H6A	0.2441	0.7895	0.5729	0.074*
H6B	0.2937	0.6414	0.5604	0.074*

C5	0.4712 (3)	0.62325 (19)	0.75996 (19)	0.0737 (7)
H5A	0.4409	0.5373	0.7260	0.088*
H5B	0.4878	0.6148	0.8500	0.088*
C8	0.7843 (2)	0.6761 (3)	0.41456 (18)	0.0696 (6)
H8	0.7424	0.5975	0.3828	0.083*
C19	0.5895 (3)	0.5365 (2)	0.12864 (18)	0.0630 (5)
H19	0.5529	0.4583	0.1595	0.076*
C9	0.9024 (3)	0.7304 (3)	0.3602 (2)	0.0862 (7)
H9	0.9399	0.6887	0.2931	0.103*
C17	0.7790 (2)	0.6507 (4)	0.03166 (18)	0.0795 (6)
C18	0.7223 (3)	0.5368 (3)	0.0783 (2)	0.0792 (7)
H18	0.7748	0.4580	0.0759	0.095*
C20	0.9241 (3)	0.6477 (6)	-0.0255 (3)	0.1361 (12)
H20A	0.9107	0.6047	-0.1054	0.204*
H20B	0.9953	0.5998	0.0285	0.204*
H20C	0.9579	0.7369	-0.0357	0.204*
F1	1.07463 (15)	0.9031 (2)	0.35259 (18)	0.1293 (7)
C13	0.2204 (3)	0.6809 (4)	0.7843 (2)	0.1021 (10)
H13A	0.1910	0.5934	0.7567	0.153*
H13B	0.1435	0.7428	0.7590	0.153*
H13C	0.2392	0.6815	0.8737	0.153*
C12	0.7948 (2)	0.8534 (2)	0.5606 (2)	0.0732 (6)
H12	0.7603	0.8954	0.6288	0.088*
C10	0.9617 (2)	0.8460 (3)	0.4074 (3)	0.0858 (7)
C16	0.7006 (3)	0.7662 (3)	0.0384 (2)	0.0751 (7)
H16	0.7389	0.8444	0.0092	0.090*
C11	0.9115 (3)	0.9079 (3)	0.5067 (3)	0.0885 (7)
H11	0.9551	0.9859	0.5379	0.106*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0645 (3)	0.0783 (3)	0.0327 (2)	-0.0102 (3)	0.00190 (16)	0.0008 (3)
O1	0.0736 (7)	0.0586 (6)	0.0328 (5)	-0.0118 (8)	0.0056 (5)	0.0006 (7)
C14	0.0649 (9)	0.0540 (9)	0.0294 (7)	-0.0033 (11)	0.0024 (6)	0.0024 (9)
O2	0.0960 (12)	0.1015 (12)	0.0521 (9)	-0.0489 (10)	0.0078 (8)	-0.0117 (8)
C2	0.0602 (10)	0.0463 (9)	0.0372 (9)	-0.0004 (8)	0.0026 (7)	-0.0028 (7)
O3	0.0729 (10)	0.1132 (13)	0.0492 (9)	0.0227 (9)	0.0005 (7)	0.0127 (9)
N1	0.0924 (13)	0.0687 (10)	0.0391 (9)	-0.0094 (9)	0.0187 (8)	-0.0068 (7)
C3	0.0747 (11)	0.0445 (11)	0.0392 (9)	0.0079 (9)	-0.0017 (8)	-0.0071 (7)
C7	0.0597 (11)	0.0611 (11)	0.0475 (11)	0.0165 (9)	-0.0053 (9)	-0.0066 (9)
C4	0.0977 (13)	0.0628 (10)	0.0419 (9)	0.0124 (14)	-0.0073 (9)	0.0007 (11)
C1	0.0588 (10)	0.0501 (10)	0.0429 (10)	-0.0019 (8)	0.0036 (8)	0.0031 (7)
C15	0.0787 (14)	0.0583 (11)	0.0409 (10)	-0.0056 (10)	0.0027 (9)	0.0054 (9)
C6	0.0715 (13)	0.0689 (11)	0.0452 (11)	-0.0076 (10)	0.0106 (9)	-0.0042 (8)
C5	0.1250 (19)	0.0585 (15)	0.0376 (10)	-0.0056 (12)	0.0081 (11)	0.0020 (8)
C8	0.0716 (12)	0.0842 (15)	0.0506 (11)	0.0210 (13)	-0.0051 (9)	-0.0139 (12)
C19	0.0824 (15)	0.0587 (12)	0.0467 (12)	-0.0019 (11)	0.0004 (10)	-0.0012 (9)

C9	0.0689 (14)	0.130 (2)	0.0600 (15)	0.0322 (14)	0.0061 (11)	-0.0080 (13)
C17	0.0626 (11)	0.1280 (19)	0.0472 (11)	-0.0052 (18)	0.0019 (8)	-0.0036 (16)
C18	0.0810 (17)	0.0940 (17)	0.0603 (14)	0.0232 (14)	-0.0049 (12)	-0.0148 (13)
C20	0.0691 (14)	0.253 (4)	0.0883 (19)	-0.003 (3)	0.0194 (13)	-0.009 (3)
F1	0.0667 (8)	0.1822 (18)	0.1440 (16)	0.0034 (10)	0.0364 (9)	0.0170 (13)
C13	0.122 (2)	0.131 (3)	0.0591 (14)	-0.040 (2)	0.0382 (13)	-0.0086 (15)
C12	0.0656 (12)	0.0774 (14)	0.0771 (15)	0.0049 (11)	0.0098 (11)	-0.0234 (11)
C10	0.0482 (12)	0.118 (2)	0.0909 (19)	0.0140 (13)	0.0073 (12)	0.0047 (16)
C16	0.0785 (16)	0.0988 (18)	0.0482 (13)	-0.0271 (14)	0.0079 (11)	0.0101 (12)
C11	0.0601 (13)	0.0933 (17)	0.112 (2)	-0.0020 (12)	0.0077 (13)	-0.0169 (15)

Geometric parameters (Å, °)

S1—O3	1.4219 (18)	C6—H6A	0.9700
S1—O2	1.4220 (17)	C6—H6B	0.9700
S1—O1	1.5796 (10)	C5—H5A	0.9700
S1—C14	1.7510 (16)	C5—H5B	0.9700
O1—C1	1.469 (2)	C8—C9	1.392 (3)
C14—C19	1.379 (3)	C8—H8	0.9300
C14—C15	1.392 (3)	C19—C18	1.380 (4)
C2—C1	1.505 (2)	C19—H19	0.9300
C2—C6	1.527 (3)	C9—C10	1.362 (4)
C2—C3	1.542 (2)	C9—H9	0.9300
C2—H2	0.9800	C17—C16	1.374 (4)
N1—C5	1.454 (3)	C17—C18	1.375 (4)
N1—C6	1.456 (2)	C17—C20	1.519 (3)
N1—C13	1.474 (3)	C18—H18	0.9300
C3—C7	1.513 (3)	C20—H20A	0.9600
C3—C4	1.535 (2)	C20—H20B	0.9600
C3—H3	0.9800	C20—H20C	0.9600
C7—C8	1.387 (3)	F1—C10	1.368 (3)
C7—C12	1.392 (3)	C13—H13A	0.9600
C4—C5	1.501 (3)	C13—H13B	0.9600
C4—H4A	0.9700	C13—H13C	0.9600
C4—H4B	0.9700	C12—C11	1.380 (3)
C1—H1A	0.9700	C12—H12	0.9300
C1—H1B	0.9700	C10—C11	1.356 (4)
C15—C16	1.375 (3)	C16—H16	0.9300
C15—H15	0.9300	C11—H11	0.9300
O3—S1—O2	119.85 (9)	C2—C6—H6B	109.3
O3—S1—O1	109.40 (9)	H6A—C6—H6B	108.0
O2—S1—O1	103.91 (9)	N1—C5—C4	111.70 (17)
O3—S1—C14	108.92 (10)	N1—C5—H5A	109.3
O2—S1—C14	109.32 (12)	C4—C5—H5A	109.3
O1—S1—C14	104.29 (6)	N1—C5—H5B	109.3
C1—O1—S1	117.67 (11)	C4—C5—H5B	109.3
C19—C14—C15	120.08 (17)	H5A—C5—H5B	107.9

C19—C14—S1	120.46 (16)	C7—C8—C9	121.4 (2)
C15—C14—S1	119.45 (17)	C7—C8—H8	119.3
C1—C2—C6	111.55 (15)	C9—C8—H8	119.3
C1—C2—C3	113.27 (14)	C14—C19—C18	119.3 (2)
C6—C2—C3	111.54 (15)	C14—C19—H19	120.4
C1—C2—H2	106.7	C18—C19—H19	120.4
C6—C2—H2	106.7	C10—C9—C8	118.5 (2)
C3—C2—H2	106.7	C10—C9—H9	120.8
C5—N1—C6	109.66 (15)	C8—C9—H9	120.8
C5—N1—C13	110.76 (19)	C16—C17—C18	117.99 (19)
C6—N1—C13	109.76 (18)	C16—C17—C20	121.4 (3)
C7—C3—C4	113.49 (15)	C18—C17—C20	120.6 (4)
C7—C3—C2	110.99 (14)	C17—C18—C19	121.7 (2)
C4—C3—C2	109.40 (15)	C17—C18—H18	119.2
C7—C3—H3	107.6	C19—C18—H18	119.2
C4—C3—H3	107.6	C17—C20—H20A	109.5
C2—C3—H3	107.6	C17—C20—H20B	109.5
C8—C7—C12	117.4 (2)	H20A—C20—H20B	109.5
C8—C7—C3	121.21 (18)	C17—C20—H20C	109.5
C12—C7—C3	121.27 (18)	H20A—C20—H20C	109.5
C5—C4—C3	112.10 (16)	H20B—C20—H20C	109.5
C5—C4—H4A	109.2	N1—C13—H13A	109.5
C3—C4—H4A	109.2	N1—C13—H13B	109.5
C5—C4—H4B	109.2	H13A—C13—H13B	109.5
C3—C4—H4B	109.2	N1—C13—H13C	109.5
H4A—C4—H4B	107.9	H13A—C13—H13C	109.5
O1—C1—C2	108.42 (13)	H13B—C13—H13C	109.5
O1—C1—H1A	110.0	C11—C12—C7	121.4 (2)
C2—C1—H1A	110.0	C11—C12—H12	119.3
O1—C1—H1B	110.0	C7—C12—H12	119.3
C2—C1—H1B	110.0	C11—C10—C9	122.2 (2)
H1A—C1—H1B	108.4	C11—C10—F1	118.5 (3)
C16—C15—C14	118.8 (2)	C9—C10—F1	119.3 (3)
C16—C15—H15	120.6	C17—C16—C15	122.2 (2)
C14—C15—H15	120.6	C17—C16—H16	118.9
N1—C6—C2	111.50 (16)	C15—C16—H16	118.9
N1—C6—H6A	109.3	C10—C11—C12	119.1 (2)
C2—C6—H6A	109.3	C10—C11—H11	120.4
N1—C6—H6B	109.3	C12—C11—H11	120.4
O3—S1—O1—C1	-38.54 (14)	C13—N1—C6—C2	177.1 (2)
O2—S1—O1—C1	-167.66 (14)	C1—C2—C6—N1	-176.31 (15)
C14—S1—O1—C1	77.84 (14)	C3—C2—C6—N1	55.9 (2)
O3—S1—C14—C19	-161.67 (15)	C6—N1—C5—C4	61.7 (2)
O2—S1—C14—C19	-29.00 (16)	C13—N1—C5—C4	-177.04 (18)
O1—S1—C14—C19	81.61 (16)	C3—C4—C5—N1	-57.2 (2)
O3—S1—C14—C15	17.09 (17)	C12—C7—C8—C9	-0.8 (3)
O2—S1—C14—C15	149.76 (15)	C3—C7—C8—C9	175.15 (18)

O1—S1—C14—C15	-99.62 (15)	C15—C14—C19—C18	-0.3 (3)
C1—C2—C3—C7	57.92 (18)	S1—C14—C19—C18	178.51 (16)
C6—C2—C3—C7	-175.24 (14)	C7—C8—C9—C10	-0.4 (3)
C1—C2—C3—C4	-176.09 (16)	C16—C17—C18—C19	1.1 (3)
C6—C2—C3—C4	-49.2 (2)	C20—C17—C18—C19	-178.8 (2)
C4—C3—C7—C8	129.5 (2)	C14—C19—C18—C17	-0.2 (3)
C2—C3—C7—C8	-106.83 (19)	C8—C7—C12—C11	1.1 (3)
C4—C3—C7—C12	-54.7 (2)	C3—C7—C12—C11	-174.8 (2)
C2—C3—C7—C12	68.9 (2)	C8—C9—C10—C11	1.3 (4)
C7—C3—C4—C5	174.59 (17)	C8—C9—C10—F1	-177.8 (2)
C2—C3—C4—C5	50.0 (2)	C18—C17—C16—C15	-1.7 (3)
S1—O1—C1—C2	-178.87 (11)	C20—C17—C16—C15	178.3 (2)
C6—C2—C1—O1	-61.63 (18)	C14—C15—C16—C17	1.3 (3)
C3—C2—C1—O1	65.21 (18)	C9—C10—C11—C12	-1.1 (4)
C19—C14—C15—C16	-0.3 (3)	F1—C10—C11—C12	178.1 (2)
S1—C14—C15—C16	-179.04 (15)	C7—C12—C11—C10	-0.2 (4)
C5—N1—C6—C2	-61.0 (2)		
