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N-(2-Amino-3,5-dibromobenzyl)-N-methylcyclohexan-1-aminium *p*-toluenesulfonate

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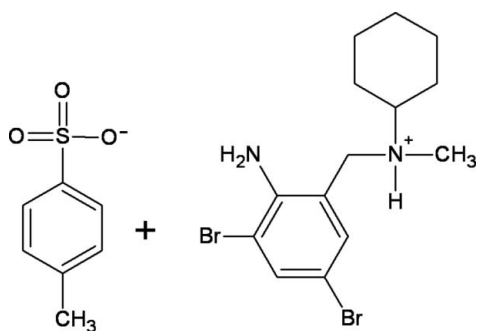
Received 26 June 2011; accepted 7 July 2011

Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.006$ Å; R factor = 0.047; wR factor = 0.114; data-to-parameter ratio = 16.9.

The title compound, $\text{C}_{14}\text{H}_{21}\text{Br}_2\text{N}_2^+\cdot\text{C}_7\text{H}_7\text{O}_3\text{S}^-$, features a salt of protonated bromhexine, a pharmaceutical used in the treatment of respiratory disorders, and the *p*-toluenesulfonate anion. The crystal packing is stabilized by intermolecular $\text{N}-\text{H}\cdots\text{O}$, $\text{N}-\text{H}\cdots\text{Br}$ and $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds.

Related literature

For salts of bromhexine, see: Koo *et al.* (1984); Shimizu & Nishigaki (1983); Shimizu *et al.* (1983, 1984).



Experimental

Crystal data

$\text{C}_{14}\text{H}_{21}\text{Br}_2\text{N}_2^+\cdot\text{C}_7\text{H}_7\text{O}_3\text{S}^-$

$M_r = 548.32$

Monoclinic, $P2_1/c$

$a = 14.008$ (5) Å

$b = 10.404$ (5) Å

$c = 17.157$ (5) Å

$\beta = 110.148$ (5)°

$V = 2347.4$ (16) Å³

$Z = 4$

Mo $K\alpha$ radiation

$\mu = 3.57$ mm⁻¹

$T = 293$ K

$0.30 \times 0.10 \times 0.08$ mm

Data collection

Bruker SMART CCD area-detector diffractometer

Absorption correction: multi-scan (SADABS; Sheldrick, 1996)

$T_{\min} = 0.414$, $T_{\max} = 0.764$

14496 measured reflections

4608 independent reflections

2343 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.052$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.047$

$wR(F^2) = 0.114$

$S = 1.00$

4608 reflections

272 parameters

4 restraints

H atoms treated by a mixture of independent and constrained refinement

$\Delta\rho_{\text{max}} = 0.78$ e Å⁻³

$\Delta\rho_{\text{min}} = -0.71$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

| $D-H\cdots A$ | $D-H$ | $H\cdots A$ | $D\cdots A$ | $D-H\cdots A$ |
|---|----------|-------------|-------------|---------------|
| $\text{N1}-\text{H1}\cdots\text{O2}^{\text{i}}$ | 0.85 (3) | 1.93 (4) | 2.756 (4) | 161 (4) |
| $\text{N2}-\text{H2A}\cdots\text{O1}^{\text{ii}}$ | 0.85 (3) | 2.11 (3) | 2.926 (4) | 162 (4) |
| $\text{N2}-\text{H2B}\cdots\text{Br1}$ | 0.84 (2) | 2.67 (3) | 3.068 (3) | 111 (2) |
| $\text{C7}-\text{H7A}\cdots\text{O2}$ | 0.97 | 2.47 | 3.257 (5) | 138 |

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

Data collection: SMART (Bruker, 1998); cell refinement: SAINT (Bruker, 1998); 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 for Windows (Farrugia, 1997); software used to prepare material for publication: PLATON (Spek, 2009).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BT5564).

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supporting information

Acta Cryst. (2011). E67, o2032 [doi:10.1107/S1600536811027358]

***N*-(2-Amino-3,5-dibromobenzyl)-*N*-methylcyclohexan-1-aminium *p*-toluene-sulfonate**

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S1. Comment

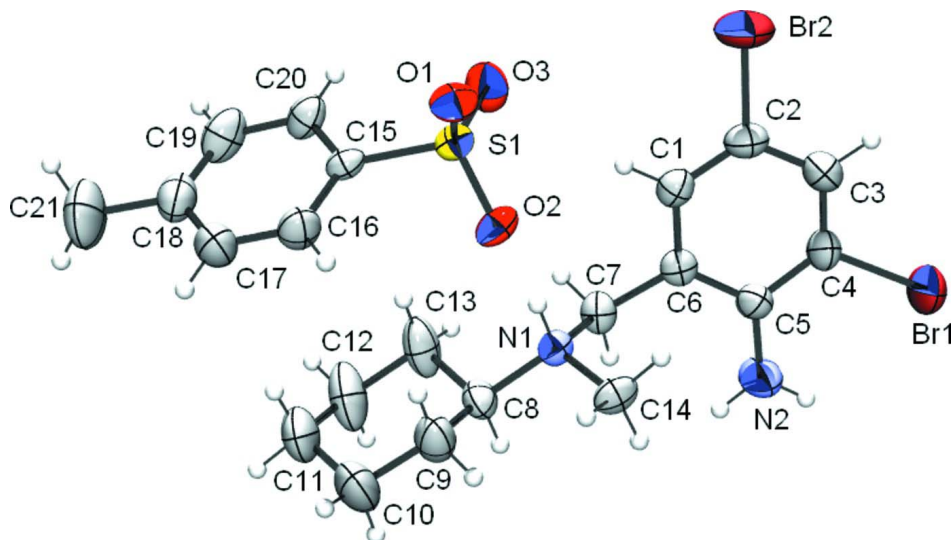
There are only three crystal structures on Bromhexine reported in the literature. Analysis of all reported structure of Bromhexine indicates that the *N*-methyl amino group of Bromhexine is basic in nature and forms a salt with HCl (Koo *et al.*, 1984), salicylic acid (Shimizu *et al.*, 1984) and 1,2-benzisothiazol-3(2*H*)-one 1,1-dioxide (Shimizu *et al.*, 1983). A similar case is found in the current study where Bromhexine forms a salt with paratoluene sulfonic acid by transferring a proton from sulfonic acid group to *N*-methyl amino group. The crystal structure is stabilized by N—H \cdots O, N—H \cdots Br and C—H \cdots O intermolecular interactions.

S2. Experimental

An equimolar ratio (1:1) of Bromhexine and para-toluene sulfonic acid were dissolved in ethanol and kept for crystallization at room temperature yielding plate shape crystals.

S3. Refinement

In the absence of significant anomalous dispersion effects, Friedel pairs were merged. H1 was freely refined, but all other H atoms were positioned geometrically and refined using a riding model.

**Figure 1**

View of the title compound with the atom numbering scheme. Displacement ellipsoids for non-H atoms are drawn at the 50% probability level.

N-(2-Amino-3,5-dibromobenzyl)-*N*-methylcyclohexan-1-aminium *p*-toluenesulfonate

Crystal data

$C_{14}H_{21}Br_2N_2^+ \cdot C_7H_7O_3S^-$

$M_r = 548.32$

Monoclinic, $P2_1/c$

Hall symbol: -P 2ybc

$a = 14.008 (5) \text{ \AA}$

$b = 10.404 (5) \text{ \AA}$

$c = 17.157 (5) \text{ \AA}$

$\beta = 110.148 (5)^\circ$

$V = 2347.4 (16) \text{ \AA}^3$

$Z = 4$

$F(000) = 1112$

$D_x = 1.551 \text{ Mg m}^{-3}$

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

Cell parameters from 2343 reflections

$\theta = 2.3\text{--}26.0^\circ$

$\mu = 3.57 \text{ mm}^{-1}$

$T = 293 \text{ K}$

Needle, colorless

$0.30 \times 0.10 \times 0.08 \text{ mm}$

Data collection

Bruker SMART CCD area-detector
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

φ and ω scans

Absorption correction: multi-scan

(*SADABS*; Sheldrick, 1996)

$T_{\min} = 0.414$, $T_{\max} = 0.764$

14496 measured reflections

4608 independent reflections

2343 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.052$

$\theta_{\max} = 26.0^\circ$, $\theta_{\min} = 2.3^\circ$

$h = -15 \rightarrow 17$

$k = -11 \rightarrow 12$

$l = -21 \rightarrow 21$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.047$

$wR(F^2) = 0.114$

$S = 1.00$

4608 reflections

272 parameters

4 restraints

Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier
map

Hydrogen site location: inferred from
neighbouring sites

H atoms treated by a mixture of independent
and constrained refinement
 $w = 1/[\sigma^2(F_o^2) + (0.050P)^2]$
where $P = (F_o^2 + 2F_c^2)/3$

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

$$\Delta\rho_{\max} = 0.78 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\min} = -0.71 \text{ e } \text{\AA}^{-3}$$

Special details

Geometry. Bond distances, angles *etc.* have been calculated using the rounded fractional coordinates. All su's are estimated from the variances of the (full) variance-covariance matrix. The cell e.s.d.'s are taken into account in the estimation of distances, angles and torsion angles

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> | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|-----|--------------|--------------|--------------|----------------------------------|
| Br1 | 0.06429 (3) | 0.35949 (7) | -0.08435 (3) | 0.0854 (3) |
| Br2 | 0.14386 (4) | 0.34597 (7) | 0.25939 (3) | 0.0893 (3) |
| N1 | 0.5129 (2) | 0.2898 (3) | 0.1723 (2) | 0.0330 (11) |
| N2 | 0.2969 (2) | 0.3732 (5) | -0.0301 (2) | 0.0666 (16) |
| C1 | 0.2959 (3) | 0.3727 (4) | 0.1844 (2) | 0.0411 (14) |
| C2 | 0.1945 (3) | 0.3605 (4) | 0.1704 (2) | 0.0500 (16) |
| C3 | 0.1249 (3) | 0.3565 (4) | 0.0899 (3) | 0.0548 (18) |
| C4 | 0.1613 (3) | 0.3629 (4) | 0.0259 (2) | 0.0466 (14) |
| C5 | 0.2651 (3) | 0.3701 (4) | 0.0360 (2) | 0.0411 (14) |
| C6 | 0.3321 (3) | 0.3795 (4) | 0.1181 (2) | 0.0356 (12) |
| C7 | 0.4449 (3) | 0.4040 (4) | 0.1372 (2) | 0.0406 (14) |
| C8 | 0.6233 (3) | 0.3192 (4) | 0.1819 (2) | 0.0424 (14) |
| C9 | 0.6906 (3) | 0.2040 (5) | 0.2153 (3) | 0.0628 (16) |
| C10 | 0.8014 (3) | 0.2361 (6) | 0.2280 (3) | 0.095 (3) |
| C11 | 0.8394 (3) | 0.3534 (5) | 0.2807 (3) | 0.080 (2) |
| C12 | 0.7710 (3) | 0.4670 (5) | 0.2473 (3) | 0.084 (2) |
| C13 | 0.6605 (3) | 0.4379 (4) | 0.2365 (3) | 0.0639 (18) |
| C14 | 0.4737 (3) | 0.1716 (4) | 0.1239 (2) | 0.0419 (14) |
| S1 | 0.48360 (7) | 0.75893 (10) | 0.09405 (5) | 0.0358 (3) |
| O1 | 0.50153 (19) | 0.6560 (3) | 0.04343 (15) | 0.0440 (10) |
| O2 | 0.48544 (18) | 0.7100 (3) | 0.17409 (14) | 0.0436 (9) |
| O3 | 0.5480 (2) | 0.8695 (3) | 0.10102 (18) | 0.0572 (11) |
| C15 | 0.3570 (3) | 0.8101 (4) | 0.0421 (2) | 0.0317 (11) |
| C16 | 0.2762 (3) | 0.7309 (4) | 0.0396 (2) | 0.0486 (16) |
| C17 | 0.1786 (3) | 0.7695 (5) | 0.0001 (3) | 0.0632 (19) |
| C18 | 0.1571 (3) | 0.8892 (5) | -0.0385 (3) | 0.0596 (19) |
| C19 | 0.2376 (3) | 0.9666 (5) | -0.0353 (2) | 0.0599 (17) |
| C20 | 0.3377 (3) | 0.9273 (4) | 0.0037 (2) | 0.0455 (16) |
| C21 | 0.0474 (3) | 0.9326 (6) | -0.0849 (3) | 0.097 (2) |
| H1 | 0.510 (3) | 0.283 (4) | 0.221 (2) | 0.039 (12)* |
| H1A | 0.34145 | 0.37656 | 0.23856 | 0.0493* |
| H2A | 0.3594 (16) | 0.376 (5) | -0.025 (2) | 0.1165* |

| | | | | |
|------|-----------|-----------|--------------|---------|
| H2B | 0.257 (2) | 0.362 (5) | -0.0791 (13) | 0.1165* |
| H3 | 0.05550 | 0.34954 | 0.08003 | 0.0656* |
| H7A | 0.46572 | 0.47432 | 0.17655 | 0.0483* |
| H7B | 0.45510 | 0.43113 | 0.08657 | 0.0483* |
| H8 | 0.62560 | 0.33869 | 0.12668 | 0.0509* |
| H9A | 0.68487 | 0.17775 | 0.26774 | 0.0753* |
| H9B | 0.66849 | 0.13292 | 0.17664 | 0.0753* |
| H10A | 0.80824 | 0.24959 | 0.17422 | 0.1137* |
| H10B | 0.84369 | 0.16334 | 0.25395 | 0.1137* |
| H11A | 0.84262 | 0.33587 | 0.33704 | 0.0963* |
| H11B | 0.90757 | 0.37357 | 0.28204 | 0.0963* |
| H12A | 0.79431 | 0.53895 | 0.28507 | 0.1008* |
| H12B | 0.77540 | 0.49181 | 0.19414 | 0.1008* |
| H13A | 0.65444 | 0.42325 | 0.29040 | 0.0774* |
| H13B | 0.61841 | 0.51114 | 0.21117 | 0.0774* |
| H14A | 0.51760 | 0.10072 | 0.14848 | 0.0629* |
| H14B | 0.40641 | 0.15390 | 0.12394 | 0.0629* |
| H14C | 0.47148 | 0.18343 | 0.06781 | 0.0629* |
| H16 | 0.28896 | 0.65083 | 0.06524 | 0.0582* |
| H17 | 0.12543 | 0.71510 | -0.00111 | 0.0757* |
| H19 | 0.22494 | 1.04752 | -0.05974 | 0.0719* |
| H20 | 0.39114 | 0.98030 | 0.00355 | 0.0546* |
| H21A | 0.00115 | 0.86721 | -0.08053 | 0.1456* |
| H21B | 0.03425 | 1.01093 | -0.06074 | 0.1456* |
| H21C | 0.03824 | 0.94686 | -0.14227 | 0.1456* |

Atomic displacement parameters (\AA^2)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|-----|-------------|------------|-------------|--------------|-------------|--------------|
| Br1 | 0.0448 (3) | 0.1514 (7) | 0.0473 (3) | 0.0073 (3) | -0.0002 (2) | -0.0107 (3) |
| Br2 | 0.0855 (4) | 0.1380 (7) | 0.0625 (3) | 0.0055 (4) | 0.0488 (3) | -0.0046 (3) |
| N1 | 0.0363 (18) | 0.030 (2) | 0.0331 (18) | -0.0035 (16) | 0.0126 (15) | -0.0005 (17) |
| N2 | 0.049 (2) | 0.114 (4) | 0.040 (2) | 0.006 (3) | 0.0193 (17) | 0.007 (2) |
| C1 | 0.043 (2) | 0.039 (3) | 0.038 (2) | 0.002 (2) | 0.0096 (18) | -0.001 (2) |
| C2 | 0.053 (3) | 0.061 (3) | 0.042 (2) | 0.001 (3) | 0.024 (2) | -0.002 (2) |
| C3 | 0.040 (2) | 0.073 (4) | 0.053 (3) | -0.001 (2) | 0.018 (2) | -0.006 (3) |
| C4 | 0.035 (2) | 0.059 (3) | 0.039 (2) | 0.005 (2) | 0.0042 (18) | -0.001 (2) |
| C5 | 0.038 (2) | 0.046 (3) | 0.039 (2) | 0.006 (2) | 0.0130 (18) | -0.002 (2) |
| C6 | 0.039 (2) | 0.026 (2) | 0.041 (2) | 0.001 (2) | 0.0128 (18) | 0.001 (2) |
| C7 | 0.039 (2) | 0.034 (3) | 0.044 (2) | -0.001 (2) | 0.0081 (18) | 0.004 (2) |
| C8 | 0.032 (2) | 0.053 (3) | 0.042 (2) | -0.004 (2) | 0.0124 (17) | 0.003 (2) |
| C9 | 0.045 (2) | 0.062 (3) | 0.072 (3) | 0.009 (3) | 0.008 (2) | -0.017 (3) |
| C10 | 0.045 (3) | 0.124 (6) | 0.107 (4) | 0.011 (4) | 0.016 (3) | -0.038 (4) |
| C11 | 0.041 (3) | 0.110 (5) | 0.080 (4) | -0.014 (3) | 0.008 (2) | -0.016 (4) |
| C12 | 0.046 (3) | 0.084 (4) | 0.101 (4) | -0.016 (3) | -0.002 (3) | 0.018 (4) |
| C13 | 0.040 (2) | 0.039 (3) | 0.096 (4) | -0.005 (2) | 0.002 (2) | 0.003 (3) |
| C14 | 0.059 (2) | 0.026 (3) | 0.042 (2) | -0.004 (2) | 0.019 (2) | -0.008 (2) |
| S1 | 0.0422 (5) | 0.0333 (7) | 0.0320 (5) | -0.0031 (6) | 0.0131 (4) | 0.0001 (5) |

| | | | | | | |
|-----|-------------|-------------|-------------|--------------|-------------|--------------|
| O1 | 0.0552 (16) | 0.044 (2) | 0.0405 (15) | 0.0049 (14) | 0.0263 (13) | -0.0070 (14) |
| O2 | 0.0583 (16) | 0.0459 (19) | 0.0260 (13) | 0.0045 (15) | 0.0137 (12) | 0.0001 (13) |
| O3 | 0.0478 (16) | 0.045 (2) | 0.070 (2) | -0.0120 (16) | 0.0089 (14) | 0.0067 (16) |
| C15 | 0.043 (2) | 0.028 (2) | 0.0253 (18) | -0.002 (2) | 0.0133 (16) | -0.0019 (19) |
| C16 | 0.051 (3) | 0.041 (3) | 0.051 (2) | -0.002 (3) | 0.014 (2) | 0.012 (2) |
| C17 | 0.045 (3) | 0.079 (4) | 0.062 (3) | -0.005 (3) | 0.014 (2) | 0.016 (3) |
| C18 | 0.050 (3) | 0.078 (4) | 0.045 (3) | 0.012 (3) | 0.009 (2) | 0.001 (3) |
| C19 | 0.076 (3) | 0.039 (3) | 0.052 (3) | 0.009 (3) | 0.006 (2) | 0.003 (2) |
| C20 | 0.053 (3) | 0.040 (3) | 0.035 (2) | -0.005 (2) | 0.0042 (19) | -0.001 (2) |
| C21 | 0.059 (3) | 0.114 (5) | 0.096 (4) | 0.027 (3) | -0.002 (3) | 0.012 (4) |

Geometric parameters (Å, °)

| | | | |
|--------------------------|-----------|--------------------------|-----------|
| Br1—C4 | 1.912 (4) | C7—H7B | 0.9700 |
| Br2—C2 | 1.898 (4) | C8—H8 | 0.9800 |
| S1—O1 | 1.454 (3) | C9—H9B | 0.9700 |
| S1—O2 | 1.456 (3) | C9—H9A | 0.9700 |
| S1—O3 | 1.441 (3) | C10—H10B | 0.9700 |
| S1—C15 | 1.770 (4) | C10—H10A | 0.9700 |
| N1—C7 | 1.512 (5) | C11—H11A | 0.9700 |
| N1—C8 | 1.529 (5) | C11—H11B | 0.9700 |
| N1—C14 | 1.480 (5) | C12—H12A | 0.9700 |
| N2—C5 | 1.354 (5) | C12—H12B | 0.9700 |
| N1—H1 | 0.85 (3) | C13—H13A | 0.9700 |
| N2—H2B | 0.84 (2) | C13—H13B | 0.9700 |
| N2—H2A | 0.85 (3) | C14—H14B | 0.9600 |
| C1—C2 | 1.363 (6) | C14—H14C | 0.9600 |
| C1—C6 | 1.398 (5) | C14—H14A | 0.9600 |
| C2—C3 | 1.390 (6) | C15—C16 | 1.389 (6) |
| C3—C4 | 1.362 (6) | C15—C20 | 1.368 (6) |
| C4—C5 | 1.406 (6) | C16—C17 | 1.361 (6) |
| C5—C6 | 1.401 (5) | C17—C18 | 1.394 (7) |
| C6—C7 | 1.521 (6) | C18—C19 | 1.371 (7) |
| C8—C9 | 1.511 (7) | C18—C21 | 1.535 (7) |
| C8—C13 | 1.529 (6) | C19—C20 | 1.391 (6) |
| C9—C10 | 1.528 (7) | C16—H16 | 0.9300 |
| C10—C11 | 1.503 (8) | C17—H17 | 0.9300 |
| C11—C12 | 1.505 (7) | C19—H19 | 0.9300 |
| C12—C13 | 1.525 (7) | C20—H20 | 0.9300 |
| C1—H1A | 0.9300 | C21—H21A | 0.9600 |
| C3—H3 | 0.9300 | C21—H21B | 0.9600 |
| C7—H7A | 0.9700 | C21—H21C | 0.9600 |
| Br1...N2 | 3.068 (3) | H1A...H7A | 2.5400 |
| Br1...Br2 ⁱ | 3.880 (2) | H1A...O3 ^{viii} | 2.6500 |
| Br2...Br1 ⁱⁱ | 3.880 (2) | H2A...H7B | 2.0100 |
| Br1...H2B | 2.67 (3) | H2A...S1 ^{iv} | 3.16 (3) |
| Br2...H19 ⁱⁱⁱ | 3.1200 | H2A...O1 ^{iv} | 2.11 (3) |

| | | | |
|--------------------------|-----------|----------------------------|----------|
| S1...H14C ^{iv} | 3.1100 | H2A...C7 | 2.64 (3) |
| S1...H1 ^v | 3.15 (3) | H2B...Br1 | 2.67 (3) |
| S1...H2A ^{iv} | 3.16 (3) | H2B...H12B ^{iv} | 2.4100 |
| O1...C7 | 3.312 (5) | H3...H17 ^x | 2.5200 |
| O1...N2 ^{iv} | 2.926 (4) | H3...H21A ^x | 2.3900 |
| O2...N1 ^v | 2.756 (4) | H7A...H1A | 2.5400 |
| O2...C14 ^v | 3.337 (4) | H7A...C13 | 2.5900 |
| O2...C7 | 3.257 (5) | H7A...H13B | 2.0500 |
| O3...C14 ^{vi} | 3.376 (5) | H7A...O2 | 2.4700 |
| O1...H2A ^{iv} | 2.11 (3) | H7B...N2 | 2.5000 |
| O1...H8 ^{iv} | 2.8500 | H7B...O1 | 2.6000 |
| O1...H7B | 2.6000 | H7B...H8 | 2.4400 |
| O1...H7B ^{iv} | 2.6600 | H7B...O1 ^{iv} | 2.6600 |
| O1...H14C ^{iv} | 2.6600 | H7B...H2A | 2.0100 |
| O2...H13B | 2.7100 | H8...H12B | 2.5700 |
| O2...H16 | 2.8100 | H8...O1 ^{iv} | 2.8500 |
| O2...H7A | 2.4700 | H8...H7B | 2.4400 |
| O2...H9A ^v | 2.9100 | H8...H10A | 2.5800 |
| O2...H1 ^v | 1.93 (4) | H9A...O2 ^{viii} | 2.9100 |
| O3...H20 | 2.5300 | H9A...H1 | 2.5500 |
| O3...H1A ^v | 2.6500 | H9B...C14 | 2.5900 |
| O3...H14C ^{iv} | 2.8700 | H9B...H14A | 2.0300 |
| O3...H20 ^{vii} | 2.7300 | H10A...C18 ^{iv} | 2.9200 |
| O3...H14A ^{vi} | 2.6200 | H10A...C17 ^{iv} | 3.0700 |
| N1...O2 ^{viii} | 2.756 (4) | H10A...H8 | 2.5800 |
| N2...Br1 | 3.068 (3) | H11A...C17 ^{viii} | 2.9900 |
| N2...O1 ^{iv} | 2.926 (4) | H12B...H2B ^{iv} | 2.4100 |
| N2...H7B | 2.5000 | H12B...H8 | 2.5700 |
| C5...C14 | 3.470 (6) | H13A...H1 | 2.4500 |
| C7...O2 | 3.257 (5) | H13B...H7A | 2.0500 |
| C7...O1 | 3.312 (5) | H13B...O2 | 2.7100 |
| C14...C20 ^{ix} | 3.411 (6) | H13B...C7 | 2.5800 |
| C14...O2 ^{viii} | 3.337 (4) | H14A...O3 ^{ix} | 2.6200 |
| C14...O3 ^{ix} | 3.376 (5) | H14A...H9B | 2.0300 |
| C14...C5 | 3.470 (6) | H14A...C9 | 2.5400 |
| C20...C14 ^{vi} | 3.411 (6) | H14B...C5 | 3.0300 |
| C1...H1 | 2.99 (4) | H14B...C20 ^{ix} | 3.0600 |
| C3...H21A ^x | 2.8900 | H14B...C6 | 2.5600 |
| C5...H16 | 2.9600 | H14C...O3 ^{iv} | 2.8700 |
| C5...H14B | 3.0300 | H14C...O1 ^{iv} | 2.6600 |
| C6...H16 | 2.9600 | H14C...H20 ^{ix} | 2.4700 |
| C6...H14B | 2.5600 | H14C...S1 ^{iv} | 3.1100 |
| C7...H2A | 2.64 (3) | H16...O2 | 2.8100 |
| C7...H13B | 2.5800 | H16...C5 | 2.9600 |
| C9...H14A | 2.5400 | H16...C6 | 2.9600 |
| C13...H7A | 2.5900 | H17...H21A | 2.4000 |
| C14...H20 ^{ix} | 2.8100 | H17...H3 ^x | 2.5200 |
| C14...H9B | 2.5900 | H19...Br2 ^{xi} | 3.1200 |

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| C17...H11A ^v | 2.9900 | H20...O3 | 2.5300 |
| C17...H10A ^{iv} | 3.0700 | H20...C14 ^{vi} | 2.8100 |
| C18...H10A ^{iv} | 2.9200 | H20...H14C ^{vi} | 2.4700 |
| C20...H14B ^{vi} | 3.0600 | H20...O3 ^{vii} | 2.7300 |
| H1...H13A | 2.4500 | H21A...H17 | 2.4000 |
| H1...S1 ^{viii} | 3.15 (3) | H21A...C3 ^x | 2.8900 |
| H1...H9A | 2.5500 | H21A...H3 ^x | 2.3900 |
| H1...C1 | 2.99 (4) | H21B...H21B ^{xii} | 2.5900 |
| H1...O2 ^{viii} | 1.93 (4) | | |
| O2—S1—O3 | 113.17 (17) | C8—C9—H9A | 110.00 |
| O2—S1—C15 | 105.61 (17) | C10—C9—H9B | 109.00 |
| O3—S1—C15 | 106.95 (19) | C11—C10—H10A | 109.00 |
| O1—S1—O3 | 113.92 (18) | C11—C10—H10B | 109.00 |
| O1—S1—C15 | 105.66 (17) | C9—C10—H10B | 109.00 |
| O1—S1—O2 | 110.80 (17) | C9—C10—H10A | 109.00 |
| C8—N1—C14 | 113.1 (3) | H10A—C10—H10B | 108.00 |
| C7—N1—C8 | 111.2 (3) | C10—C11—H11B | 109.00 |
| C7—N1—C14 | 111.6 (3) | C10—C11—H11A | 109.00 |
| C7—N1—H1 | 103 (3) | H11A—C11—H11B | 108.00 |
| C8—N1—H1 | 107 (3) | C12—C11—H11A | 109.00 |
| C14—N1—H1 | 110 (3) | C12—C11—H11B | 109.00 |
| C5—N2—H2A | 123 (2) | C11—C12—H12A | 109.00 |
| H2A—N2—H2B | 114 (3) | C13—C12—H12A | 109.00 |
| C5—N2—H2B | 122 (2) | C13—C12—H12B | 109.00 |
| C2—C1—C6 | 120.6 (3) | C11—C12—H12B | 109.00 |
| Br2—C2—C3 | 117.9 (3) | H12A—C12—H12B | 108.00 |
| Br2—C2—C1 | 121.4 (3) | C12—C13—H13A | 110.00 |
| C1—C2—C3 | 120.7 (4) | C12—C13—H13B | 109.00 |
| C2—C3—C4 | 118.1 (4) | C8—C13—H13A | 110.00 |
| Br1—C4—C3 | 117.4 (3) | C8—C13—H13B | 109.00 |
| C3—C4—C5 | 124.2 (3) | H13A—C13—H13B | 108.00 |
| Br1—C4—C5 | 118.4 (3) | H14B—C14—H14C | 109.00 |
| N2—C5—C4 | 121.5 (3) | N1—C14—H14A | 110.00 |
| N2—C5—C6 | 122.7 (4) | N1—C14—H14B | 109.00 |
| C4—C5—C6 | 115.7 (3) | N1—C14—H14C | 109.00 |
| C1—C6—C5 | 120.6 (4) | H14A—C14—H14B | 109.00 |
| C5—C6—C7 | 121.0 (3) | H14A—C14—H14C | 109.00 |
| C1—C6—C7 | 118.3 (3) | S1—C15—C16 | 120.1 (3) |
| N1—C7—C6 | 114.8 (3) | S1—C15—C20 | 120.5 (3) |
| C9—C8—C13 | 111.8 (3) | C16—C15—C20 | 119.4 (4) |
| N1—C8—C9 | 111.0 (3) | C15—C16—C17 | 120.6 (4) |
| N1—C8—C13 | 110.3 (3) | C16—C17—C18 | 121.1 (4) |
| C8—C9—C10 | 110.6 (4) | C17—C18—C19 | 117.7 (4) |
| C9—C10—C11 | 113.3 (4) | C17—C18—C21 | 121.5 (4) |
| C10—C11—C12 | 111.2 (4) | C19—C18—C21 | 120.7 (5) |
| C11—C12—C13 | 112.1 (4) | C18—C19—C20 | 121.7 (4) |
| C8—C13—C12 | 110.6 (4) | C15—C20—C19 | 119.5 (4) |

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| C6—C1—H1A | 120.00 | C15—C16—H16 | 120.00 |
| C2—C1—H1A | 120.00 | C17—C16—H16 | 120.00 |
| C2—C3—H3 | 121.00 | C16—C17—H17 | 119.00 |
| C4—C3—H3 | 121.00 | C18—C17—H17 | 120.00 |
| N1—C7—H7A | 109.00 | C18—C19—H19 | 119.00 |
| C6—C7—H7B | 109.00 | C20—C19—H19 | 119.00 |
| N1—C7—H7B | 109.00 | C15—C20—H20 | 120.00 |
| C6—C7—H7A | 109.00 | C19—C20—H20 | 120.00 |
| H7A—C7—H7B | 108.00 | C18—C21—H21A | 109.00 |
| C13—C8—H8 | 108.00 | C18—C21—H21B | 109.00 |
| N1—C8—H8 | 108.00 | C18—C21—H21C | 110.00 |
| C9—C8—H8 | 108.00 | H21A—C21—H21B | 109.00 |
| H9A—C9—H9B | 108.00 | H21A—C21—H21C | 110.00 |
| C8—C9—H9B | 110.00 | H21B—C21—H21C | 110.00 |
| C10—C9—H9A | 110.00 | | |
| O2—S1—C15—C16 | -47.4 (3) | N2—C5—C6—C1 | -178.4 (4) |
| O1—S1—C15—C16 | 70.1 (3) | N2—C5—C6—C7 | 4.7 (7) |
| O1—S1—C15—C20 | -109.6 (3) | C4—C5—C6—C7 | -172.6 (4) |
| O3—S1—C15—C20 | 12.1 (4) | C4—C5—C6—C1 | 4.3 (6) |
| O2—S1—C15—C20 | 133.0 (3) | C1—C6—C7—N1 | 75.0 (5) |
| O3—S1—C15—C16 | -168.2 (3) | C5—C6—C7—N1 | -108.1 (4) |
| C8—N1—C7—C6 | 174.7 (3) | N1—C8—C9—C10 | -177.7 (3) |
| C14—N1—C8—C13 | -176.8 (3) | C9—C8—C13—C12 | 55.3 (5) |
| C7—N1—C8—C9 | -178.8 (3) | C13—C8—C9—C10 | -54.1 (5) |
| C7—N1—C8—C13 | 56.7 (4) | N1—C8—C13—C12 | 179.3 (3) |
| C14—N1—C7—C6 | 47.3 (4) | C8—C9—C10—C11 | 53.7 (5) |
| C14—N1—C8—C9 | -52.3 (4) | C9—C10—C11—C12 | -53.7 (6) |
| C2—C1—C6—C5 | -1.9 (6) | C10—C11—C12—C13 | 54.2 (5) |
| C2—C1—C6—C7 | 175.1 (4) | C11—C12—C13—C8 | -55.1 (5) |
| C6—C1—C2—C3 | -1.0 (6) | S1—C15—C16—C17 | 179.8 (3) |
| C6—C1—C2—Br2 | 178.1 (3) | C20—C15—C16—C17 | -0.6 (6) |
| Br2—C2—C3—C4 | -178.0 (3) | S1—C15—C20—C19 | -178.7 (3) |
| C1—C2—C3—C4 | 1.1 (6) | C16—C15—C20—C19 | 1.7 (5) |
| C2—C3—C4—Br1 | -179.2 (3) | C15—C16—C17—C18 | -0.2 (7) |
| C2—C3—C4—C5 | 1.7 (6) | C16—C17—C18—C19 | 0.0 (7) |
| C3—C4—C5—N2 | 178.4 (4) | C16—C17—C18—C21 | 178.6 (4) |
| Br1—C4—C5—N2 | -0.7 (6) | C17—C18—C19—C20 | 1.1 (6) |
| Br1—C4—C5—C6 | 176.6 (3) | C21—C18—C19—C20 | -177.6 (4) |
| C3—C4—C5—C6 | -4.3 (6) | C18—C19—C20—C15 | -1.9 (6) |

Symmetry codes: (i) $x, -y+1/2, z-1/2$; (ii) $x, -y+1/2, z+1/2$; (iii) $x, -y+3/2, z+1/2$; (iv) $-x+1, -y+1, -z$; (v) $-x+1, y+1/2, -z+1/2$; (vi) $x, y+1, z$; (vii) $-x+1, -y+2, -z$; (viii) $-x+1, y-1/2, -z+1/2$; (ix) $x, y-1, z$; (x) $-x, -y+1, -z$; (xi) $x, -y+3/2, z-1/2$; (xii) $-x, -y+2, -z$.

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

| $D-H\cdots A$ | $D-H$ | $H\cdots A$ | $D\cdots A$ | $D-H\cdots A$ |
|-----------------------------------|----------|-------------|-------------|---------------|
| N1—H1 \cdots O2 ^{viii} | 0.85 (3) | 1.93 (4) | 2.756 (4) | 161 (4) |
| N2—H2A \cdots O1 ^{iv} | 0.85 (3) | 2.11 (3) | 2.926 (4) | 162 (4) |

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|--------------|----------|----------|-----------|---------|
| N2—H2B···Br1 | 0.84 (2) | 2.67 (3) | 3.068 (3) | 111 (2) |
| C7—H7A···O2 | 0.9700 | 2.4700 | 3.257 (5) | 138.00 |
| C20—H20···O3 | 0.9300 | 2.5300 | 2.905 (5) | 104.00 |

Symmetry codes: (iv) $-x+1, -y+1, -z$; (viii) $-x+1, y-1/2, -z+1/2$.