| C2-C3 | $1.50(2)$ | C7-C8 | $1.36(2)$ |
| :--- | :---: | :--- | :--- |
| C3-C9 | $1.50(2)$ | C8-C9 | $1.37(2)$ |
| C3-C11 | $1.53(2)$ | C12-C13 | $1.53(2)$ |
| C3-C10 | $1.56(2)$ | C12-C15 | $1.55(3)$ |
| C4-C5 | $1.41(2)$ | C12-C14 | $1.56(3)$ |
| C4-C9 | $1.43(2)$ |  |  |
| C2-N1-O | $129(1)$ | C6-C5-C4 | $123(1)$ |
| C2-N1-C8 | $110(1)$ | C5-C6-C7 | $119(1)$ |
| O-N1-C8 | $122(1)$ | C5-C6-C12 | $121(1)$ |
| N1-C2-C3 | $114(1)$ | C7-C6-C12 | $120(1)$ |
| C9-C3-C2 | $98(1)$ | C8-C7-C6 | $117(1)$ |
| C9-C3-C11 | $111(1)$ | C7-C8-C9 | $126(1)$ |
| C9-C3-C10 | $113(1)$ | C7-C8-N1 | $127(1)$ |
| C2-C3-C11 | $111(1)$ | C9-C8-N1 | $107(1)$ |
| C2-C3-C10 | $113(1)$ | C8-C9-C4 | $118(1)$ |
| C11-C3-C10 | $110(1)$ | C8-C9-C3 | $111(1)$ |
| C5-C4-C9 | $117(1)$ | C4-C9-C3 | $131(1)$ |

The structure was solved by direct methods (SIMPEL; Schenk \& Hall, 1990) and refined by full-matrix least-squares calculations; anisotropic for non- H atoms and isotropic for H atoms. The H atoms were positioned geometrically and included as riding atoms in the structure-factor calculations. After isotropic refinement an empirical absorption correction (DIFABS; Walker \& Stuart, 1983) was applied. Data collection: CAD-4 Software (Enraf-Nonius, 1989). Cell refinement: CELCON program comparable to Xtal LATCON (Hall \& Stewart, 1990). Data reduction: Xtal ADDREF, SORTRF. Program(s) used to solve structure: Xtal SIMPEL. Program(s) used to refine structure: Xtal CRYLSQ. Molecular graphics: PLUTO (Motherwell \& Clegg, 1978). Software used to prepare material for publication: Xtal BONDLA, CIFIO.

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sits 0.506 (4) $\AA$ out of plane from the other four atoms; the ring displacement asymmetry parameter $\Delta C_{s}(\mathrm{C} 3 \mathrm{a})$ is $0.8^{\circ}$. The title compound forms the ring skeleton in the crystal structures of a number of bromo substituted desmotroposantonin compounds: in 2-bromo-(-)- $\alpha-$ desmotroposantonin (White \& Sim, 1977a), 2,7-dibromo-$(-)-\beta$-desmotroposantonin (White \& Sim, 1977b) and 2-bromo-(-)- $\beta$-desmotroposantonin (McPhail, Rimmer, Robertson \& Sim, 1967) the $B / C$ ring junction is always found to be cis.


Fig. 1. PLUTO (Motherwell \& Clegg, 1978) drawing showing the numbering system of the title compound. The H atoms are shown but not labelled.

## Experimental

Crystal data
$\mathrm{C}_{12} \mathrm{H}_{12} \mathrm{O}_{2}$
$M_{r}=188.23$
Monoclinic
$P 2_{1} / c$
$a=9.433$ (3) $\AA$
$b=12.674$ (3) $\AA$
$c=8.276$ (3) $\AA$
$\beta=101.99$ (3) ${ }^{\circ}$
$V=967.8(5) \AA^{3}$
$Z=4$
$D_{x}=1.292 \mathrm{Mg} \mathrm{m}^{-3}$
Mo $K \alpha$ radiation
$\lambda=0.71069 \AA$
Cell parameters from 23 reflections
$\theta=20.0-21.5^{\circ}$
$\mu=0.081 \mathrm{~mm}^{-1}$
$T=293 \mathrm{~K}$
Block
$0.43 \times 0.43 \times 0.25 \mathrm{~mm}$
Colourless
Data collection
Enraf-Nonius CAD-4 diffractometer
$\theta / 2 \theta$ scans
Absorption correction: None
2963 measured reflections
2927 independent reflections
1514 observed reflections $[I>2.5 \sigma(I)]$

## Refinement

Refinement on $F$
Extinction correction: Zachariasen (1967)
Final $R=0.049$
$w R=0.072$
$S=0.169$
1514 reflections
176 parameters
$w=1 /\left(2.01+F_{o}\right.$ $\left.+0.0091 F_{o}^{2}\right)$
$(\Delta / \sigma)_{\text {max }}=0.092$
$\Delta \rho_{\text {max }}=0.21 \mathrm{e}_{\AA^{-3}}$
$\Delta \rho_{\text {min }}=-0.183 \mathrm{e}^{-3}$

Table 1. Fractional atomic coordinates and equivalent isotropic thermal parameters $\left(\AA^{2}\right)$

| $U_{\mathrm{eq}}=\frac{1}{3} \Sigma_{i} \Sigma_{j} U_{i j} a_{i}^{*} a_{j}^{*} \mathbf{a}_{i} \cdot \mathbf{a}_{j}$. |  |  |  |  |
| :---: | :---: | :---: | :---: | :---: |
|  | $x$ | $y$ | $z$ | $U_{\text {eq }}$ |
| Ol | 0.7167 (2) | 0.7584 (1) | 0.9241 (2) | 0.0578 (9) |
| C2 | 0.8589 (3) | 0.7507 (3) | 0.9235 (3) | 0.062 (1) |
| C3 | 0.9113 (3) | 0.6417 (3) | 0.9729 (5) | 0.075 (2) |
| C3a | 0.7739 (3) | 0.5759 (2) | 0.9440 (3) | 0.058 (1) |
| C4 | 0.7326 (3) | 0.5316 (2) | 0.7696 (4) | 0.064 (2) |
| C5 | 0.5835 (3) | 0.4818 (2) | 0.7396 (4) | 0.064 (1) |
| C5a | 0.4711 (3) | 0.5600 (2) | 0.7699 (3) | 0.049 (1) |
| C6 | 0.3262 (3) | 0.5519 (2) | 0.6870 (3) | 0.064 (2) |
| C7 | 0.2251 (3) | 0.6234 (3) | 0.7122 (4) | 0.070 (2) |
| C8 | 0.2629 (3) | 0.7056 (2) | 0.8223 (4) | 0.066 (2) |
| C9 | 0.4045 (3) | 0.7140 (2) | 0.9082 (3) | 0.052 (1) |
| C9a | 0.5095 (2) | 0.6429 (2) | 0.8813 (3) | 0.043 (1) |
| C9b | 0.6627 (3) | 0.6572 (2) | 0.9750 (3) | 0.049 (1) |
| O2 | 0.9250 (2) | 0.8254 (2) | 0.8859 (3) | 0.087 (1) |

Table 2. Geometric parameters ( $\left(\AA,{ }^{\circ}\right.$ )

| $\mathrm{Ol}-\mathrm{C} 2$ | 1.345 (3) | C5-C5a | 1.510 (4) |
| :---: | :---: | :---: | :---: |
| O1-C9b | 1.474 (3) | C5a-C6 | 1.400 (4) |
| C2-C3 | 1.495 (5) | C5a-C9a | 1.395 (3) |
| C2-02 | 1.209 (4) | C6-C7 | 1.364 (5) |
| C3-C3a | 1.518 (4) | C7-C8 | 1.381 (4) |
| C3a-C4 | 1.523 (4) | C8-C9 | 1.380 (4) |
| C3a-C9b | 1.530 (4) | C9-C9a | 1.390 (3) |
| C4-C5 | 1.514 (4) | C9a-C9b | 1.502 (3) |
| C2-O1-C9b | 110.0 (2) | C6-C5a-C9a | 118.2 (2) |
| $\mathrm{O} 1-\mathrm{C} 2-\mathrm{C} 3$ | 109.8 (2) | C5a-C6-C7 | 121.3 (3) |
| $\mathrm{O} 1-\mathrm{C} 2-\mathrm{O} 2$ | 120.8 (3) | C6-C7-C8 | 120.6 (3) |
| C3-C2-02 | 129.4 (3) | C7-C8-C9 | 119.2 (3) |
| C2-C3-C3a | 104.0 (2) | C8-C9-C9a | 120.9 (2) |
| C3-C3a-C4 | 113.2 (3) | C5a-C9a-C9 | 119.9 (2) |
| C3-C3a-C9b | 101.3 (2) | C5a-C9a-C9b | 121.4 (2) |
| C4-C3a--C9b | 111.0 (2) | C9-C9a-C9b | 118.7 (2) |
| C3a-C4-C5 | 110.8 (3) | O1-C9b-C3a | 104.4 (2) |
| C4-C5-C5a | 111.0 (2) | O1-C9b-C9a | 108.1 (2) |
| C5-C5a-C6 | 121.2 (2) | C3a-C9b-C9a | 116.9 (2) |
| C5-C5a-C9a | 120.6 (2) |  |  |

The structure was solved by direct methods (SIMPEL; Schenk \& Hall, 1990) and refined by full-matrix least-squares calculations; anisotropic for non- H atoms and isotropic for H atoms. The H atoms were positioned geometrically and refined in the structure-factor calculations. Data collection: CAD-4 Software (Enraf-Nonius, 1989). Cell refinement: CELCON comparable to Xtal LATCON (Hall \& Stewart, 1990). Data reduction: Xtal ADDREF, SORTRF. Program(s) used to solve structure: Xtal SIMPEL. Program(s) used to refine structure: Xtal CRYLSQ. Molecular graphics: PLUTO (Motherwell \& Clegg, 1978). Software used to prepare material for publication: Xtal BONDLA, CIFIO.

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# Structure of $N, N^{\prime}$-Dimethylpiperazine-2,3dithione: Space Group Correction 

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#### Abstract

The crystal structure of $N, N^{\prime}$-dimethylpiperazine-2,3dithione, $\mathrm{C}_{6} \mathrm{H}_{10} \mathrm{~N}_{2} \mathrm{~S}_{2}$, has been described and refined in space group $A a$ [Servaas, Stufkens, Oskam, Vernooijs, Baerends, De Ridder \& Stam (1989). Inorg. Chem. 28,


[^0]4104-4113]. It is described properly in space group $A 2 / a$. Revised coordinates and bond lengths and angles are given.

## Comment

The structure of the title compound was reported in space group $A a$ (Servaas et al., 1989). The six-membered ring of the molecule was found to adopt a half-chair conformation with the twofold axis running through the midpoints of the $\mathrm{C}-\mathrm{C}$ bonds, parallel to the $b$ axis. Furthermore, the two $\mathrm{N}-\mathrm{C}$ (methyl) bond lengths were very dissimilar [1.439 (5) and 1.489 (5) A]]; there is no reason for them to be unequal. This unusual spread of distances undoubtedly resulted from the refinement of a centrosymmetric structure in a non-centrosymmetric space group (Ermer \& Dunitz, 1970; Schomaker \& Marsh, 1979). Therefore it seemed likely that the structure should properly be described in space group $A 2 / a$.

Starting coordinates were from Table 3 of Servaas et al. (1989) and averaged in accordance with $A 2 / a$. Fullmatrix minimization of the quantity $\Sigma w\left(F_{o}-F_{c}\right)^{2}$ with $w=1 /\left[\sigma^{2}(F)+0.0035 F^{2}\right]$ was performed. Refinement was anisotropic for the non- H atoms and isotropic for the H atoms. An isotropic extinction coefficient was included in the parameters [Zachariasen, final value $3.4(5) \times 10^{-5}$ ]. Anomalous dispersion was taken into account but no correction for absorption was applied. The H -atom positions were calculated initially on the basis of standard geometry and refined. A final difference Fourier map revealed residual electron density within the range $-0.22-0.30$ e $\AA^{-3}$.
The values obtained for parameters, bond lengths and angles are close to the averages of pairs of equivalent values obtained in the earlier $A a$ refinement. The two $\mathrm{N}-\mathrm{C}$ (methyl) bond lengths are now equal by symmetry at 1.473 (3) $\AA$. The general description of the structure remains unchanged.


Fig. 1. A PLUTO (Motherwell \& Clegg, 1978) drawing of $\mathrm{C}_{6} \mathrm{H}_{10} \mathrm{~N}_{2} \mathrm{~S}_{2}$ showing the labelling of the independent non- H atoms. The view is down the twofold axis, with the crystallographic $2_{1}$ axis vertical.


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