These ring conformations are consistent with the *trans-syn-trans-anti-trans* ring junctions (Boeyens, Bull & Van Rooyen, 1980). For related Westphalen compounds chemical-reactivity studies were interpreted in terms of a half-chair conformation for ring B whilst NMR studies have shown this ring to be relatively flexible (Glotter *et al.*, 1975).

The 17β -side chain exists in the extended conformation which is the most common conformation in cholestane derivatives (Duax, Griffin, Rohrer & Weeks, 1980). All intermolecular contacts between non-H atoms are >3.2 Å.

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Structure of 1,4-Dinitrosopiperazine, $C_4H_8N_4O_2$

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Abstract. $M_r = 144 \cdot 13$, orthorhombic, *Pbca*, $a = 9 \cdot 495$ (4), $b = 6 \cdot 570$ (3), $c = 10 \cdot 521$ (7) Å, $V = 656 \cdot 3$ (9) Å³, Z = 4, $D_x = 1 \cdot 459$, $D_m = 1 \cdot 427$ Mg m⁻³, $\lambda = 0 \cdot 7107$ Å, $\mu = 0 \cdot 0877$ mm⁻¹, F(000) = 304, T = 293 K, final $R = 0 \cdot 089$ for 959 independent reflections. The molecule is centrosymmetric, the piperazine ring having a chair form. The two nitroso groups are in the *anti* form. The N–O bond length is 1 \cdot 202 (4) Å, and the N–N–O angle 113 \cdot 4 (3)°. The molecules are packed together to form infinite ribbons along the [110] direction.

Introduction. As the piperazine system offers a rich selection of conformational questions, molecular shapes of this system have been investigated by NMR, dipole moment, and electron diffraction. Since piperazine derivatives are of interest due to their pharmacological activity and natural occurrence, we have studied the crystal structures of these derivatives by the X-ray method under a current program of this laboratory (Okamoto, Sekido, Itoh, Noguchi & Hirokawa, 1979;

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Okamoto, Sekido, Ono, Noguchi & Hirokawa, 1982). 1,4-Dinitrosopiperazine (DNP) was chosen because it possesses several modes of conformational flexibility and it is a potent carcinogen in various animals. Lambert, Gosnell, Bailey & Henkin (1969) reported that the centrosymmetric *anti* form of DNP appeared to be present exclusively in the solid state, but a clear choice between a chair form and a boat form was not provided. The present work was undertaken to obtain the unambiguous structure of DNP.

Experimental. Colorless rectangular prismatic crystals (m.p. 430 K) obtained by slow evaporation from an ethanol solution. D_m measured by flotation in a mixture of toluene and tetrachloromethane. Crystal $0.5 \times 0.5 \times 0.6$ mm. Rigaku AFC-III four-circle diffractometer, Mo $K\alpha$ radiation monochromatized by a graphite plate. X-ray intensity data measured by $\omega-2\theta$ scan, scan speed 4° min⁻¹ in 2θ . Cell dimensions determined by least-squares method RSLC-3 (UNICS, 1967) using 13 reflections with $25 \cdot 0^{\circ} < \theta < 29 \cdot 9^{\circ}$. Systematic absences h = 2n + 1 in hk0, k = 2n + 1 in 0kl, and l = 2n + 1 in h0l. 959 independent reflections

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measured; $0 \le h \le 13$, $0 \le k \le 9$, $0 \le l \le 14$; standard reflections 400, 020, 006, intensity variation 3.0%; $(\sin\theta_{\rm max})/\lambda = 0.7030 \text{ Å}^{-1}, 2\theta_{\rm max} = 59.99^{\circ}; 644 \text{ unique}$ reflections, $R_{\text{int}} = 0.067$; 315 unobserved reflections with $I < 3\sigma(I)$. Data corrected for Lorentz and polarization factors but not for absorption. Structure solved by a three-dimensional Patterson synthesis, and refined by block-diagonal least squares using HBLS-5 (Ashida, 1973) and RSSFR-5 (UNICS, 1967). H-atom positions located on a difference Fourier map. Final cycles of refinement with anisotropic temperature factors for C, N, and O, and isotropic for H atoms. $R = 0.089, wR = 0.090, S = 1.62. \sum w(|F_o| - k|F_c|)^2$ minimized; weighting scheme $w = 1 \cdot 0/[\sigma(F_o)]^2$ for $F_o \neq 0$ and $w = 1 \cdot 0$ for $F_o = 0$. In final refinement cycle $(\Delta/\sigma)_{max} = 0.09$ and max. peak height in final difference map 0.3 e Å⁻³. Scattering factors from *International* Tables for X-ray Crystallography (1974). All calculations performed on a HITAC M-200 (H) computer in the National Defense Academy.

Discussion. The numbering scheme and the thermal ellipsoids of the non-hydrogen atoms are illustrated in Fig. 1. The atomic parameters with their standard deviations are shown in Table 1.* The bond distances and angles are listed in Table 2. Since the molecule is centrosymmetric, the piperazine ring has a chair form (Armarego, 1977), and the nitroso groups are in the anti form. The N-C bond length in the piperazine ring, 1.450 (5) Å (mean), is in good agreement with 1.463 Å in 2,5-dimethylpiperazine (DP) (Okamoto, Sekido, Ono, Noguchi & Hirokawa, 1982), 1.459 Å in trans-1,4-dibenzoyl-2,5-dimethylpiperazine (trans-DDP) (Okamoto, Sekido, Itoh, Noguchi & Hirokawa, 1979), 1.467 Å in piperazine (PZ) (Yokozeki & Kuchitsu, 1971), 1.464 Å in cis-1,4-dibenzoyl-2,5-dimethylpiperazine (cis-DDP) (Sakurai, Nakamaru, Tsuboyama

^{*} Lists of structure factors, anisotropic thermal parameters, H-atom parameters, least-squares planes, and torsion angles have been deposited with the British Library Lending Division as Supplementary Publication No. SUP 39978 (8 pp.). Copies may be obtained through The Executive Secretary, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.



Fig. 1. Numbering scheme and the thermal ellipsoids of the non-hydrogen atoms drawn by *ORTEP* (Johnson, 1965). The thermal-vibration ellipsoids are set to 50% probability.

Table 1. Fractional atomic coordinates $(\times 10^4)$ and equivalent isotropic thermal parameters for non-H atoms

The estimated standard deviations given in parentheses refer to the last decimal position. $B_{eq} = \frac{1}{2} \sum_{i} \sum_{j} B_{ij} a_i^* a_j^* \mathbf{a}_i \cdot \mathbf{a}_j$.

	x	у	z	$B_{eq}(\dot{A}^2)$
N(1)†	1195 (3)	612 (3)	629 (3)	4.5
C(2)	-140(4)	710 (4)	1292 (3)	5.2
C(3)	-1148(3)		743 (3)	5.1
N(7)	2337 (3)	422 (3)	1361 (3)	5.8
O(8)	3422 (3)	422 (3)	772 (3)	7.3

 † N(1), C(2), C(3), N(7), and O(8) are related by a center of symmetry to N(4), C(5), C(6), N(9), and O(10), respectively.

 Table 2. Intramolecular distances (Å) and angles (°)

 between non-H atoms with e.s.d.'s in parentheses

N(1)-C(2)	1.448 (5)	C(2) - C(3)	1.515 (5)
N(1)C(6)	1.451 (5)	N(7)-O(8)	1.202 (4)
N(1)-N(7)	1.336 (4)		
N(1)···C(3)	2.425 (5)	C(2)····N(7)	2-361 (5)
$N(1) \cdots N(4)$	2.746 (4)	$C(3)\cdots C(5)$	2.467 (5)
$N(1)\cdots C(5)$	2.416 (5)	$C(6) \cdots N(7)$	2.500(5)
N(1)···O(8)	2.124 (4)	C(6)····O(8)	2.697 (5)
N(1)-C(2)-C(3)	109.8 (3)	C(2)-N(1)-N(7)	115-9 (3)
C(2)-C(3)-N(4)	109.1 (3)	C(6)-N(1)-N(7)	127.4 (3)
C(2)-N(1)-C(6)	116.6 (3)	N(1)-N(7)-O(8)	113-4 (3)

& Tsuboyama, 1977), and 1.463 Å in 1.4-bis(1.2dimethyl-1-propenyl)piperazine (Van Meerssche, Germain, Declercq & Colens, 1979). The bond distance of 1.515 (5) Å for C–C is shorter than the $C(sp^3)$ – $C(sp^3)$ covalent bond length. But this is comparable to 1.521 Å in DP, 1.517 Å in trans-DDP, 1.535 Å in cis-DDP, and 1.521 Å in N.N'-dimethylpiperazine (Davis & Hassel, 1963). The N····N distance of 2.746 (4) Å is slightly shorter than 2.806 Å in trans-DDP, and 2.820 Å in trans-DP. The bond angles of $C(sp^3)$ atoms in the piperazine ring are approximately tetrahedral and close to the angles of PZ. The value of 116.6 (3)° for C(2)–N(1)–C(6) agrees with 115.7° in cis-DDP, and is larger than 113.8° in trans-DDP, and 112.6° in piperidine (Blackburne, Duke, Jones, Katritzky & Record, 1973).

The ring torsion angles are $-51 \cdot 1$ (4), $55 \cdot 0$ (4), and $-55 \cdot 4$ (4)° for N(1)-C(2)-C(3)-N(4), C(2)-C(3)-N(4)-C(5), and C(3)-N(4)-C(5)-C(6), respectively. The C(2), C(6), N(1), N(7), and O(8) atoms are coplanar within 0.023 Å. The dihedral angle between the mean plane (1) formed by C(2), C(3), C(5) and C(6) and the mean plane (2) composed of N(1), C(2), C(6), N(7), and O(8) is 48.4 (2)°, which is comparable to those found in *trans*-DP (about 55°) and *trans*-DDP (about 53°).

The value of 113.4 (3)° for N(1)–N(7)–O(8) agrees with 113.6° for N–N–O in dimethylnitrosamine (MNA) (Rademacher & Stolevik, 1969). The N–O bond distance of 1.202 (4) Å is comparable to that



Fig. 2. The molecular packing in the unit cell viewed down the b axis. Symmetry codes: (i) x, y, z; (ii) 0.5 + x, 0.5 - y, -z.

found in MNA (1.234 Å). The N–N bond length of 1.336 (4) Å is in good agreement with that of 1.344 Å in MNA, and is significantly shorter than the N–N single bond (Liminga & Olovsson, 1964). This shows the N–N double-bond character. Thus the C_2NNO group results in hindered rotation about the N–N bond (Cooney, Brownstein & ApSimon, 1974).

The molecular packing viewed down the *b* axis is shown in Fig. 2. The closest non-bonded approach of $3 \cdot 185$ (4) Å is between N(1ⁱ) and O(10ⁱⁱ). The secondshortest contact of $3 \cdot 300$ (6) Å is between N(7ⁱ) and N(9ⁱⁱ). The triangular plane formed by N(1ⁱ), N(7ⁱ), and O(8ⁱ) is approximately parallel to the neighboring triangular plane formed by N(4ⁱⁱ), N(9ⁱⁱ), and O(10ⁱⁱ), the dihedral angle being $10 \cdot 2$ (1)°. No other short contacts can be seen in the structure. The molecules are packed together to form infinite ribbons along the [110] direction, and the ribbons are held together in the crystal by van der Waals forces.

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Structure of 2-[(Hydroxyimino)(phenylthio)methyl]-1-methylpyridinium Chloride, $C_{13}H_{13}N_2OS^+.Cl^-$

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Abstract. $M_r = 280.78$, $P2_1/c$, a = 9.30 (1), b = 11.433 (7), c = 14.02 (2) Å, $\beta = 114.8$ (1)°, V = 1353 (3) Å³, Z = 4, $D_x = 1.378$ Mg m⁻³, Mo Ka, $\lambda = 0.71069$ Å, $\mu = 0.373$ mm⁻¹, F(000) = 584, room

temperature, R = 0.037, 2513 unique reflections. The oxime moiety of the molecule is in the (E) configuration. The molecule has a folded conformation, which may be responsible for its inability to reactivate

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