355 Å<sup>3</sup> at 100% RH using areas per lipid of 59.3 and 65.5 Å<sup>2</sup>, respectively. Terminal methyl-group volumes increased with increased levels of humidity from 51 (1) at 70% RH to 56 (2) Å<sup>3</sup> at 100% RH.

In conclusion, we have clearly demonstrated, using one-dimensional absolute electron-density profiles calculated by means of direct-phasing methods, quantitative structural differences between two phases of DLPC. In addition to the changes observed by other researchers when comparing gel and liquid-crystalline bilayers, our profiles clearly show the phosphorylcholine headgroup reorienting when the DLPC bilayers undergo a transition from the 3D- $L_{\delta}$  phase to the  $L_{\alpha}$  phase. Although a 2D- $L_{\delta}$ phase has been previously observed (Tardieu *et al.*, 1973), we have reported and quantified the physical characteristics for the first time of a DLPC 3D- $L_{\delta}$ phase which seems to differ substantially from the two-dimensional phase.

We are indebted to Dr Douglas L. Dorset for discussions and his critical reading of the manuscript. We would also like to thank D. S. C. Yang for use of his rotating-anode two-dimensional detector. JK is the recipient of a Natural Sciences and Engineering Research Council of Canada postdoctoral fellowship (NSERC). This research was supported by grants from NSERC.

### References

- ABRAHAMSSON, S., DAHLEN, B., LÖFGREN, H. & PASCHER, I. (1978). Prog. Chem. Fats Other Lipids, 16, 125–143.
- CHAPMAN, D., URBINA, J. & KEOUGH, K. M. (1974). J. Biol. Chem. 249, 2512-2521.

DORSET, D. L. (1990). Biophys. J. 58, 1077-1087.

- DORSET, D. L. (1991a). Biophys. J. 60, 1356-1365.
- DORSET, D. L. (1991b). Biophys. J. 60, 1366-1373.
- DORSET, D. L., BECKMANN, E. & ZEMLIN, F. (1990). Proc. Natl Acad. Sci. USA, 87, 7570-7573.
- DOUCET, J., LEVELUT, A. M. & LAMBERT, M. (1983). Acta Cryst. B39, 724-731.
- GENNIS, R. B. (1989). Biomembranes: Molecular Structure and Function. New York: Springer-Verlag.
- HAUPTMAN, H. & KARLE, J. (1953). Solution of the Phase Problem I. The Centrosymmetric Crystal. ACA Monograph No. 3. Ann Arbor, MI: Edwards Brothers.
- JENDRASIAK, G. L. & HASTY, J. H. (1974). Biochim. Biophys. Acta, 337, 79-91.
- KATSARAS, J. & STINSON, R. H. (1990). Biophys. J. 57, 649-655.
- KATSARAS, J., STINSON, R. H., DAVIS, J. H. & KENDALL, E. J. (1991). *Biophys. J.* **59**, 645–653.
- KATSARAS, J., YANG, D. S.-C. & EPAND, R. M. (1992). *Biophys. J.* 63, 1179–1175.
- LUZZATI, V., GULIK-KRZYWICKI, T. & TARDIEU, A. (1968). Nature (London), 218, 1031-1034.
- MCINTOSH, T. J. (1978). Biochim. Biophys. Acta, 513, 43-58.
- MORROW, M. R. & DAVIS, J. H. (1987). Biochim. Biophys. Acta, 904, 61-70.
- NAGLE, J. F. & WIENER, M. C. (1988). Biochim. Biophys. Acta, 942, 1-10.
- PEARSON, R. H. & PASCHER, I. (1979). Nature (London), 281, 499-501.
- RUOCCO, M. J. & SHIPLEY, G. G. (1982). Biochim. Biophys. Acta, 691, 309-320.
- SAYRE, D. (1952). Acta Cryst. 5, 843.
- SHANNON, C. E. (1949). Proc. Inst. Radio Eng. NY, 37, 10-21.
- SHERWOOD, D. H. (1976). Crystals, X-rays and Proteins. London: Longman.
- SMALL, D. M. (1984). J. Lipid Res. 25, 1490-1500.
- TANFORD, C. (1973). The Hydrophobic Effect: Formation of Micelles and Biological Membranes. New York: John Wiley. TARDIEU, A., LUZZATI, V. & REMAN, F. C. (1973). J. Mol. Biol.
- **75**, 711–733. **76**, 711–733.
- TORBET, J. & WILKINS, M. H. F. (1976). J. Theor. Biol. 62, 447–458.
- WIENER, M. C. & WHITE, S. H. (1992). Biophys. J. 61, 434-447.
- WIENER, M. C., SUTER, R. M. & NAGLE, J. F. (1989). *Biophys. J.* 55, 315-325.

Acta Cryst. (1994). B50, 216-220

# Structure of 1,1,5,5-Tetranitro-[4]peristylane. Structure Solution from Molecular Packing Analysis

BY HERMAN L. AMMON, ZUYUE DU AND JAMES R. HOLDEN

Department of Chemistry and Biochemistry, University of Maryland, College Park, MD 20742, USA

AND LEO A. PAQUETTE

Department of Chemistry, The Ohio State University, Columbus, OH 43210, USA

(Received 4 January 1993; accepted 31 August 1993)

## Abstract

The structure of the title compound, decahydro-2,2,5,5-tetranitro-1,6:3,4-dimethanocyclobuta[1,2:3,-

© 1994 International Union of Crystallography Printed in Great Britain – all rights reserved 4]dicyclopentene,  $C_{12}H_{12}N_4O_8$ , was solved with the molecular packing program *MOLPAK*, starting with an *AM*1-geometry-optimized model of an isolated molecule. The 20 best predicted crystal structures

from the *MOLPAK* procedure were subjected to lattice energy refinement with the *WMIN* program. A structure-factor calculation with the top *MOLPAK/WMIN* derived structure and the 197  $F_o$ data to  $\sin\theta/\lambda = 0.324$  Å<sup>-1</sup> ( $\theta = 30^\circ$ ) gave an *R* value of 0.29. Final R = 0.057, wR = 0.067 for 730 reflections with  $I > 3\sigma(I)$ . There are no unusual intermolecular distances and crystal packing is normal.

## Introduction

One of the most important properties of an energetic material is the solid-state density. The performance characteristics of propellants and explosives, such as detonation velocity and pressure, are functions of density and at an even more basic level density determines the quantity of material that can be packed into a limited volume. Polynitro-cage compounds, such as the cubanes and homocubanes, are of interest as potential high-energy density materials. The energy contained in the strained ring systems together with their rather compact shapes holds the promise of energetic materials with exceptional performance characteristics.

We have undertaken crystal structure determinations of several of the [4]peristylanes and bishomopentaprismanes to enable detailed crystal-packing studies to be made and to confirm, in some cases, reported crystal densities. In this paper we report the structure of (1). The solution was achieved by an *ab initio* molecular packing procedure that has been under development in our laboratory over the last several years; this is described in some detail.



## Experimental

Experimental details are given in Table 1. Data were collected with Enraf–Nonius diffractometer software (Enraf–Nonius, 1989). The  $\theta$ -scan width was 1.5(0.8 + 0.14 tan $\theta$ )°, recorded as 96 steps with two outermost 16-step blocks for background determination; the  $\theta$ -scan speed was 2.24–0.82° min<sup>-1</sup>. Refinement was by full-matrix least squares, minimizing  $\Sigma[1/\sigma^2(F_o)(F_o - F_c)^2]$  with anisotropic temperature factors for C, O and N; H-atom positions were obtained

### Table 1. Experimental details

Crystal data	
Chemical formula	C <sub>12</sub> H <sub>12</sub> N <sub>4</sub> O <sub>8</sub>
Aolecular weight	340.3
Crystal system	Monoclinic
Space group	P21
(Å)	7.073 (1)
(Å)	11.624 (1)
(Å)	8.231 (1)
3 (°)	96.79 (1)
$(\dot{A}^3)$	671.7 (8)
	2
adjustion type	Cu Kα
Vavelength (Å)	1.54178
No. of reflections for cell	25
parameters	
range for cell parameters (°)	7.4-25.2
$(mm^{-1})$	1.20
Comperature (K)	293
Printel color	Colorless
Crystal Color	Rod
Crystal rollin	$0.40 \times 0.037 \times 0.033$
rystar size (nun)	1.682
$D_x$ (Mg III )	Supplied by I A Paquette
rystal source	Supplied by E. A. Faquette
Data collection	
Diffractometer	Enraf-Nonius CAD-4
Absorption correction	None
No. of measured reflections	1285
No. of independent reflections	1054
No. of observed reflections	730
boundation criterion	$[l > 3\sigma(l)]$
	0.030 for 78 pairs
Tint D (9)	60.0
max ()	$-7 \rightarrow h \rightarrow 7$
cange of <i>n</i> , <i>k</i> , <i>i</i>	$-11 \rightarrow k \rightarrow 0$
	$0 \rightarrow l \rightarrow 8$
a for a standard and a stand	7
No. of standard reflections	60 min of X-ray exposure
-requency or standard reflections	$-28 \rightarrow 0.7\%$ (-1.8%)
ntensity variation (%)	-2.0-0.7%; (-1.0%)
Refinement	
Refinement on	F
Final <i>R</i> factor	0.057
	0.067
~A	1.7
)	730
refinement	
lo of porameters used	228
Latom refinement	Only H-atom U's refined
Veighting	$w = 1/\sigma^2(F_0)$
	0.09
$\Delta \sigma_{\text{max}}$	0.28
$\Delta \rho_{\text{max}} (c A^{-})$	_0.31
$\Delta \rho_{\rm min}$ (e A <sup>-</sup> )	International Tables for Y ray
source of atomic scattering factors	Crystallography (1974, Vol. IV,
	nn ())-(/)

from the C-atom framework, with individual isotropic temperature factors refined.\*

All crystallographic calculations were performed with the *TEXSAN* (Molecular Structure Corporation, 1989) program system on DEC MicroVAX II, VAXStation II and 3100 computers; structure solution was performed with the molecular packing program *MOLPAK* (Holden, Du & Ammon, 1993). Figs. 1 and 2 were drawn using the *Chem*3D pro-

<sup>\*</sup> Lists of structure factors, anisotropic thermal parameters and H-atom parameters have been deposited with the British Library Document Supply Centre as Supplementary Publication No. SUP 71570 (11 pp.). Copies may be obtained through The Technical Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England. [CIF reference: CR1059]

gram (Cambridge Scientific Computing, 1990) on an Apple MacIntosh IIx workstation and printed on an Apple LaserWriter IINTX. Figs. 3 and 4 were drawn with the ORTEP (Johnson, 1976) subprogram of TEXSAN; the PLOTMD program (Luo, Ammon & Gilliland, 1989) was used to display the drawings on a VAXStation monitor, add labels and prepare print files for a Hewlett-Packard Laser-Jet II printer.

The procedure used to predict possible crystal structures has been previously described in detail (Holden, Du & Ammon, 1993). A brief outline will be given here. Investigations of the molecular coordination environments in a number of triclinicorthorhombic crystal structures with  $Z \le 4$  and one molecule per asymmetric unit have revealed common patterns in the coordination geometries for each of the space groups. The most common coordination for  $P2_1$  is illustrated in Fig. 1. The molecular packing program MOLPAK was written to reconstruct the various geometries, starting with models for the compounds of interest. MOLPAK uses a rigid-body model of the compound (the search probe) to build a number of packing arrangements (possible crystal structures) with the predetermined coordination geometry. All unique orientations of the search probe (with respect to fixed Cartesian axes) were examined and the molecules were fit together (in the appropriate coordination geometry) with а repulsion-only potential function. The best (smallest volume) of the MOLPAK packing arrangements are refined with the WMIN procedure (Busing, 1981).

For this investigation, the search probe structure was constructed with the *Macromodel* (Still, 1990) system and subjected to geometry optimization with the AM1 procedure (Dewar, Zoebisch, Healy & Stewart, 1985; Stewart, 1990). The probe and refined X-ray structures, which are shown in Fig. 2, differ primarily in C—NO<sub>2</sub> conformations. In the former, the pairs of geminal nitro substituents are arranged with the upper and lower nitro O atoms approximately equidistant (the four O…O distances range from 3.37 to 3.48 Å). A similar orientation is observed on the left side of the X-ray structure (O…O = 2.98 and 3.47 Å), whereas the bottom right nitro group is twisted by approximately 90° from this arrangement.

The 20 smallest-volume MOLPAK packing arrangements [a total of 19<sup>3</sup> (6859) were created] were refined with the WMIN procedure (lattice energy minimization), which adjusts the search-probe orientation and position and unit-cell constants. The structure with the lowest lattice energy is generally the best (correct structure), but in a situation where experimental cell constants are available, these data can be used to indicate which of the hypothetical structures is most probably correct. Both the lattice energy and cell-parameter comparison criteria pointed to the same structure in the present case. Portions of the refined X-ray and predicted crystal structures are shown in Fig. 3. The predicted structure shown is based on the AM1-optimized search probe; differences in the C-NO<sub>2</sub> conformations alter the crystal packing somewhat and presumably are the cause of the orientation and position differences in the crystal structures. A WMIN refinement,



Fig. 1. Coordination geometry for space group  $P_{2_1}$ . The threeand four-digit numbers are similar to ORTEP (Johnson, 1976) ADC codes. The central molecule is O(555) and the  $\pm$  b-axis directions are indicated with B(565) and B(545). The first number of the four-digit codes refers to the symmetry operations: 1 = x, y, z; 2 = -x,  $\frac{1}{2} \pm y$ , -z. This coordination is characterized by six I (identity) molecules in the *ac* plane and four A (axis) molecules in planes above and below the *ac* plane. The positions labeled I(1655), I(1455), I(1556) and I(1554) show the *a* and *c* axes.



Fig. 2. Chem3D (Cambridge Scientific Computing, 1990) views of the refined X-ray (a) and AM1-optimized search probe (b) structures.

## Table 2. Comparison of the cell constants from the X-ray and predicted structures\*

	X-ray	$\Delta_1$ (%)	$\Delta_2$ (%)
a (Å)	7.073	3.1	2.1
b (Å)	11.625	0.1	0.5
c (Å)	8.231	-0.3	- 1.6
B(°)	96.79°	-4.7	- 0.5

\* X-ray = experimental cell constants.  $\Delta_1$  and  $\Delta_2$  give per cent differences between the predicted unit-cell parameters (1 and 2 indicate without and with WMIN C-NO<sub>2</sub> optimization). % =  $100 \times (\text{predicted} - \text{X-ray})/\text{X-ray}.$ 

which included optimization of the four C-NO<sub>2</sub> twist angles, yielded a hypothetical structure with an r.m.s. deviation of 0.26 Å from the refined X-ray structure [the deviation is from a least-squares fit of the 24 C. N and O atoms and 8 unit-cell corners (000, 100, 010 etc.) in the two structures]. The cell constant information for the experiment and two predicted structures is listed in Table 2. A structurefactor calculation with the predicted structure (WMIN-optimized C-NO<sub>2</sub> twist) and the 197 data to  $\sin \theta/\lambda = 0.324 \text{ Å}^{-1}$  ( $\theta = 30^{\circ}$ ) gave R = 0.29 and wR = 0.37 (no attempt was made to correct the model coordinates for differences between the experimental and predicted cell constants; this calculation was performed with the unmodified predicted fractional coordinates); several cycles of least-squares



Fig. 3. ORTEP (Johnson, 1976) drawings of the X-ray (a) and predicted (b) crystal structures. The predicted structure shown was obtained from the MOLPAK/WMIN procedure and does not include refinement of the C-NO<sub>2</sub> conformations.

Table 3. Fractional atomic coordinates and equivalent isotropic displacement parameters  $(Å^2)$  with e.s.d.'s in parentheses

$B_{\text{eq}} = (8\pi^2/3)\sum_i\sum_j U_{ij}a_i^*a_j^*\mathbf{a}_i.\mathbf{a}_j.$					
x	у	z	$B_{eq}$		
-0.364 (1)	0.0620	0.670(1)	5.3 (5)		
-0.538 (2)	0.206(1)	0.591(1)	7.6 (6)		
-0.306(1)	-0.041(1)	0.354 (1)	5.3 (5)		
-0.041 (1)	0.034 (1)	0.474 (1)	4.9 (5)		
0.049 (2)	0.452(1)	-0.090(1)	6.6 (6)		
-0.251 (1)	0.454 (1)	-0.154 (1)	7.0 (6)		
-0.060(1)	0.177 (1)	-0.148(1)	4.7 (4)		
0.167 (1)	0.220(1)	0.043 (1)	5.2 (4)		
-0.415 (1)	0.140(1)	0.574 (1)	3.8 (5)		
-0.206 (2)	0.037 (1)	0.416(1)	3.8 (5)		
-0.113 (2)	0.418 (1)	-0.081 (1)	3.4 (5)		
0.003 (1)	0.227 (1)	-0.024(1)	2.9 (4)		
-0.308 (1)	0.155 (1)	0.421 (1)	2.8 (4)		
-0.178 (1)	0.259 (1)	0.454 (1)	2.6 (4)		
-0.016(1)	0.268 (1)	0.343 (1)	2.3 (4)		
-0.066 (1)	0.360(1)	0.211 (1)	2.5 (5)		
-0.129(1)	0.317 (1)	0.042 (1)	2.4 (4)		
-0.337 (1)	0.279 (1)	0.031 (1)	2.0 (4)		
-0.370 (1)	0.162(1)	0.109(1)	2.6 (5)		
-0.449 (1)	0.178(1)	0.272 (1)	2.9 (5)		
-0.317 (2)	0.359(1)	0.401 (1)	2.5 (4)		
-0.246 (1)	0.421 (1)	0.251 (1)	2.4 (4)		
-0.418 (1)	0.368 (1)	0.139(1)	2.5 (4)		
-0.485 (1)	0.310(1)	0.285 (1)	2.6 (4)		
	$B_{eq} = \frac{x}{-0.364 (1)}$ $-0.364 (1)$ $-0.036 (1)$ $-0.041 (1)$ $-0.041 (1)$ $-0.050 (1)$ $-0.251 (1)$ $-0.060 (1)$ $-0.167 (1)$ $-0.206 (2)$ $-0.113 (2)$ $-0.003 (1)$ $-0.308 (1)$ $-0.308 (1)$ $-0.178 (1)$ $-0.066 (1)$ $-0.129 (1)$ $-0.370 (1)$ $-0.370 (1)$ $-0.317 (2)$ $-0.246 (1)$ $-0.418 (1)$ $-0.485 (1)$	$B_{eq} = (8\pi^2/3)\Sigma_i\Sigma_jU_i$ $x y$ $-0.364 (1) 0.0620$ $-0.538 (2) 0.206 (1)$ $-0.306 (1) -0.041 (1)$ $-0.041 (1) 0.043 (1)$ $-0.041 (1) 0.0454 (1)$ $-0.251 (1) 0.452 (1)$ $-0.251 (1) 0.452 (1)$ $-0.415 (1) 0.140 (1)$ $-0.415 (1) 0.140 (1)$ $-0.206 (2) 0.037 (1)$ $-0.113 (2) 0.418 (1)$ $-0.003 (1) 0.227 (1)$ $-0.308 (1) 0.155 (1)$ $-0.016 (1) 0.268 (1)$ $-0.066 (1) 0.360 (1)$ $-0.129 (1) 0.317 (1)$ $-0.377 (1) 0.229 (1)$ $-0.370 (1) 0.162 (1)$ $-0.449 (1) 0.178 (1)$ $-0.246 (1) 0.421 (1)$ $-0.246 (1) 0.421 (1)$ $-0.418 (1) 0.368 (1)$ $-0.485 (1) 0.310 (1)$	$B_{eq} = (8\pi^2/3)\sum_i\sum_j U_{ij}a_i^*a_j^*a_i.a_j.$ $x y z$ $-0.364 (1) 0.0620 0.670 (1)$ $-0.538 (2) 0.206 (1) 0.591 (1)$ $-0.306 (1) -0.041 (1) 0.354 (1)$ $-0.041 (1) 0.034 (1) 0.474 (1)$ $0.049 (2) 0.452 (1) -0.090 (1)$ $-0.251 (1) 0.454 (1) -0.154 (1)$ $-0.060 (1) 0.177 (1) -0.148 (1)$ $0.167 (1) 0.220 (1) 0.043 (1)$ $-0.415 (1) 0.140 (1) 0.574 (1)$ $-0.206 (2) 0.037 (1) 0.416 (1)$ $-0.308 (1) 0.155 (1) 0.421 (1)$ $-0.308 (1) 0.155 (1) 0.421 (1)$ $-0.066 (1) 0.377 (1) 0.416 (1)$ $-0.178 (1) 0.259 (1) 0.434 (1)$ $-0.066 (1) 0.360 (1) 0.211 (1)$ $-0.370 (1) 0.162 (1) 0.031 (1)$ $-0.371 (2) 0.359 (1) 0.401 (1)$ $-0.317 (2) 0.359 (1) 0.401 (1)$ $-0.276 (1) 0.316 (1) 0.272 (1)$ $-0.317 (2) 0.359 (1) 0.401 (1)$ $-0.246 (1) 0.421 (1) 0.272 (1)$ $-0.317 (2) 0.359 (1) 0.401 (1)$ $-0.448 (1) 0.368 (1) 0.139 (1)$ $-0.448 (1) 0.368 (1) 0.139 (1)$ $-0.485 (1) 0.310 (1) 0.285 (1)$		

refinement with individual isotropic temperature factors and the 376 data to  $\sin \theta / \lambda = 0.417 \text{ Å}^{-1}$  ( $\theta =$ 40°) gave R = 0.103 and wR = 0.122. The refinement was completed in the usual way.

#### Discussion

An ORTEP (Johnson, 1976) drawing is shown in Fig. 4, fractional atomic coordinates are given in Table 3, and bond lengths, bond angles and selected torsion angles are given in Table 4. There is substantial similarity with the structures of other [4]peristylanes [(II)-(IX)], but the rather high parameter standard deviations in (I) preclude detailed comparisons. The structures of only two other [4]peristylanes [(VI) and (IX)] with  $sp^3$  hybridization at all four crown positions (C1, C3, C5, C7) are known. The



	ic parameters (A. °)	Table 4. <i>Geometric</i>
--	----------------------	---------------------------

		-	
01-N1	1.24 (1)	C2-C3	1.55(1)
O2-N1	1.18 (1)	C2-C9	1.56(1)
O3—N2	1.23 (1)	C3-C4	1.53 (1)
04—N2	1.21 (1)	C4-C5	1.50(1)
05—N3	121(1)	C4 - C10	1.52 (1)
06—N3	1.16(1)	C5C6	1.52(1)
07_N4	1.10(1)	C6-C7	1.52(1)
09_N4	1.22(1)	C6_C11	1.55(1)
	1.22(1)	C7_C1	1.52(1)
N2 C1	1.50(1)		1.52 (1)
N2-C1	1.55 (1)	$C_{0} = C_{12}$	1.50(2)
NJ-CJ	1.57(1)	C9 - C10	1.50(1)
N4-C3	1.54(1)	09-012	1.54(1)
CI = C2	1.52(1)	C10-C11	1.56(1)
C1-C8	1.51 (1)	C11–C12	1.50 (2)
01-N1-02	125 (1)	N3-C5-N4	100.8 (7)
01-N1-C1	118(1)	N3-C5-C4	107.9 (8)
02-N1-C1	118 (1)	N3-C5-C6	108.9 (8)
O3-N2-O4	128 (1)	N4-C5-C4	115.1 (8)
O3-N2-C1	115(1)	N4-C5-C6	114 2 (8)
04—N2—C1	117(1)	C4C5C6	109.4 (8)
05-N3-06	127 (1)	$C_{5} - C_{6} - C_{7}$	115.0 (8)
05-N3-C5	115 (1)	C5-C6-C11	101 4 (8)
06-N3-C5	119(1)	C7-C6-C11	105.6 (8)
07—N4—08	126 (1)	C6 - C7 - C8	110 7 (8)
07-N4-C5	1159(9)	$C_{1} - C_{8} - C_{7}$	114 7 (8)
08-N4-C5	118.2 (0)	$C_1 = C_2 = C_1^2$	102.2 (0)
N1 - C1 - N2	101 5 (8)	$C_{1}^{-}$ $C_{2}^{-}$ $C_{12}^{-}$	102.3 (9)
N1 - C1 - C2	106.6 (0)	$C_{1}^{2} = C_{0}^{2} = C_{12}^{2}$	109.1 (9)
NI = CI = C2	100.0 (9)	$C_2 = C_3 = C_{10}$	106.5 (6)
$N_{1} - C_{1} - C_{2}$	109.7 (8)	$C_2 = C_3 = C_{12}$	107.6 (9)
$N_2 = C_1 = C_2$	113.7 (6)	C10 - C9 - C12	88.0 (8)
$m_2 - c_1 - c_0$	113.4 (9)	C4 - C10 - C9	108.2 (8)
$C_2 = C_1 = C_0$	109.4 (9)		107.8 (8)
$C_1 = C_2 = C_3$	113.1 (9)		89.4 (7)
C1 - C2 - C9	101.4 (8)		107.1 (8)
$C_{3} - C_{2} - C_{9}$	105.7 (9)	$C_0 = C_1 = C_1 Z_2$	110.0 (9)
$C_2 - C_3 - C_4$	109.9 (8)		90.2 (8)
$C_{3} = C_{4} = C_{3}$	115.9 (9)	12 - 12 - 19	106.8 (9)
$C_{3} - C_{4} - C_{10}$	107.5 (7)	C8-C12-C11	108.3 (9)
C3-C4-C10	100.8 (8)	C9-C12-C11	91.9 (8)
01-N1-C1-N2	17 (1)	O5-N3-C5-N4	-56(1)
01-N1-C1-C2	-105(1)	O5-N3-C5-C4	65 (1)
01-N1-C1-C8	137 (1)	O5-N3-C5-C6	-176(1)
02-N1-C1-N2	-164(1)	O6-N3-C5-N4	124 (1)
02-N1-C1-C2	74 (1)	O6-N3-C5-C4	-115(1)
O2-N1-C1-C8	-44(1)	O6-N3-C5-C6	3 (1)
O3-N2-C1-N1	81 (1)	07-N4-C5-N3	-76(1)
O3-N2-C1-C2	-164(1)	07-N4-C5-C4	168.4 (9)
O3-N2-C1-C8	-37(1)	07-N4-C5-C6	41 (1)
04-N2-C1-N1	-99 (1)	08-N4-C5-N3	97 (1)
O4-N2-C1-C2	16(1)	08-N4-C5-C4	-18(1)
04-N2-C1-C8	144 (1)	08-N4-C5-C6	-146.0(9)
	• · · (•)	50 05 00	. (0.0 (9)

X-ray derived crystal density of  $1.682 \text{ g cm}^{-3}$  is lower than the  $1.70 \text{ g cm}^{-3}$  reported from solvent density measurements (Waykole, Shen & Paquette, 1988). The relative densities of (I) and the 2,2,5,5,9pentanitro derivative (VI;  $1.731 \text{ g cm}^{-3}$ ; unpublished) are in line with the replacement of a basal H atom by NO<sub>2</sub>. The crystal packing is normal. There are no intermolecular distances less than the appropriate van der Waals contact values.



Fig. 4. ORTEP (Johnson, 1976) drawing. The C, N and O atoms are shown as 50% ellipsoids and the H atoms are drawn as spheres with  $B = 1.5 \text{ Å}^2$ .

This work was supported at the University of Maryland by the Office of Naval Research (N00014-90-J-1388) and in part by National Science Foundation Award No. CHE-85-02155, which provided funds for the purchase of a diffractometer system, and the National Institutes of Health for Shared Instrumentation Award No. RR-03354 to purchase a graphics workstation/computer system. Our thanks also to Dr R. S. Miller for his support and encouragement.

#### References

- BUSING, W. R. (1981). WMIN. A Computer Program to Model Molecules and Crystals in Terms of Potential Energy Functions. Report ORNL-5747. Oak Ridge National Laboratory, Tennessee, USA.
- Cambridge Scientific Computing (1990). Chem3D. The Molecular Modeling System. Version 3.0 Cambridge Scientific Computing, Massachusetts, USA.
- DEWAR, M. J. S., ZOEBISCH, E. G., HEALY, E. F. & STEWART, J. J. P. (1985). J. Am. Chem. Soc. 107, 3902–3909.
- Enraf-Nonius (1989). CAD-4 Software. Version 5.0. Enraf-Nonius, Delft, The Netherlands.
- HOLDEN, J. R., DU, Z. & AMMON, H. L. (1993). J. Comput. Chem. 14, 422-437.
- JOHNSON, C. K. (1976). ORTEPII. Report ORNL-5138. Oak Ridge National Laboratory, Tennessee, USA.
- LUO, J., AMMON, H. L. & GILLILAND, G. J. (1989). J. Appl. Cryst. 22, 186.
- Molecular Structure Corporation (1989). TEXSAN. Single Crystal Structure Analysis Software. MSC, Research Forest Drive, The Woodlands, TX 77381, USA.
- STEWART, J. J. P. (1990). MOPAC. A General Molecular Orbital Package. Version 6.0. QCPE Program 455.
- STILL, C. (1990). Macromodel, Columbia Chemistry Molecular Modeling System. Version 3.0. Columbia University, New York, USA.
- WAYKOLE, L. M., SHEN, C. & PAQUETTE, L. A. (1988). J. Org. Chem. 53, 4969–4972.