## 263. Correct Structure of a Condensation Product C<sub>6</sub>H<sub>12</sub>N<sub>4</sub>O<sub>4</sub> from Hydrazine, Formaldehyde and Hydrogen Peroxide

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## Summary

The compound mentioned in the title is shown by X-ray analysis to be 1, 2, 7, 8tetraaza-4, 5, 10, 11-tetraoxatricyclo [6.4.1.1<sup>2,7</sup>]tetradecane, contrary to previous suggestions. The torsion angle about the peroxo bonds in the seven-membered rings in 100°, close to the value in hydrogen peroxide, which would not be possible in a six-membered ring.

The colourless crystalline compound obtained by the reaction of hydrazine with formaldehyde in the presence of hydrogen peroxide [1] was first assigned an incorrect monocyclic structure **1**. The compound was later shown to have the double molecular weight and was assigned the tricyclic structure **2** mainly from its IR. and UV. spectra [2]. More recently, the conformational equilibrium of the compound has been discussed [3] on the basis of the <sup>13</sup>C- and <sup>1</sup>H-(300 MHz)NMR. spectra measured at temperatures down to 200 K. Our attention was directed to **2** as a possible candidate for a low-temperature X-ray crystallographic study aimed at the characterization of the bonding electron density in polar and nonpolar bonds [4].



The results of a low-temperature (96 K) X-ray analysis leave no doubt that the structure is not 2 but 3. A stereo-view of the centrosymmetric molecule is shown in *Figure 1*, bond lengths and angles in *Figure 2*. The identity of our compound with that studied by *Katritzsky et al.* [3] is established by the similarity of measured with published NMR. spectra, which appear to be just as compatible with structure 3 as with 2, or even better, since at low-enough temperature one would expect the high-resolution <sup>1</sup>H-NMR. spectrum of 3 to consist of three *AB*-systems, which is what is observed [3].



Fig. 1. Stereo-view of 3 prepared by program ORTEP [9] (Vibration ellipsoids are shown at the 50% probability level)

Since the reaction leading to 3 is reversible, structure 3 should be thermodynamically more stable than 2, although it contains the two seven-membered rings whose existence was discounted in the arguments leading to the incorrect structure [2]. The tetraaza-cyclohexane ring has the chair conformation with all four N-lone pairs pointing in equatorial directions. The exocyclic N(1), C(1)-bond is



Fig. 2. Bond distances (Å) and angles (deg.) in the centrosymmetric molecule 3 (Standard deviations are about 0.001 Å and 0.1°)

0.025 (2) Å shorter than N(2), C(2), while C(1), O(1) is 0.017 (2) Å longer than C(2), O(2), differences that clearly depend on stereoelectronic factors, the lone pair at N(1) being antiperiplanar to the C(1), O(1)-bond while the lone pair at N(2) is anticlinal to C(2), O(2). In the seven-membered ring the torsion angle about the O, O-bond is 100°, close to the equilibrium torsion angle of about 112° in hydrogen peroxide [5]. In a six-membered ring with approximately tetrahedral bond angles it is almost impossible to obtain a torsion angle much greater than 60°, and this restriction may be the main reason for the thermodynamic stability of **3** compared with **2** since the energy penalty for deviating so far from the equilibrium conformation being about 30 kJ mol<sup>-1</sup> [5]. This argument implies that seven-membered peroxides should generally be more stable than their better known, six-membered counterparts, a generalization that might provoke an experimental enquiry into the matter.

**Crystal Structure of 1,2,7,8-tetraaza-4,5,10,11-tetraoxatricyclo[6.4.1.1<sup>2,7</sup>]tetradecane (3).**  $-C_{6}H_{12}N_{4}O_{4}$ , M=204.18. Crystals are triclinic, space group PI, Z=1 with cell dimensions a = 6.065(3), b = 6.136(1), c = 6.574(1) Å, a = 105.99(1),  $\beta = 102.52(2)$ ,  $\gamma = 112.77(3)^{\circ}$  at 96 K (6.123, 6.194, 6.645 Å, 105.82, 102.92, 112.67° at room temperature). Intensity measurements were made at 96 K with an *Enraf-Nonius* CAD4-diffractometer equipped with graphite monochromator (MoKa.  $\lambda = 0.7107$  Å) and cooling device. 1773 independent reflexions were measured, 1413 with I>5 $\sigma$  (I). The structure was solved by direct methods using the program-system SHELX [6] and refined by full-matrix least-squares analysis using SHELX and XRAY 72 [7] programs with a modified weighting scheme with r = 4.0 Å<sup>2</sup> [8] and an extinction correction. The final R factor was 0.027,  $R_W$  was 0.033.

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