

may find application in the early determination of those phases which can at best be but crudely approximated by (1·1).

4. Concluding remarks

Squared-tangent formulas have been found for the space groups $P\bar{1}$ and $P_{2\bar{1}}2_12_1$. Analogous formulas exist for the remaining noncentrosymmetric space groups. It is anticipated that, at least for the space group $P_{2\bar{1}}2_12_1$ and other, selected, noncentrosymmetric space groups, these formulas will find application in facilitating the early determination of certain phases. The results obtained here raise two questions. First does (3·20) hold also in the space group $P\bar{1}$? Secondly, does (3·20) retain its validity if the \mathbf{k} are restricted so

that the $|E_{\mathbf{k}}$ and $|E_{\mathbf{h}-\mathbf{k}}$ are large, say greater than unity? Although the available evidence indicates strongly that both of these questions are to be answered in the affirmative, rigorous proofs have been elusive so far. With respect to practical applications, the second question is particularly significant since, owing to a reduction of the errors arising from finite sampling, it is better to restrict the vectors \mathbf{k} so that the $|E_{\mathbf{k}}$ and $|E_{\mathbf{h}-\mathbf{k}}$ are large.

References

- International Tables for X-ray Crystallography* (1952). Vol. I, p. 385. Birmingham: Kynoch Press.
 KARLE, J. & HAUPTMAN, H. (1956). *Acta Cryst.* **9**, 635.

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Neutron and X-ray Diffraction Studies of Hydrazinium Sulfate, $\text{N}_2\text{H}_6\text{SO}_4^*$

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Hydrazinium sulfate has been studied by X-ray and neutron diffraction. The crystals are orthorhombic, space group $P_{2\bar{1}}2_12_1$, with four formula units in a cell of dimensions: $a = 8\cdot251$ (5), $b = 9\cdot159$ (1), $c = 5\cdot532$ (1) Å. Both the X-ray and neutron data were affected by extinction; this effect was very severe in the neutron case. By inclusion of six anisotropic extinction parameters in the least-squares refinement it was possible to obtain a good fit between observed and calculated structure factors. The structure consists of $\text{N}_2\text{H}_6^{2+}$ ions and SO_4^{2-} ions held together by a three-dimensional system of N-H···O hydrogen bonds. A number of the hydrogen bonds are weak and bifurcated or trifurcated. The $\text{N}_2\text{H}_6^{2+}$ ion has an almost perfectly staggered conformation. The neutron and X-ray diffraction results are in good agreement with respect to bond lengths involving the non-hydrogen atoms. Although the hydrogen atoms were located with a precision of 0·03 Å in the X-ray study, the mean N-H bond lengths were 0·2 Å shorter than those found from the neutron study, confirming the existence of large systematic errors in the location of hydrogen atoms from X-ray results when a spherical atomic electron distribution is assumed.

Introduction

The crystal structures of a number of hydrazine compounds have been studied; a review of some recent work has been given by Liminga (1968). These compounds often contain a quite complicated arrangement of hydrogen bonds, and an accurate location of the hydrogen atoms is thus essential. Only two hydrazine compounds studied by neutron diffraction have been reported: lithium hydrazinium sulfate, $\text{LiN}_2\text{H}_5\text{SO}_4$, by Padmanabhan & Balasubramanian (1967) and hydra-

zinium hydrogenoxalate, $\text{N}_2\text{H}_5\text{HC}_2\text{O}_4$, by Nilsson, Liminga & Olovsson (1968).

The present investigation involves the refinement of the hydrazinium sulfate structure by both X-ray and neutron diffraction methods. An X-ray diffraction study of this compound has been carried out by Nitta, Sakurai & Tomie (1951); the accuracy of this determination was, however, rather low.

Crystal data

Hydrazinium sulfate, $\text{N}_2\text{H}_6\text{SO}_4$, F.W. 130·12. Orthorhombic, $a = 8\cdot251(5)$, $* b = 9\cdot159(1)$, $c = 5\cdot532(1)$ Å. (Nit-

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* Numbers in parentheses here and throughout this paper are estimated standard deviations in the least significant digits.

eters. The models tried were isotropic extinction and anisotropic extinction for type I and type II crystals. The anisotropic extinction refinements were also carried out with the anisotropic extinction tensors constrained to have orthorhombic symmetry. The parameters refined were positional and anisotropic thermal parameters for all atoms, a scale factor and one, three or six extinction parameters. The function minimized and the weighting scheme were the same as in the X-ray study. The un-averaged set of 858 reflections was now used as data since the extinction components were not equal for the equivalent forms of a reflection. A summary of the resulting R values is given in Table 4; the final values of the extinction parameters are presented in Table 2. A discussion of the physical interpretation of the extinction parameters is given in the paper by Coppens & Hamilton (1970). The unconstrained type II model, particle-size dominated, was concluded to be the most appropriate for this crystal. The final positional and thermal parameters for the unconstrained type II model are presented in Table 1,* and the observed and calculated structure factors together with the extinction coefficients are listed in Table 5. In the last cycle of least-squares refinement no parameter shifted by more than 0.1σ ; the final agreement indices were $R=0.074$ and $R_w=0.096$. The corresponding conventional R index based on F was 0.040. The error in an observation of unit weight was 1.46. The neutron scattering lengths used were $b_S=0.28$, $b_O=0.58$, $b_N=0.94$, $b_H=-0.372$ (10^{-12} cm).

* There was little correlation of the positional and thermal parameters with the type of crystal assumed for the extinction refinement.

Table 4. R values (based on F^2) for neutron data

	R	R_w
Isotropic extinction	0.085	0.109
Constrained Type I	0.081	0.103
Type I	0.074	0.097
Constrained Type II	0.081	0.103
Type II	0.074	0.096

Computer programs

The calculations were carried out on the CDC 6600 computer at the Brookhaven National Laboratory. The following programs from the BNL crystallographic program library were used: *DATAPH* for absorption correction, *FORDAP* for Patterson and Fourier calculations, *LINUS* for least-squares refinement, *NANOVA* for analysis of the weighting scheme at the end of the least-squares refinement, and *ORFFE* and *ORTEP* for structure description.

Comparison of the neutron and X-ray results

Only a few compounds have been studied by both neutron and X-ray diffraction using counter methods; a comparison of the results of the two determinations is therefore of some interest.

There is very satisfactory agreement between the neutron and X-ray results for the positional parameters of the heavy atoms, as can be seen from the values in Table 1. The estimated standard deviations are lower in the X-ray results for the heavy atoms; this reflects the larger number of observations in the X-ray study. The thermal parameters for the two methods are also compared in Table 1; there are clearly differences. The corresponding diagonal elements of the thermal vibratio-

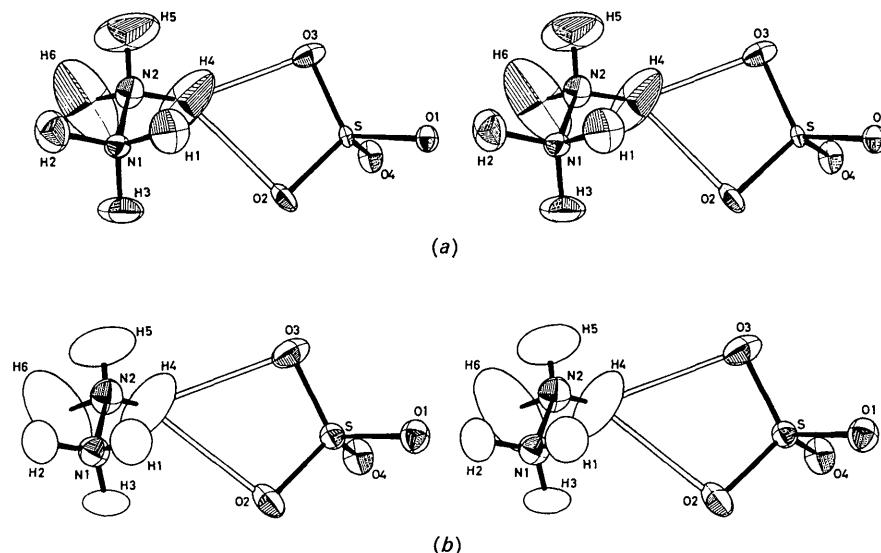


Fig. 1. Stereoscopic illustration of an asymmetric unit of $\text{N}_2\text{H}_6\text{SO}_4$. Thermal ellipsoids are scaled to include 50% probability. (a) Neutron results, (b) X-ray results; the boundary ellipsoids for the hydrogen atoms are taken from the neutron study.

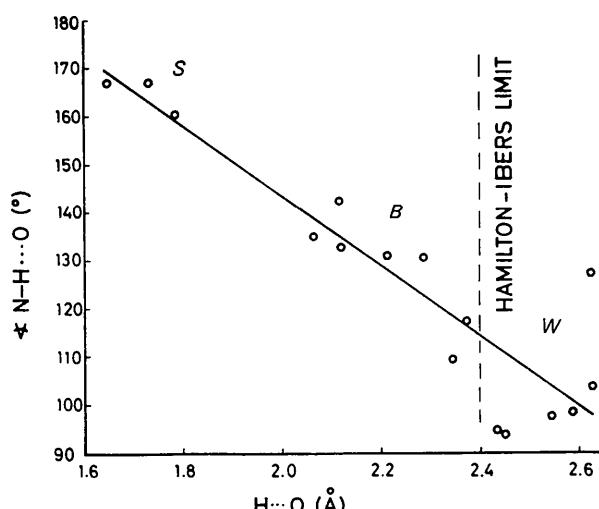


Fig. 3. Hydrogen bond angle-distance correlation in hydrazinium sulfate. *S* — single hydrogen bonds; *B* — bifurcated or trifurcated hydrogen bonds; *W* — weak hydrogen bonds or van der Waals contacts.

thermal motion. An explanation of the large thermal parameters of the hydrogen atoms attached to N(2) in terms of static disorder seems less likely.

The sulfate ion

For the following discussion, we prefer to use the more precise bond lengths from the X-ray study. The neutron values do not, however, differ significantly from the X-ray values.

The geometry of the sulfate ion shows a small but significant departure from a regular tetrahedron as indicated by some variation in the S—O bond lengths and also by the departure of the O—S—O bond angles from the tetrahedral value. This departure can be related to the hydrogen bonding situation around the sulfate group (see Fig. 4). The longest S—O distance,

1.491 Å, involves O(1) which acts as an acceptor for three rather short hydrogen bonds (1.73, 2.06 and 2.12 Å). The next longest S—O distance, 1.483 Å, is to O(2) which also accepts three hydrogen bonds (1.65, 2.21 and 2.37 Å); the longest of these is probably rather weak. The S—O distance to O(3) and O(4) is in both cases 1.468 Å; each of these oxygen atoms acts as an acceptor for two hydrogen bonds.

The mean S—O bond length of 1.477(1) Å is slightly longer than the average of 1.473(1) Å from the mean values of five different sulfate hydrates (Baur, 1964). When a correction for thermal riding motion is applied, the mean value becomes 1.487 Å which is in good agreement with the values of 1.485 and 1.486 given by Baur for the three compounds where the assumptions for this correction were considered to be valid.

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References

- BAUR, W. H. (1964). *Acta Cryst.* **17**, 1361.
- BEAUCAGE, D. R., KELLEY, M. A., OPHIR, D., RANKOWITZ, S., SPINRAD, R. J. & NORTON, R. VAN (1966). *Nucl. Instrum. Methods*, **40**, 26.
- BUSING, W. R. & LEVY, H. A. (1964). *Acta Cryst.* **17**, 142.
- COPPENS, P. & HAMILTON, W. C. (1970). *Acta Cryst.* **A26**, 71.
- COPPENS, P. (1969). *Acta Cryst.* **A25**, 180.
- HAMILTON, W. C. (1965). *Acta Cryst.* **18**, 502.
- HAMILTON, W. C. (1969). *Acta Cryst.* **A25**, 194.
- HAMILTON, W. C. & IBERS, J. A. (1968). *Hydrogen Bonding in Solids: Methods of Molecular Structure Determination*. New York: Benjamin.
- HANSON, H. P., HERMAN, F., LEA, J. D. & SKILLMAN, S. (1964). *Acta Cryst.* **17**, 1040.
- IBERS, J. A. & HAMILTON, W. C. (1964). *Acta Cryst.* **17**, 781.
- International Tables for X-ray Crystallography* (1962). Vol. III, p. 215. Birmingham: Kynoch Press.

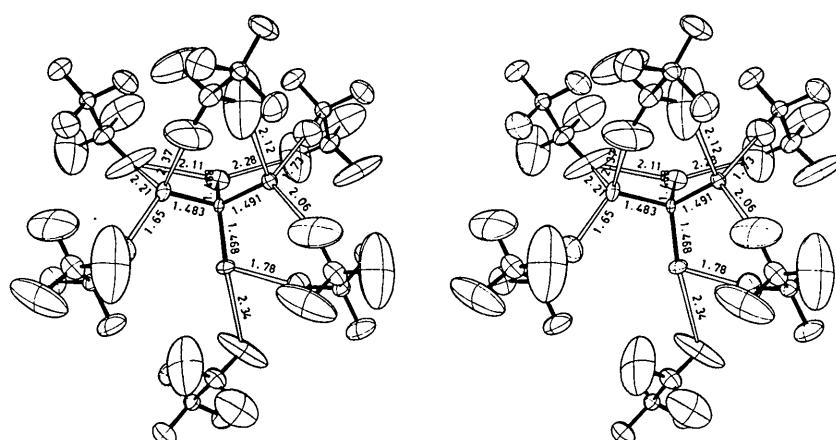


Fig. 4. Stereoscopic illustration of the bonding situation around a sulfate ion. The positional parameters for the sulfate ion are taken from the X-ray study; all other parameters are from the neutron study.

- LIMINGA, R. (1966). *Acta Chem. Scand.* **20**, 2483.
LIMINGA, R. (1968). *Ark. Kemi*, **28**, 483.
NILSSON, Å., LIMINGA, R. & OLOVSSON, I. (1968). *Acta Chem. Scand.* **22**, 719.
NITTA, I., SAKURAI, K. & TOMIIE, Y. (1951). *Acta Cryst.* **4**, 289.
PADMANABHAN, V. M. & BALASUBRAMANIAN, R. (1967). *Acta Cryst.* **22**, 532.
SEQUERA, A., BERKEBILE, C. A. & HAMILTON, W. C. (1968). *J. Mol. Structure*, **1**, 283.
STEWART, R. F., DAVIDSON, E. R. & SIMPSON, W. T. (1965). *J. Chem. Phys.* **42**, 3175.

Acta Cryst. (1970). B**26**, 546

Stereochemistry of Nucleic Acids and their Constituents. VI. The Crystal Structure and Conformation of Dihydrouracil: a Minor Base of Transfer-Ribonucleic Acid*

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Crystals of dihydrouracil are monoclinic, space group *P2*₁/*c*, *Z* = 4, *a* = 4.210 ± 0.001, *b* = 5.816 ± 0.001, *c* = 19.777 ± 0.003 Å, β = 95.15 ± 0.01°, *D*_m = 1.575 g.cm⁻³, *D*_x = 1.577 g.cm⁻³. The structure was solved by direct methods and refined by least squares to an *R* of 0.073, with 639 diffractometric intensities. The saturated ring displays a twist half-chair conformation with C(5) and C(6) displaced by 0.14 and 0.45 on either side of the dihydrouracil plane. The maximum torsion angle of 45.4° is about the saturated C(5)–C(6) bond. The molecule exists in the diketo form, and forms infinite zigzag hydrogen-bonded ribbons such that each base is hydrogen bonded to two adjacent bases related by centers of inversion. O(4) is not involved in hydrogen bonding, whereas O(2) is involved in two hydrogen bonds. The interplanar separation of the bases is 3.34 Å. The base stacking in dihydrouracil is typical of the ketopyrimidines.

Introduction

Transfer ribonucleic acid (*t*-RNA) is probably the smallest (about eighty nucleotides) of the naturally occurring nucleic acids. In addition to the common nucleotides, guanylic, adenylic, cytidylic and uridylic acids, *t*-RNA is composed of approximately 15–20 per cent of a unique class of minor nucleotides (Holley, Apgar, Everett, Madison, Marquisee, Merrill, Penswick & Zamir, 1965; Madison, Everett & Kung, 1966; Zachau, Dütting & Feldman, 1966; RajBhandary, Chang, Stuart, Faulkner, Hoskinson & Khorana, 1967). The majority of the minor nucleotides differ from the common forms only in the composition of the base, although sometimes 2'-*O*-methylated derivatives are also isolated. It is believed that the minor components influence significantly the secondary and tertiary structures of *t*-RNA. To determine the role of the minor components in *t*-RNA, the structural analyses of several minor constituents are presently being

carried out in our laboratory (Rao & Sundaralingam, 1969). In this article, we present the results of the crystal structure of the minor base dihydrouracil (DiHU), which is one of the more common minor bases and is located in the loops of the cloverleaf model for *t*-RNA proposed by Holley *et al.* (1965). A preliminary report on this work has already appeared (Rohrer & Sundaralingam, 1968). Dihydrouracil differs from the common base uracil in that the C(5)–C(6) bond is saturated.

Experimental

Dihydrouracil, purchased from Sigma Chemical Company, St. Louis, Missouri, was crystallized in dimethyl sulfoxide. The crystals showed preferred elongation in the *a*-axis direction and very pronounced cleavage parallel to the (001) plane, thus making selection of a suitable crystal difficult. More than 20 crystals were examined before one without partial cleavage was found. The crystal data for dihydrouracil are given in Table 1. Three-dimensional diffractometric intensities were collected up to $2\theta = 134^\circ$, on a crystal mounted with the long direction parallel to the ϕ axis of the diffractometer, using the θ – 2θ scan mode and a scan speed of 1° per minute. Altogether 823 independent reflections were scanned, and 690 were con-

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