

Acta Cryst. (1956). **9**, 974

The crystal structure of 5'-Br-3':5'-dideoxythymidine. By M. M. WOOLFSON,* *Crystallographic Laboratory, Cavendish Laboratory, Cambridge, England*

(Received 30 July 1956)

The X-ray analysis of 5'-Br-3':5'-dideoxythymidine, which was produced during the study of the chemistry

of certain thymidine derivatives (Michelson & Todd, 1955), was undertaken for the purpose of determining its chemical structure.

The space group was found to be $P2_12_12_1$ with

$$a = 14.78, b = 4.89, c = 15.17 \text{ \AA},$$

there being four molecules per unit cell. The bromine parameters were readily found for the $h0l$ projection from the two-dimensional Patterson synthesis. With the signs of the structure factors taken from the bromine contribution, an electron-density map was calculated which showed clearly the remainder of the atoms. One stage of refinement gave the map shown in Fig. 1, and, since the object of the investigation had been achieved, no further refinement was attempted. The atomic coordinates found for this projection are:

	x	z		x	z
C ₁	0.533	0.304	O ₁	0.364	0.330
C ₂	0.430	0.274	O ₂	0.091	0.318
C ₃	0.416	0.202	O ₃	0.089	0.535
C ₄	0.314	0.221	N ₁	0.236	0.356
C ₅	0.280	0.294	N ₂	0.093	0.434
C ₆	0.266	0.413	Br	0.547	0.400
C ₇	0.226	0.471			
C ₈	0.274	0.522			
C ₉	0.124	0.367			
C ₁₀	0.128	0.486			

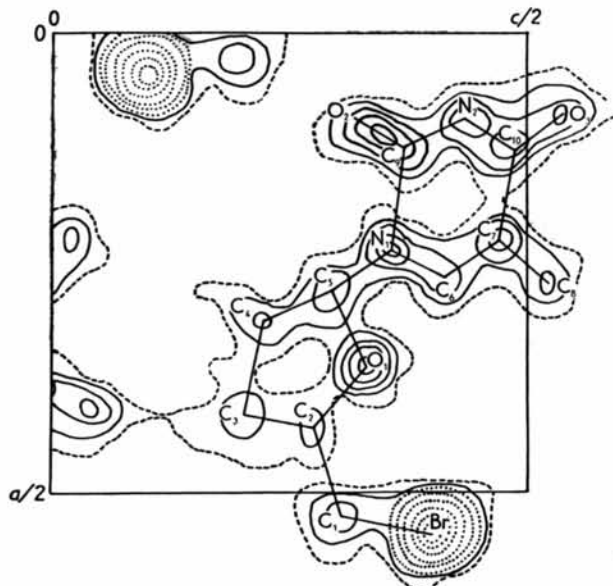


Fig. 1. 5'-Br-3':5'-dideoxythymidine, electron-density map. The contours are at equal arbitrary intervals with the lowest contour broken. The dotted contours round the bromine atom are at twice the interval elsewhere.

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Reference

MICHELSON, A. M. & TODD, A. R. (1955). *J. Chem. Soc.* p. 816.

Acta Cryst. (1956). **9**, 974

The shrinkage of ribonuclease. By C. H. CARLISLE, *Crystallography Laboratory, Birkbeck College, London W. C. 1, England*

(Received 23 May 1956)

In the course of taking an X-ray photograph to record the $h0l$ reflexions of a wet ribonuclease crystal of the monoclinic form (ex aqueous alcohol) in a sealed capillary in the normal manner, a small leak allowed the crystal to dry during its exposure. There was recorded on the photograph both the diffraction pattern of the wet crystal and that of the dry one, with smear lines indicating that a change had taken place. This is most clearly seen (Fig. 1) for the 003 reflexion. The unit-cell dimensions of this form of ribonuclease are:

	a (Å)	b (Å)	c (Å)	β (°)
Wet	30.90	38.8	54.06	106
Dry	29.10	30.08	51.03	114.0

It was possible to confirm from this photograph that the angle β increases when the crystal goes from the wet to the dry state.

The crystal did not move off the capillary wall during the drying process and so Fig. 1 shows the correct orientation.

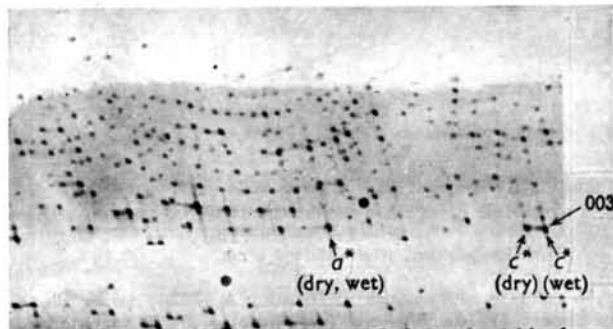


Fig. 1. Weissenberg photograph of ribonuclease, $h0l$ reflexions.