118 K may be ascribed to the anharmonicities which still remain at this temperature.

torsion angles of the C-Ph bond was observed. This is

expected to be accompanied by the shortening of the

ethylenic bond since its bond order is increased by the

twisting of the C-Ph bond. But, in fact, a lengthening

of the ethylenic bond was observed on lowering the

temperature. This result may be explained by the

assumption that on lowering temperature the shortening

of the bond caused by the increase of the torsion

angle of the C-Ph bond was outweighed by the

lengthening due to the decrease of the above-mentioned

The present study demonstrates that X-ray crystal-

lographic data which apparently seem to be normal

may not always give fully reliable geometry for a

molecule and that the molecular structure may be

changed without phase transition in a crystal on

On lowering the temperature, an increase in the

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varving the temperature.

## Structure of 3-Isoadenosine

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Abstract. 3- $\beta$ -D-Ribofuranosyladenine,  $C_{10}H_{13}N_5O_4$ ,  $M_r = 267.25$ , orthorhombic,  $P2_12_12_1$ , a = 13.470 (4), b = 16.054 (6), c = 5.141 (2) Å, V = 1111.8 (7) Å<sup>3</sup>, Z = 4,  $D_x = 1.596 \text{ g cm}^{-3}$ ,  $\lambda (\text{Mo } K\overline{\alpha}) = 0.71073 \text{ Å}$ ,  $\mu$  $= 1.18 \text{ cm}^{-1}$ , F(000) = 560, T = 298 K, R = 0.043, wR = 0.042 for 1136 reflections. The tautomeric form present in the crystal is 6-NH<sub>2</sub>. The adenine moiety is nearly planar and the torsional angle of the glycosidic linkage O(4')-C(1')-N(3)-C(4) is  $-161.5^{\circ}$ . All N and O atoms except N(3) and O(4') participate in a three-dimensional hydrogen-bonding system.

Introduction. 3-Isoadenosine (Leonard & Laursen, 1963, 1965) is an isomer of adenosine in which the D-ribofuranosyl moiety is attached to N(3) of adenine. Because of the close spatial relationship between these two compounds (Leonard, Cruickshank, Groziak, Clauson & Devadas, 1986), the determination of the structure of 3-isoadenosine is desired in order to furnish information concerning the preferred tautomeric form

in the crystal, the orientation of the ribose ring, and the intermolecular hydrogen-bonding pattern, and to compare its structure with adenosine (Lai & Marsh, 1972) and other known N(3)-substituted adenines (Petersen & Furberg, 1975; Kistenmacher, Urmey & Rossi, 1977).

Experimental. The title compound was recrystallized from water as transparent, colorless acicular crystals of dimensions approximately  $0.1 \times 0.2 \times 0.8$  mm;  $D_m$ was not measured. The X-ray data were measured with a Syntex P2, automated four-circle diffractometer using graphite-monochromatized Mo  $K\bar{\alpha}$  radiation. The lattice parameters were refined using 15 reflections with  $2\theta$  values between 17.8 and 19.5°; 1448 unique reflections were measured (+h, +k, +l) to the limit of  $2\theta < 54.0^{\circ}$  using the  $\omega/2\theta$  scan technique, scan rate  $2 \cdot 0 - 15 \cdot 0^{\circ} \text{ min}^{-1}$ , scan range  $1 \cdot 7^{\circ}$  with a scan-tobackground ratio of 0.25, 1136 reflections with  $I > 2.58\sigma(I)$ ,  $\sigma(F_o^2)$  based on counting statistics plus the term  $0.02F_{o}^{2}$ . Anomalous-dispersion, Lorentz, and polarization corrections were applied. Three standard intensities monitored per 100 reflections showed no

anharmonicity.

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significant intensity variation. The structure was solved using the direct-methods program SHELX86 (Sheldrick, 1985). The correct positions for all non-hydrogen atoms were deduced from an E map and were refined (SHELX76; Sheldrick, 1976) with anisotropic thermal coefficients in a process that minimized  $\sum [w(|F_o| - |F_c|)^2]$ , where  $w = 1/[\sigma(F_o)]^2$ . The atomic scattering factors used were taken from International Tables for X-ray Crystallography (1974). H atoms were located by difference electron-density maps and were refined with a group isotropic thermal parameter; an empirical isotropic extinction parameter converged to  $6.15(7) \times 10^{-7}$ . The final agreement factors for 1136 reflections were R = 0.043 and wR = 0.042 and goodness-of-fit E = 1.26; max.  $|\Delta/\sigma| < 0.01$ ; range of residual electron density =  $+0.27 > \Delta \rho > -0.27$  e Å<sup>-3</sup>. Anomalous dispersion using molybdenum radiation would probably be insufficient to distinguish between enantiomers but it is known that the sugar moiety has the D configuration.

Discussion. An ORTEP view (Johnson, 1965) of the molecule of 3-isoadenosine, together with the standard adenosine atom numbering, is given in Fig. 1. The final atomic coordinates for the non-hydrogen atoms are given in Table 1, and selected bond lengths and bond angles are listed in Table 2.\* As has been found for other N(3)-substituted adenines, 3-isoadenosine exists in the 6-NH<sub>2</sub> tautomeric form. Adenosine (Lai & Marsh, 1972) and other N(9)-substituted adenines also exist in 6-NH<sub>2</sub> form. The bond lengths and bond angles of the aglycone of 3-isoadenosine match those of other known N(3)-substituted adenines, namely 3-ethyladenine and tricanthine  $(3-\Delta^2$ -isopentenyladenine)

\* Lists of structure factors, anisotropic thermal parameters, mean-planes data and H-atom parameters have been deposited with the British Library Document Supply Centre as Supplementary Publication No. SUP 44513 (17 pp.). Copies may be obtained through The Executive Secretary, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.



Fig. 1. An ORTEP drawing (Johnson, 1965) of the molecule with the atom numbering.

(Petersen & Furberg, 1975; Kistenmacher et al., 1977). The N(3)-C(1') bond length, 1.488 (5) Å, in 3-isoadenosine, is close to that in tricanthine (1.486 Å) (Kistenmacher et al., 1977) and comparable to the N(9)-C(1') bond length in adenosine (1.466 Å) (Lai & Marsh, 1972).

## Table 1. Final atomic parameters for non-H atoms with e.s.d.'s in parentheses and equivalent isotropic thermal parameters

The equivalent isotropic U is defined as one-third the value of the orthogonalized  $U_{ii}$  tensor.

	х	У	Z	$U_{eq}(Å^2)$
N(1)	0.3453 (2)	0.0656 (2)	-0.0682 (7)	0.028(2)
N(3)	0.4768 (2)	0.1361 (2)	0.1469 (6)	0.023(1)
N(6)	0.3712 (2)	-0.0223(2)	-0.4167 (7)	0.033 (2)
N(7)	0.5952 (2)	0.0036 (2)	-0.3095(6)	0.025(1)
N(9)	0.6436 (2)	0.0976 (2)	0.0048 (6)	0.025(1)
C(2)	0.3799 (3)	0.1175 (2)	0.1069 (8)	0.029 (2)
C(4)	0.5432 (2)	0.0934 (2)	-0.0050 (7)	0.021(2)
C(5)	0.5120(2)	0.0382 (2)	-0·1930 (8)	0.022 (2)
C(6)	0.4091 (2)	0.0255 (2)	-0.2331 (7)	0.024 (2)
C(8)	0.6683 (2)	0.0414 (2)	-0·1826 (8)	0.024 (2)
C(1')	0.5135 (2)	0.1984 (2)	0.3384 (7)	0.023 (2)
C(2')	0.5400 (2)	0.2811 (2)	0.2041 (8)	0.025 (2)
C(3')	0.4431 (3)	0.3294(2)	0.2418(7)	0.024 (2)
C(4′)	0.4102 (2)	0.3015 (2)	0.5097 (7)	0.020 (2)
C(5')	0.2999 (2)	0.3097 (3)	0.5501 (8)	0.027 (2)
O(2')	0.6158 (2)	0.3237 (2)	0.3405 (7)	0.039 (2)
O(3')	0.4550 (2)	0.4169 (2)	0.2324 (5)	0.032(1)
O(4′)	0.4394 (2)	0.2144 (1)	0.5234 (5)	0.027(1)
O(5′)	0.2690 (2)	0.2819 (2)	0.7997 (5)	0.030(2)

Table 2. Selected bond lengths (Å), bond angles (°) and hydrogen-bond parameters

N(1)-C(2)	1.312 (5)	N(3) - C(1')	1.488 (5)
C(2) - N(3)	1.354 (4)	C(1') - C(2')	1.538 (5)
N(3) - C(4)	1.371 (4)	C(2') = O(2')	1.415 (4)
C(4) - C(5)	1.378 (5)	C(2') - C(3')	1.531 (5)
C(5)-C(6)	1.416 (4)	C(3') - O(3')	1.414 (4)
C(6)N(1)	1.368 (4)	C(3') - C(4')	1.515 (5)
C(6)-N(6)	1.320 (5)	C(4')- O(4')	1.453 (4)
C(5)N(7)	1.387 (4)	C(4')-C(5')	1.506 (4)
N(7)C(8)	1.327 (4)	C(5')-O(5')	1.421 (5)
C(8)-N(9)	1.362 (5)	C(1')–O(4')	1.403 (4)
N(9)-C(4)	1.355 (4)		
C(2) - N(1) - C(6)	120.0 (3)	N(3)-C(4)-N(9)	127-2 (3)
N(1)-C(2)-N(3)	125-9 (3)	N(7)-C(5)-C(6)	132-2 (3)
C(2) - N(3) - C(4)	115.6 (3)	C(2)-N(3)-C(1')	124.7 (3)
N(3)-C(4)-C(5)	121.5 (3)	C(4)-N(3)-C(1')	119.7 (3)
C(4) - C(5) - C(6)	119.5 (3)	C(1')-C(2')-C(3')	100.5 (3)
C(5)-C(6)-N(1)	117.2 (3)	C(2')-C(3')-C(4')	102.5 (3)
N(1)-C(6)-N(6)	118.3 (3)	C(3')-C(4')-O(4')	104.5 (3)
N(6) - C(6) - C(5)	124.5 (3)	O(4')-C(1')-C(2')	108-1 (3)
C(5) - N(7) - C(8)	101.8 (3)	C(4') - O(4') - C(1')	109.6 (2)
N(7)C(8)N(9)	118.0 (3)	N(3)-C(1')-O(4')	109.6 (3)
N(9) - C(4) - C(5)	111.3 (3)	C(1')-C(2')-O(2')	111.3 (3)
C(4) - C(5) - N(7)	108.3 (3)	C(2')-C(3')-O(3')	113.8 (3)
C(4)N(9)C(8)	100.6 (3)		
N(6H)AN(1)	2.21 (4)	N(6)A = N(1)	2 007 (4)
N(6H)RO(2')	2.21(4)	N(6)R = O(2')	2.012 (4)
H(01)D(0(3))	1.07(3)	O(2') = O(5')	2 766 (4)
H(2') = H(3')	1.37 (3)	$O(2) \cdots O(3)$	2.700 (4)
H(5')N(9)	1.08 (4)	$O(5') \cdots N(7)$ $O(5') \cdots N(9)$	2.009 (4)
N(6) - H(6) A - N(1)	1.90(4)	O(2') = U(2') O(5')	2.137 (4)
N(6) = H(6)RO(3)	(3)	$O(2') = H(2') \cdots O(3')$	158 (3)
11(0) 11(0) <b>D</b> O(3	, 155(5)	$O(5') = H(5') \dots N(0)$	170 (4)



Fig. 2. Stereoscopic view of the unit cell with hydrogen bonding.

The adenine ring of the molecule is nearly planar (Fig. 1), but is slightly folded about the C(4)–C(5) vector with an interplanar angle of  $2.5^{\circ}$ . The torsionangle values of N(1)–C(6)–N(6)–HN(6)A and N(1)– C(6)–N(6)–HN(6)B, 1 (3) and 171 (3)°, respectively, are indicative of the approximate coplanarity of the two amino H atoms at N(6) with the bicyclic ring. As in adenosine, the furanose ring is puckered with a C(3') *endo* conformation.

A stereoview of the crystal packing arrangement is shown in Fig. 2. As with adenosine, all N and O atoms except N(3) and O(4') [O(1')] participate in a threedimensional hydrogen-bonding scheme. In the crystal structure of adenosine the 6-amino group donates a hydrogen bond to both the N(7) and O(5') atoms of a neighboring adenosine molecule. In tricanthine and 3-ethyladenine, this same amino group participates in a hydrogen-bonding scheme with two neighboring molecules. In tricanthine the N(1) atom of one neighbor and the N(7) atom of the other participate in this scheme, whereas in 3-ethyladenine it is the N(7) atom of one neighbor and the N(9) atom of the second neighbor that participate. 3-Isoadenosine represents yet another variation in the hydrogen-bonding pattern. The 6amino-group H bonds with the N(1) atom of one neighbor and the O(3') atom of the second neighboring molecule. The hydrogen-bond-related lengths and angles are provided in Table 2.

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# Structure of (4SR,6aRS)-4-[2-(2-Methyl-1,3-dioxolan-2-yl)ethyl]-4,5,6,6a-tetrahydro-5,5,6a-trimethyl-2(1H)-pentalenone

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(Received 30 March 1987; accepted 28 October 1987)

Abstract.  $C_{17}H_{26}O_3$ ,  $M_r = 278 \cdot 21$ , triclinic,  $P\overline{1}$ , a = 7.545 (1), b = 10.465 (2), c = 10.629 (1) Å,  $\alpha = 95.41$  (1),  $\beta = 103.32$  (1),  $\gamma = 104.30$  (1)°, V = 781.1 (2) Å<sup>3</sup>, Z = 2, F(000) = 304,  $D_x = 1.18$  Mg m<sup>-3</sup>,  $\lambda$ (Mo K $\alpha$ ) = 0.71073 Å,  $\mu$ (Mo K $\alpha$ ) = 0.74 cm<sup>-1</sup>, T = 133 K, R = 0.046, wR = 0.053 with 1659 independent

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non-zero reflections. The stereochemistry is 4SR, 6aRS, the two fused rings form a boat-like arrangement and the crystal structure is stabilized by distinctive directionally-specific C-H···O hydrogen bonds.

Introduction. In the course of the stereoselective synthesis of  $(\pm)$ -silphenene from isophorone (Rao & Nagarajan, unpublished results) the pentalenone (1)

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