Fig. 2. Projection of the crystal structure down **b**.

$C(12)$  and  $C(14)$ . This interaction is also responsible for the flattening of ring *B* at  $C(9)$ . Analogously, the widening of angle  $C(9)-C(8)-C(13)$  to  $114.2(4)^\circ$  is a demand of the interaction  $C(12)\cdots C(13)$ , while the short contact ( $1.96\text{ \AA}$ ) between two H atoms at  $C(1)$  and  $C(11)$  causes the angle  $C(1)-C(10)-C(9)$  to widen to  $114.4(3)^\circ$ .

Otherwise, the widening of the angle  $C(9)-C(11)-C(16)$  to  $119.1(4)^\circ$  is probably due to requirements of the packing of the hydroquinone fragment which, in this way, becomes perpendicular to the mean plane through the sesquiterpene system

[interplanar angles:  $91(1)$  for molecule 1 and  $86(1)^\circ$  for molecule 2].

The crystal packing, illustrated in Fig. 2, shows weak van der Waals contacts.

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#### References

- BUCHOURT, R. & HAINAUT, D. (1965). *Bull. Soc. Chim.* pp. 1366–1377.
- CARIELLO, L., DE NICOLA GIUDICI, M. & ZANETTI, L. (1980). *Comp. Biochem. Physiol. C*, **65**, 37–41.
- CREMER, D. & POPLE, J. A. (1975). *J. Am. Chem. Soc.* **97**, 1354–1358.
- DE ROSA, S., MINALE, L., RICCIO, R. & SODANO, G. (1976). *J. Chem. Soc. Perkin Trans. 1*, pp. 1408–1414.
- DJURA, P., STERLE, D. B., SULLIVAN, B., FAULKNER, D. J., ARNOLD, E. & CLARDY, J. (1980). *J. Org. Chem.* **45**, 1435–1441.
- DUAX, W. L., WEEKS, C. M. & ROHRER, D. C. (1976). *Top. Stereochem.* **9**, 271–383.
- ENRAF-NONIUS (1979). *Structure Determination Package*. Enraf-Nonius, Delft.
- INTERNATIONAL TABLES FOR X-RAY CRYSTALLOGRAPHY (1974). Vol IV. Birmingham: Kynoch Press. (Present distributor D. Reidel, Dordrecht.)
- MAIN, P., HULL, S. E., LESSINGER, L., GERMAIN, G., DECLERCQ, J.-P. & WOOLFSON, M. M. (1978). *MULTAN78. A System of Computer Programs for the Automatic Solutions of Crystal Structures from X-ray Diffraction Data*. Univs. of York, England, and Louvain, Belgium.
- MINALE, L., RICCIO, R. & SODANO, G. (1974). *Tetrahedron Lett.* **38**, 3401–3404.

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## Structure of *N*-Benzoyl-5'-*O*-*tert*-butyldimethylsilyl-2'-deoxyadenosine Monohydrate

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**Abstract.**  $C_{23}H_{31}N_5O_4Si.H_2O$ ,  $M_r = 487.6$ , orthorhombic,  $P2_12_12_1$ ,  $a = 18.642(2)$ ,  $b = 8.032(1)$ ,  $c = 16.825(2)\text{ \AA}$ ,  $V = 2519.3(1)\text{ \AA}^3$ ,  $Z = 4$ ,  $D_x = 1.286\text{ g cm}^{-3}$ ,  $\lambda(\text{Cu } K\alpha) = 1.5418\text{ \AA}$ ,  $\mu = 10.85\text{ cm}^{-1}$ ,  $F(000) = 1040$ ,  $T = 283\text{ K}$ ,  $R = 0.067$  for 1664 unique observed reflections. The glycosidic torsion angle,  $\chi = 161.8(10)^\circ$ , is in the *anti* conformation. The deoxyribose ring has a  $C3'$ -*exo* ( $C2'$ -*endo*) pucker and the conformation around the  $C5'$ – $C4'$  bond is *gauche*–

*gauche*. The mean Si–C bond length is  $1.861(6)\text{ \AA}$  (three evaluations), and the Si–O $5'$  bond length is  $1.658(6)\text{ \AA}$ . The crystal structure is stabilized by a fully satisfied hydrogen-bond network involving the water of crystallization and by stacking interactions between the phenyl ring and the base. Finally, there are weakly polar C–H $\cdots$ O interactions between C8 and O $5'$  and between the  $\delta(+)$  edge of the phenyl ring and the water of crystallization.

**Introduction.** The crystal structures of 2'-deoxyadenosine monohydrate (Watson, Sutor & Tollin, 1965) and 2'-deoxyadenosine (Sato, 1984) have been reported. We describe the solution of the structure of a monohydrate of a protected form of 2'-deoxyadenosine, which is part of a series of crystallographic investigations of the reagents for oligonucleotide synthesis.

**Experimental.** Rectangular crystals from a water/methanol mixture by slow evaporation,  $0.2 \times 0.2 \times 0.8$  mm, Nicolet P3 diffractometer, Ni-filtered radiation,  $\omega$ -scan method,  $(\sin\theta)/\lambda < 0.58 \text{ \AA}^{-1}$ , lattice parameters from the  $2\theta$  values of ten reflections with  $30 < 2\theta < 50^\circ$ , no absorption corrections,  $h = 0$  to 20,  $k = 0$  to 8,  $l = 0$  to 19, three reflections measured as intensity standards with variation  $< 2\%$ . 2395 unique reflections measured, 731 excluded during refinement [ $|F| < 3\sigma(F)$ ]. Structure solved by heavy-atom phasing from silicon atom, first  $F$  map revealed the positions of all non-hydrogen atoms, least-squares refinement using SHELLX76 (Sheldrick, 1976),  $F$  magnitudes, unit weights; isotropic and then anisotropic temperature factors gave  $R = 0.067$  and  $S = 1.77$  [calculated with  $\sigma(F)$ ]. 311 parameters varied:  $x$ ,  $y$ ,  $z$ ,  $U_{ij}$  for non-H atoms and H atoms at calculated positions allowed to 'ride' on non-H atom, with a  $U$  varied for each group of H atoms bound to the phenyl ring, the base, the sugar and the methyl carbons, respectively. In final cycle  $(\Delta/\sigma)_{\max} = 0.015$ . Final difference synthesis  $(\Delta\rho)_{\max} = 0.29$ ,  $(\Delta\rho)_{\min} = -0.48 \text{ e \AA}^{-3}$ . Scattering factors from International Tables for X-ray Crystallography (1974).

**Discussion.** Table 1\* gives the final atom parameters, and Fig. 1 shows the molecular structure drawn by ORTEPII (Johnson, 1976) and the atomic numbering scheme.

The 2'-deoxyribose ring adopts a C3'-*exo* ( $\text{C}_3T^2$ ) ( $\text{C}_2'$ -*endo*) pucker with a pseudorotation angle  $P \approx 190^\circ$ , which is often found in 2'-deoxynucleosides and nucleotides (Olson & Sussman, 1982). The glycosidic torsion angle defined by O4'-C1'-N9-C4,  $\chi = 161.8(10)^\circ$ , is in the *anti* conformation, giving a close contact between sugar and base [O4'...C8 = 2.799(11) Å] similar to that seen in 2'-deoxyadenosine (Sato, 1984). In addition, the conformation about the C4'-C5' bond defined by O5'-C5'-C4'-C3',  $\gamma = 56.0(11)^\circ$ , is *gauche-gauche* (+sc), which is stabilized by an intramolecular 'hydrogen bond' [C8-H...O5' = 3.241(11) Å] that is only rarely observed in

purine nucleotides and nucleosides (Amidon, Anik & Rubin, 1975). The bond lengths in the adenine base are in good agreement with the mean values for the neutral base calculated by Taylor & Kennard (1982), except for C2-N3 = 1.359(13) Å and C6-N6 = 1.386(10) Å, which exceed their mean values, and C4-C5 = 1.353(12) Å and C6-N1 = 1.317(11) Å, which are less than their mean values. The three Si-C bond lengths vary between 1.857(11) and 1.864(11) Å [mean = 1.861(6) Å] and the Si-O5'

Table 1. *Atomic coordinates and equivalent isotropic thermal parameters with e.s.d.'s in parentheses*

|       | $x$         | $y$         | $z$         | $U_{eq}(\text{\AA}^2)$ |
|-------|-------------|-------------|-------------|------------------------|
| M1    | 0.3011(8)   | 0.2071(16)  | 0.3222(7)   | 0.103                  |
| M2    | 0.2498(7)   | 0.592(2)    | 0.3139(8)   | 0.117                  |
| M3    | 0.3762(7)   | 0.6115(19)  | 0.3548(7)   | 0.107                  |
| M4    | 0.3434(10)  | 0.6973(15)  | 0.2141(7)   | 0.126                  |
| M5    | 0.4356(6)   | 0.304(2)    | 0.2299(8)   | 0.119                  |
| Si    | 0.33964(15) | 0.3555(4)   | 0.24887(16) | 0.055                  |
| CSi   | 0.3278(6)   | 0.5723(14)  | 0.2850(6)   | 0.065                  |
| O5'   | 0.2931(4)   | 0.3234(9)   | 0.1663(3)   | 0.058                  |
| C1'   | 0.2194(5)   | 0.0114(11)  | 0.0185(6)   | 0.055                  |
| C2'   | 0.2952(5)   | -0.0615(13) | 0.0059(6)   | 0.058                  |
| C3'   | 0.3415(5)   | 0.0697(12)  | 0.0516(5)   | 0.051                  |
| O3'   | 0.4137(3)   | 0.0688(9)   | 0.0288(4)   | 0.063                  |
| C4'   | 0.3028(5)   | 0.2325(12)  | 0.0312(5)   | 0.050                  |
| O4'   | 0.2275(3)   | 0.1886(7)   | 0.0283(4)   | 0.050                  |
| C5'   | 0.3134(6)   | 0.3729(13)  | 0.0899(5)   | 0.059                  |
| N1    | 0.0799(5)   | -0.4738(10) | 0.1375(4)   | 0.056                  |
| C2    | 0.1086(6)   | -0.4494(16) | 0.0660(6)   | 0.066                  |
| N3    | 0.1472(4)   | -0.3169(11) | 0.0398(4)   | 0.057                  |
| C4    | 0.1517(5)   | -0.2040(12) | 0.0983(5)   | 0.047                  |
| C5    | 0.1255(5)   | -0.2126(13) | 0.1732(5)   | 0.046                  |
| C6    | 0.0899(5)   | -0.3620(13) | 0.1937(5)   | 0.048                  |
| N7    | 0.1408(4)   | -0.0676(10) | 0.2160(4)   | 0.050                  |
| C8    | 0.1764(5)   | 0.0243(14)  | 0.1637(5)   | 0.057                  |
| N9    | 0.1857(4)   | -0.0513(11) | 0.0924(4)   | 0.050                  |
| N6    | 0.0658(4)   | -0.3844(10) | 0.2709(4)   | 0.049                  |
| C     | 0.0566(5)   | -0.5394(14) | 0.3083(5)   | 0.054                  |
| O     | 0.0708(5)   | -0.6717(10) | 0.2752(4)   | 0.079                  |
| C1(B) | 0.0296(5)   | -0.5314(13) | 0.3894(5)   | 0.045                  |
| C2(B) | 0.0477(5)   | -0.6596(14) | 0.4389(5)   | 0.058                  |
| C3(B) | 0.0232(6)   | -0.6615(16) | 0.5181(6)   | 0.070                  |
| C4(B) | -0.0198(6)  | -0.5335(16) | 0.5465(6)   | 0.074                  |
| C5(B) | -0.0377(6)  | -0.4038(16) | 0.4945(6)   | 0.070                  |
| C6(B) | -0.0133(5)  | -0.4010(14) | 0.4160(6)   | 0.056                  |
| O(W)  | 0.0701(3)   | -0.0728(8)  | 0.3611(3)   | 0.052                  |

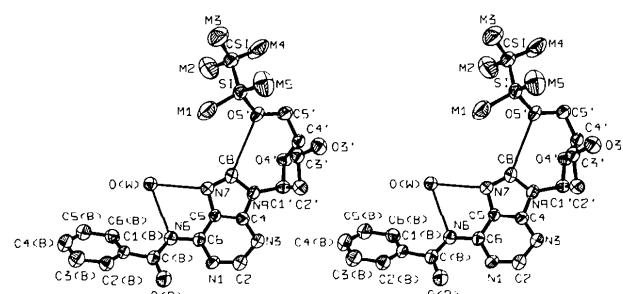


Fig. 1. Stereodrawing of the molecular geometry with the atom numbering. The thermal ellipses are drawn at the 50% level.

\* Lists of structure-factor amplitudes, anisotropic thermal parameters, bond lengths and bond angles, and H-atom parameters have been deposited with the British Library Document Supply Centre as Supplementary Publication No. SUP 43678 (16 pp.). Copies may be obtained through The Executive Secretary, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

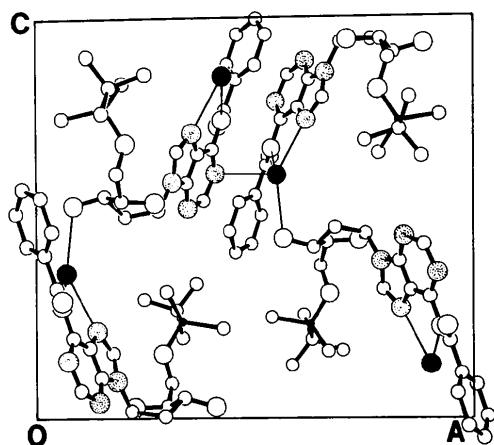


Fig. 2. Packing of the molecules in the unit cell. Atoms are identified by the following: N = large shaded circles, O = large open circles, H<sub>2</sub>O = large black circles, C = small open circles, Si = small black circles. Hydrogen bonds are indicated with narrow lines.

bond length is 1.658 (6) Å, which are in good agreement with previously published values for Si-X distances given in *International Tables for X-ray Crystallography* (1974).

The crystal structure is illustrated in Fig. 2. All available H atoms participate in hydrogen bonds [N6—H···O(W) = 2.929 (11), N7—H···O(W) = 2.775 (11) Å, O3'—H···O(W') = 2.838 (10) and N1—H···O(W') = 2.907 (11) Å], and there are no adenine—adenine base pairs. There is a stacking interaction between the phenyl ring and the base

[C5···C1(B) = 3.339 (11), C8···C6(B) = 3.377 (10), N9···C5(B) = 3.339 (11), N9···C6(B) = 3.435 (11) Å]. Finally, there is a close contact between C6 of the phenyl ring and O of the water of crystallization [C6(B)—H···O(W) = 3.197 (11) Å], which brings the δ(−) oxygen atom close to one of the δ(+) hydrogen atoms of the phenyl ring and is enthalpically favorable (Thomas, Smith, Thomas & Feldman, 1982).

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#### References

- AMIDON, G. L., ANIK, S. & RUBIN, J. (1975). *Structure and Conformation of Nucleic Acids and Protein–Nucleic Acid Interactions*, edited by M. SUNDRAMINGHAM & S. T. RAO, pp. 729–744. Univ. Park Press, Baltimore.
- International Tables for X-ray Crystallography* (1974). Vol. IV, pp. 72–75. Birmingham: Kynoch Press. (Present distributor D. Reidel, Dordrecht.)
- JOHNSON, C. K. (1976). ORTEPII. Report ORNL-5138. Oak Ridge National Laboratory, Tennessee.
- OLSON, W. K. & SUSSMAN, J. L. (1982). *J. Am. Chem. Soc.* **104**, 270–278.
- SATO, T. (1984). *Acta Cryst. C* **40**, 880–882.
- SHEDDRICK, G. M. (1976). SHELLX76. Program for crystal structure determination. Univ. of Cambridge, England.
- TAYLOR, R. & KENNARD, O. (1982). *J. Mol. Struct.* **78**, 1–28.
- THOMAS, K. A., SMITH, G. M., THOMAS, T. M. & FELDMAN, R. J. (1982). *Proc. Natl Acad. Sci. USA*, **79**, 4843–4847.
- WATSON, D. G., SUTOR, D. J. & TOLLIN, P. (1965). *Acta Cryst.* **19**, 111–124.

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## Structure and Stereochemistry of the Methyl Ester of (5α,13α,14β,17α,20S,24Z)-3-Oxolanosta-7,24-dien-26-oic Acid (Masticadienonic Acid)\*†

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(Received 1 October 1986; accepted 23 December 1986)

**Abstract.** C<sub>31</sub>H<sub>48</sub>O<sub>3</sub>, M<sub>r</sub> = 468.7, orthorhombic, P2<sub>1</sub>2<sub>1</sub>2<sub>1</sub>, a = 7.118 (1), b = 10.165 (3), c =

38.244 (10) Å, V = 2767 (1) Å<sup>3</sup>, Z = 4, D<sub>x</sub> = 1.125 Mg m<sup>-3</sup>, λ(Mo Kα) = 0.7107 Å, μ = 0.065 mm<sup>-1</sup>, F(000) = 1032, T = 293 K. Final R = 0.066 for 1636 observed reflections. The molecular structure determined from the X-ray data confirms the structure previously assigned on the basis of chemical and spectroscopic evidence. All the rings are *trans*

\* Contribution No. 838 of the Instituto de Química, UNAM.

† Note that the crystallographic numbering differs from that used to name the compound.

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