

Lists of structure factors, anisotropic displacement parameters, H-atom coordinates and complete geometry, together with a stereopacking diagram, have been deposited with the IUCr (Reference: TA1032). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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Acta Cryst. (1996). **C52**, 250–252

2,8-Dithia-1-phospha-5-arsabicyclo[3.3.0]-octane 1-Oxide

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(Received 3 May 1995; accepted 16 August 1995)

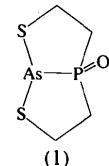
Abstract

The title compound, $C_4H_8AsOPS_2$, obtained by oxidation of $As(SCH_2CH_2)_2P$, shows a bicyclooctane structure with an As—P bond distance of 2.345 (1) Å. The compound shows weak intermolecular As···O interactions of 2.882 (2) Å.

Comment

In 1982, we reported a number of 2,8-dithia-1-phospha-5-arsabicyclo[3.3.0]octanes (Jurkschat, Mügge, Tzschach, Uhlig & Zschunke, 1982). A detailed conformational study of these compounds in solution, followed later by NMR spectroscopic analysis (Zschunke, Mügge, Meyer, Tzschach & Jurkschat,

1983), suggested that they exist in a preferred conformer with strong puckering of the fused five-membered rings. No single crystals of any of these compounds has been obtained thus far. However, oxide $As(SCH_2CH_2)_2PO$, hereafter (1), was obtained as a minor product by recrystallization of $As(SCH_2CH_2)_2P$ from toluene in the presence of air. Its ^{31}P NMR spectrum shows a singlet at 128.8 p.p.m., which is consistent with the presence of pentavalent phosphorus.



The twist conformation of the two fused five-membered rings, $As—S1—C1—C2—P1$ [$\varphi(2) = -118.9(4)^\circ$] and $As—S2—C4—C3—P1$ [$\varphi(2) = 94.6(3)^\circ$], of (1) in the solid state resembles that predicted for $As(SCH_2CH_2)_2P$ in solution (Zschunke *et al.*, 1983). The As—P distance of 2.345 (1) Å may be regarded as normal; it resembles those reported for other As—P compounds (Baudler, Aktalay, Heinlein & Tebbe, 1982; Weber, Bungardt & Boese, 1989; Jutzi, Meyer, Opiela, Olmstead & Power, 1990; Märkl, Dietl, Ziegler & Nuber, 1988; Sheldrick, 1975). The same holds for the As—S bond distances (Dräger, 1975). An interesting feature of the structure of (1) is the weak intermolecular As···O interaction of 2.882 (2) Å resulting in head-to-tail dimers. This distance is at the upper limit of such contacts (Dräger, 1975) and reflects the ambivalent character of (1), with weak Lewis acidity at the As atom and Lewis basicity at the P=O function. The P=O bond length of 1.490 (2) Å is not affected by the weak coordination and resembles those found in functional substituted organotin compounds (Weichmann, Rensch, Dargatz & Meunier-Piret, 1989; Weichmann & Meunier-Piret, 1993).

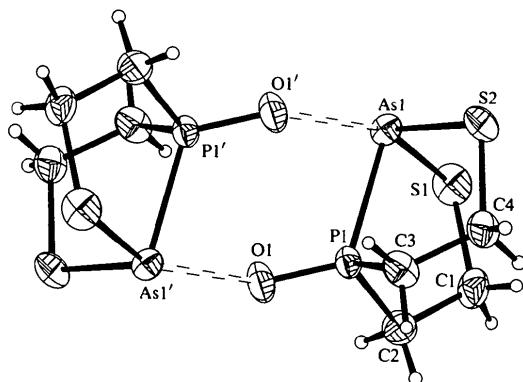


Fig. 1. ORTEPII (Johnson, 1976) plot of the title compound showing the atom-numbering scheme with 50% probability ellipsoids.

Experimental

Crystals of the title compound were obtained by recrystallization of As(SCH₂CH₂)₂P from toluene in the presence of air.

Crystal data



*M*_r = 242.11

Triclinic

P̄*I*

a = 6.522 (3) Å

b = 6.852 (1) Å

c = 10.539 (1) Å

α = 71.24 (1)°

β = 78.88 (3)°

γ = 63.29 (3)°

V = 397.7 (2) Å³

Z = 2

*D*_x = 2.022 Mg m⁻³

Data collection

Enraf–Nonius CAD-4
diffractometer

$\omega/2\theta$ scans

Absorption correction:
refined from ΔF

(DIFABS; Walker &
Stuart, 1983)

T_{\min} = 0.330, T_{\max} =
1.000

2495 measured reflections

2307 independent reflections

2069 observed reflections

[*I* > 2σ(*I*)]

Refinement

Refinement on *F*²

R[*F*_o > 4σ(*F*_o)] = 0.0297

wR(*F*²) = 0.0826

S = 1.120

2307 reflections

86 parameters

Only H-atom *U*'s refined

$w = 1/[\sigma^2(F_o^2) + (0.0467P)^2
+ 0.3211P]$

where *P* = (*F*_o² + 2*F*_c²)/3

Mo K α radiation

λ = 0.7107 Å

Cell parameters from 75
reflections

θ = 14–15°

μ = 4.917 mm⁻¹

T = 298 (2) K

Rectangular prism

0.90 × 0.64 × 0.50 mm

Colourless

*R*_{int} = 0.0201

θ_{\max} = 14.99°

h = 0 → 9

k = -8 → 9

l = -14 → 14

3 standard reflections

frequency: 50 min

intensity decay: 5%

Extinction correction: none

Atomic scattering factors

from International Tables

for Crystallography (1992,

Vol. C, Tables 4.2.6.8 and

6.1.1.4)

(Δ/σ)_{max} < 0.001

$\Delta\rho_{\max}$ = 0.543 e Å⁻³

$\Delta\rho_{\min}$ = -0.682 e Å⁻³

Extinction correction: none

Atomic scattering factors

from International Tables

for Crystallography (1992,

Vol. C, Tables 4.2.6.8 and

6.1.1.4)

Table 2. Selected geometric parameters (Å, °)

As—S2	2.2354 (12)	P—O1	1.490 (2)
As—S1	2.2623 (8)	P—C3	1.811 (3)
As—P	2.3447 (11)	P—C2	1.817 (3)
As—O1 ⁱ	2.882 (2)	C1—C2	1.521 (4)
S1—C1	1.824 (3)	C3—C4	1.516 (4)
S2—C4	1.834 (3)		
S2—As—S1	103.13 (4)	O1—P—C2	110.83 (13)
S2—As—P	90.83 (4)	C3—P—C2	108.9 (2)
S1—As—P	90.09 (4)	O1—P—As	118.06 (10)
S2—As—O1 ⁱ	85.77 (5)	C3—P—As	100.97 (10)
S1—As—O1 ⁱ	160.32 (5)	C2—P—As	102.01 (10)
P—As—O1 ⁱ	72.05 (5)	C2—C1—S1	111.6 (2)
C1—S1—As	101.05 (10)	C1—C2—P	114.5 (2)
C4—S2—As	103.58 (9)	C4—C3—P	112.8 (2)
O1—P—C3	114.86 (14)	C3—C4—S2	112.7 (2)
S2—As—S1—C1	-66.38 (11)	S1—As—P—C2	-3.51 (11)
P—As—S1—C1	24.50 (11)	O1 ⁱ —As—P—C2	-175.09 (12)
O1 ⁱ —As—S1—C1	48.9 (2)	As—S1—C1—C2	-46.8 (2)
S1—As—S2—C4	81.24 (11)	S1—C1—C2—P	49.4 (3)
P—As—S2—C4	-9.05 (11)	O1—P—C2—C1	-151.4 (2)
O1 ⁱ —As—S2—C4	-80.99 (12)	C3—P—C2—C1	81.4 (3)
S2—As—P—O1	-138.69 (10)	As—P—C2—C1	-24.8 (3)
S1—As—P—O1	118.18 (10)	O1—P—C3—C4	166.3 (2)
O1 ⁱ —As—P—O1	-53.40 (13)	C2—P—C3—C4	-68.8 (2)
S2—As—P—C3	-12.63 (11)	As—P—C3—C4	38.1 (2)
S1—As—P—C3	-115.76 (11)	P—C3—C4—S2	-50.8 (3)
O1 ⁱ —As—P—C3	72.65 (12)	As—S2—C4—C3	35.4 (2)
S2—As—P—C2	99.63 (11)		

Symmetry code: (i) $1 - x, 1 - y, -z$.

Data collection: CAD-4 Software (Enraf–Nonius, 1989).

Cell refinement: CELSIUS (Svenson, 1974). Data reduction:

CORINC (Dräger & Gattow, 1971). Program(s) used to solve

structure: SHELLXS86 (Sheldrick, 1985). Program(s) used to refine structure: SHELLXL93 (Sheldrick, 1993). Molecular graph-

ics: ORTEPII (Johnson, 1976).

Lists of structure factors, anisotropic displacement parameters, H-atom coordinates and complete geometry have been deposited with the IUCr (Reference: JZ1062). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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Table 1. Fractional atomic coordinates and equivalent isotropic displacement parameters (Å²)

$$U_{\text{eq}} = (1/3) \sum_i \sum_j U_{ij} a_i^* a_j^* \mathbf{a}_i \cdot \mathbf{a}_j$$

	<i>x</i>	<i>y</i>	<i>z</i>	<i>U</i> _{eq}
As	0.17165 (4)	0.39448 (4)	0.16387 (3)	0.02818 (9)
S1	0.18837 (14)	0.19937 (13)	0.38312 (8)	0.0399 (2)
S2	-0.08575 (11)	0.74476 (12)	0.16791 (8)	0.0356 (2)
P	0.47165 (10)	0.48049 (10)	0.18397 (6)	0.02427 (13)
O1	0.6894 (3)	0.3959 (4)	0.1013 (2)	0.0376 (4)
C1	0.3144 (5)	0.3337 (5)	0.4495 (3)	0.0377 (6)
C2	0.5210 (5)	0.3568 (6)	0.3619 (3)	0.0372 (6)
C3	0.3264 (5)	0.7861 (4)	0.1514 (3)	0.0346 (6)
C4	0.0819 (5)	0.8663 (5)	0.2119 (3)	0.0348 (6)

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Acta Cryst. (1996). C52, 252–254

N-(p-Tolyl)-amine-1-D-fructose from a Small Crystal

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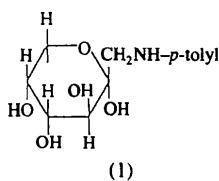
(Received 30 March 1994; accepted 11 August 1995)

Abstract

Non-enzymatic glycation (reaction of an amino group with a sugar) is the first step in a complex, poorly understood, series of Maillard reactions. The reaction can also serve as a model for the general non-enzymatic processing of proteins. The structure of the title compound, C₁₃H₁₉NO₅, is a model for the product in such a process. Despite the small volume of the crystal, the use of a Cu K α rotating-anode source allowed the collection of sufficient data to solve and refine the structure.

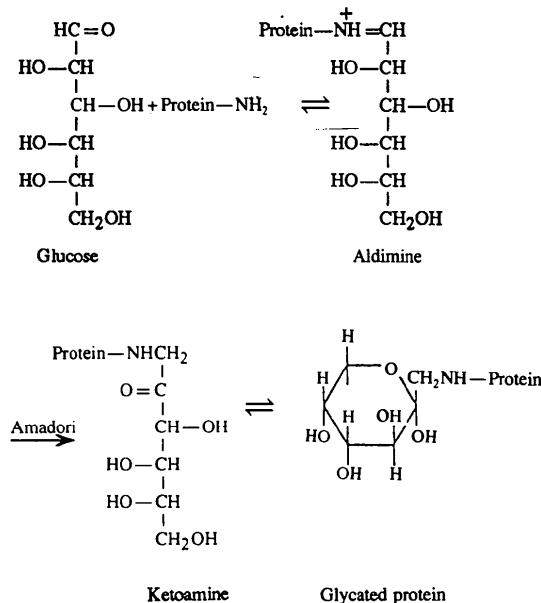
Comment

The title compound (1) is formed by a process like that occurring in the reaction of glucose with amino groups in proteins, which occurs non-enzymatically, *in vivo* (Cohen, 1986).



This intrinsically very slow process appears to be critical in the pathogenesis of various secondary complications associated with diabetes Mellitus (Cohen, 1986; Baynes, Thorpe & Murtiashaw, 1984; Lowrey, Lyness & Soeldner, 1985) and the process of aging (Monnier,

Kohn & Cerami, 1984). Glycation is initiated by the condensation of the acyclic form of glucose with protein amino groups to yield a Schiff-base intermediate (aldimine). This aldimine can undergo the practically irreversible Amadori rearrangement to a stable ketoamine derivative, which then cyclizes to the hemiketal structure (Neglia, Cohen, Garber, Thorpe & Baynes, 1985). Although the Schiff-base and Amadori compounds of small organic amines are known to assume cyclic pyranose and furanose conformations in solution (Funcke & Klemer, 1976), relatively little is known about the structures which exist in proteins either *in vivo* or *in vitro*. ^{13}C NMR spectroscopy has been used to characterize Amadori ketoamine adducts formed by the reaction of glucose with amino groups of proteins (Neglia, Cohen, Garber, Thorpe & Baynes, 1985) and with small molecules (Neglia, Cohen, Garber, Ellis, Thorpe & Baynes, 1983). The pyranose form appears to be the predominant structure in solution.



Glycated proteins have been reported to undergo either conformational changes (Shaklai, Garlick & Bunn, 1984) and/or functional changes (Watkins, Thorpe & Baynes, 1987; Cerami, Vlassara & Browlee, 1987). Non-enzymatic glycation of a large polypeptide at a single site can have marked effects on both the conformation as well as the biological properties of the protein. In order to have a much clearer understanding, we have prepared (1) and performed an X-ray crystallographic analysis which can serve as a structural model for the product in the general non-enzymatic processing of proteins. A diagram showing the molecular structure of C₁₃H₁₉NO₅ is shown in Fig. 1. Short intermolecular distances between a number of atom pairs indicate hydrogen-bonding interactions between these pairs (Table 2).