Lists of structure factors, anisotropic displacement parameters, Hatom coordinates and complete geometry have been deposited with the IUCr (Reference: FG1204). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

References

- Brussani, G., Ley, S. V., Wright, J. L. & Williams, D. J. (1986). J. Chem. Soc. Perkin Trans. 1, pp. 303-307.
- Dupont, L., Dideberg, O. & Jacquemin, P. (1990). Acta Cryst. C46, 484-486.
- Iwaoka, M. & Tomoda, S. (1994). J. Am. Chem. Soc. **116**, 4463–4464. Kienitz, C. O., Thöne, C. & Jones, P. G. (1996). *Inorg. Chem.* In the
- press. Mautner, H. G., Chu, S.-H. & Lee, C. M. (1962). J. Org. Chem. 27, 3671–3673.
- Sheldrick, G. M. (1990). Acta Cryst. A46, 467-473.
- Sheldrick, G. M. (1993). SHELXL93. Program for the Refinement of Crystal Structures. University of Göttingen, Germany.
- Siemens (1994a). XEMP. Empirical Absorption Correction Program. Siemens Analytical X-ray Instruments Inc., Madison, Wisconsin, USA.
- Siemens (1994b). XP. Molecular Graphics Program. Version 5.03. Siemens Analytical X-ray Instruments Inc., Madison, Wisconsin, USA.
- Stoe & Cie (1994a). DIF4. Diffractometer Control Program. Version 7.08. Stoe & Cie, Darmstadt, Germany.
- Stoe & Cie (1994b). REDU4. Data Reduction Program. Version 7.08. Stoe & Cie, Darmstadt, Germany.

Comment

Numerous routes to the indolizidine ring system have been reported (for a recent example, see Carretero & Arrayas, 1995, and references therein), largely stimulated by their extremely potent inhibition of glycosidases (Elbein, 1987). Our synthesis of the indolizidine ring system stems from our ability to perform conjugate additions to unsaturated nitriles (Fleming & Pak, 1995) that serves to assemble rapidly heterocyclic indolizidines, quinolizidines (Hussain, Fleming, Norman & Chang, 1996) and azulenes.

The X-ray structure of the title compound, (I), shows that the indolizidine ring system adopts the envelopechair orientation found in both the crystalline state (Koh, Lee, Sim & Zhu, 1993) and in solution (Reymond, Pinkerton & Vogel, 1991). The metric parameters of the rings are quite similar to the indolizidine 1-deoxycastanospermine (Koh, Lee, Sim & Zhu, 1993) and, in the present case, differ mainly in the shortening of the N(1)—C(9) bond [1.459 (4) versus 1.479 (5) Å] and lengthening of C(7)—C(8) [1.528 (4) versus 1.514 (5) Å]. A similar though less-pronounced trend is seen in the homologous quinolizine (Hussain, Fleming, Norman & Chang, 1996).



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(8*R*,8a*S*)-Indolizidine-1-spiro-2'-(1',3'dithiane)-8-carbonitrile

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Abstract

The title compound, $C_{12}H_{18}N_2S_2$, is an indolizidine whose structure was determined unequivocably by X-ray diffraction. The indolizidine ring adopts a *trans*-fused envelope-chair conformation while the dithiane ring adopts a chair conformation distal to the nitrile moiety.

© 1996 International Union of Crystallography Printed in Great Britain – all rights reserved The exact reason for these trends are unclear though we may explain this phenomenon by an interaction between the electron-rich N atom and the strongly electron-withdrawing nitrile group (Reddy, Goldstein &



Fig. I. Perspective drawing of (8R,8aS)-indolizidine-1-spiro-2'-(1',3'- dithiane)-8-carbonitrile with displacement ellipsoids drawn at the 50% probability level.

2849

Acta Crystallographica Section C ISSN 0108-2701 © 1996 Mandell, 1961). There is a good precedent for this type of interaction in α -cyanoamines as it is manifest in the facile elimination of cyanide (Yue, Royer & Husson, 1992), but prior difficulties in synthesizing β -cyanoamines may have precluded this effect from being observed previously.

Experimental

The synthesis of the title compound follows our earlier preparation (Hussain, Fleming, Norman & Chang, 1996) of the homologous octahydroquinolizine. Recrystallization from hot hexane afforded a crystalline material from which a single crystal was selected for diffraction.

Crystal data

$C_{12}H_{18}N_2S_2$	Mo $K\alpha$ radiation
$M_r = 254.41$	$\lambda = 0.7107 \text{ Å}$
Monoclinic	Cell parameters from 25
$P2_1/n$	reflections
a = 10.513(3) Å	$\theta = 16.81 - 20.50^{\circ}$
b = 11.182(2) Å	$\mu = 0.367 \text{ mm}^{-1}$
c = 11.206(2) Å	T = 294 K
$\beta = 97.18(2)^{\circ}$	Triangular pyramid
$V = 1307.0(5) \text{ Å}^3$	$0.35 \times 0.30 \times 0.30$ mm
Z = 4	White
$D_x = 1.293 \text{ Mg m}^{-3}$	
D_m not measured	

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Data collection
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AFC-7 <i>R</i> diffractometer	$R_{\rm int} = 0.044$
$\omega/2\theta$ scans	$\theta_{\rm max} = 32.50^{\circ}$
Absorption correction:	$h = 0 \rightarrow 14$
ψ scans (TEXSAN;	$k = 0 \rightarrow 15$
Molecular Structure	$l = -15 \rightarrow 15$
Corporation, 1992)	3 standard reflections
$T_{\rm min} = 0.83, \ T_{\rm max} = 0.90$	monitored every 150
4997 measured reflections	reflections
4787 independent reflections	intensity decay: -1.8%
2248 observed reflections	
$[I > 3.00\sigma(I)]$	

Refinement

Refinement on F $(\Delta/\sigma)_{\rm max} = 0.124$ $\Delta \rho_{\rm max} = 0.61 \ {\rm e} \ {\rm \AA}^{-3}$ R = 0.060 $\Delta \rho_{\rm min} = -0.55 \ {\rm e} \ {\rm \AA}^{-3}$ wR = 0.076S = 1.92Extinction correction: none 2248 reflections Atomic scattering factors 217 parameters from International Tables All H-atom parameters for X-ray Crystallography refined (1974, Vol. IV) $w = 4F^2/\sigma^2(F^2)$

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

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

	x	у	c	U_{eq}
S(1)	0.46482 (9)	0.06809 (8)	0.17378 (8)	0.0599 (3)
S(2)	0.3473(1)	0.30739(7)	0.21052 (7)	0.0576 (2)
N(1)	0.2379 (2)	0.1821 (2)	-0.0841 (2)	0.0453 (7)

N(2)	0.2944 (4)	0.5086 (3)	-0.0264(3)	0.072(1)
C(1)	0.3390 (3)	0.1726(2)	0.1189(2)	0.0413 (7)
C(2)	().2()42 (4)	0.1156 (4)	0.1047 (3)	0.059(1)
C(3)	0.1804 (4)	0.0818 (3)	-0.0276(3)	0.060(1)
C(5)	0.2497 (4)	0.1668 (4)	-0.2119 (3)	0.061(1)
C(6)	0.3143 (4)	().2742 (4)	-0.2582(3)	0.069(1)
C(7)	().4413 (4)	0.3024 (4)	-0.1835 (3)	0.066(1)
C(8)	0.4246(3)	0.3140(3)	-0.0484 (3)	0.0481 (9)
C(9)	0.3627 (3)	0.1978 (3)	-0.0129 (2)	0.0417 (7)
C(10)	0.3495 (3)	0.4214 (3)	-0.0323(3)	0.0510(9)
C(11)	0.4404 (5)	().0470 (4)	0.3295 (3)	0.064(1)
C(12)	0.4426 (4)	0.1601 (4)	0.4016 (3)	0.063(1)
C(13)	0.3342 (5)	0.2433 (4)	0.3574 (3)	0.064(1)

Table 2. Selected geometric parameters (Å, °)

S(1) - C(1)	1.814 (3)	C(1)—C(9)	1.555 (4)
S(1) - C(11)	1.810 (4)	C(2) - C(3)	1.520 (5)
S(2) - C(1)	1.820(3)	C(5)—C(6)	1.503 (6)
S(2) - C(13)	1.816 (3)	C(6)—C(7)	1.518 (6)
N(1) - C(3)	1.456 (4)	C(7)—C(8)	1.550(4)
N(1)—C(5)	1.463 (4)	C(8)—C(9)	1.528 (4)
N(1)—C(9)	1.459 (4)	C(8) - C(10)	1.462 (5)
N(2) - C(10)	1.140(4)	C(11) - C(12)	1.500 (6)
C(1)—C(2)	1.544 (5)	C(12)—C(13)	1.506 (6)
C(1)S(1)C(11)	103.0 (2)	N(1)-C(5)-C(6)	110.0 (3)
C(1) = S(2) = C(13)	100.4 (2)	C(5)—C(6)—C(7)	112.3 (3)
C(3) - N(1) - C(5)	115.2 (3)	C(6)—C(7)—C(8)	110.9 (3)
C(3) = N(1) = C(9)	104.4 (2)	C(7)C(8)C(9)	106.7 (3)
C(5) - N(1) - C(9)	111.6 (3)	C(7) - C(8) - C(10)	108.6 (3)
S(1) - C(1) - S(2)	111.2 (2)	C(9)—C(8)—C(10)	114.4 (3)
S(1) - C(1) - C(2)	113.0(2)	N(1)—C(9)—C(1)	104.9 (2)
S(1)C(1)C(9)	104.0 (2)	N(1)—C(9)—C(8)	110.1 (2)
S(2) - C(1) - C(2)	112.3 (2)	C(1)—C(9)—C(8)	121.7 (2)
S(2) - C(1) - C(9)	112.6 (2)	N(2) - C(10) - C(8)	175.2 (3)
C(2)—C(1)—C(9)	103.4 (2)	S(1) - C(11) - C(12)	114.5 (3)
C(1) - C(2) - C(3)	103.9 (3)	C(11)—C(12)—C(13)	112.6(3)
N(1) - C(3) - C(2)	101.9 (3)	S(2)—C(13)—C(12)	113.7 (3)

The C—H bonds range from 0.89 (3) to 1.07 (4) Å and the Hatom displacement parameters from 0.043 (8) to 0.10 (1) Å².

Data collection: *MSC/AFC Diffractometer Control Software* (Molecular Structure Corporation, 1988). Cell refinement: *MSC/AFC Diffractometer Control Software*. Data reduction: *TEXSAN* (Molecular Structure Corporation, 1992). Program(s) used to solve structure: *SIR*92 (Altomare *et al.*, 1994). Program(s) used to refine structure: *TEXSAN*. Software used to prepare material for publication: *TEXSAN*.

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Lists of structure factors, anisotropic displacement parameters, Hatom coordinates and complete geometry have been deposited with the IUCr (Reference: BK1249). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

References

- Altomare, A., Cascarano, M., Giacovazzo, C., Guagliardi, A., Burla, M. C., Polidori, G. & Camalli, M. (1994). J. Appl. Cryst. 27, 435– 436.
- Carretero, J. C. & Arrayas, R. G. (1995). J. Org. Chem. 60, 6000-6001.
- Elbein, A. D. (1987). Ann. Rev. Biochem. 56, 497-534.
- Fleming, F. F. & Pak. J. J. (1995). J. Org. Chem. 60, 4299-4301.
- Hussain, Z., Fleming, F. F., Norman, R. E. & Chang, S.-C. (1996). Acta Cryst. C52, 1296–1298.
- Koh, L.L, Lee, C. K., Sim, K. Y. & Zhu, J. (1993). Acta Cryst. C49, 391–392.

- Molecular Structure Corporation (1988). MSC/AFC Diffractometer Control Software. MSC, 3200 Research Forest Drive, The Woodlands, TX 77381, USA.
- Molecular Structure Corporation (1992). TEXSAN. Crystal Structure Analysis Package. MSC, 3200 Research Forest Drive, The Woodlands, TX 77381, USA.
- Reddy, G. S., Goldstein, J. H. & Mandell, L. (1961). J. Am. Chem. Soc. 83, 1300-1306.
- Reymond, J.-L., Pinkerton, A. A. & Vogel, P. (1991). J. Org. Chem. 56, 2128–2135.
- Yue, C., Royer, J. & Husson, H.-P. (1992). J. Org. Chem. 57, 4211-4214.



9-Dicyclohexylphenylphosphino-*arachno*-6-thiadecaborane(11)

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Abstract

The {SB₉} cage residue in the title compound, C₁₈H₃₈B₉PS, has the expected *arachno* ten-vertex geometry and the PCy₂Ph substituent occupies an *exo* position on B9, with B9—P1 = 1.953 (6) Å.

Comment

There is current interest in bis(phosphine)rhodathiaboranes whose molecular structures appear to be at variance with those expected by electron-counting rules (Ferguson et al., 1990; Murphy, Spalding, Ferguson & Gallagher, 1992) unless intramolecular agostic interactions are invoked (Adams, McGrath & Welch, 1995; Adams, McGrath, Thomas, Weller & Welch, 1996). As part of our studies in this area, we attempted the synthesis of $8,8-(Cy_2PhP)_2-8,7-nido-RhSB_9H_{10}$ (Cy = cyclohexyl) by reaction between $[RhCl(PCy_2Ph)_2(C_2H_4)]$, generated in situ from $[RhCl(C_2H_4)_2]_2$ and PCy_2Ph , and $Cs[SB_9H_{12}]$ in Et₂O. However, the major tractable product afforded by work-up proved to be the title compound. Since there is current interest in 9-substituted arachno-6-SB₉H₁₁ species (Stibr et al., 1996) and since 9-PPh₃-arachno-6-SB₉H₁₁ has been variously described (by the same workers) as endo-9- (Nestor, Fontaine, Greenwood, Kennedy & Thornton-Pett, 1991) and exo-9- (Stibr et al., 1996), we undertook a crystallographic study of the title compound.



The compound crystallizes with no short intermolecular contacts. A perspective view of a single molecule is shown in Fig. 1.



Fig. 1. Perspective view with 40% ellipsoids for non-H atoms. Ring C atoms are numbered in sequence.

This is the first crystallographic characterization of a 9-phosphino-6-SB₉H₁₁ species. The {SB₉} residue has the same basic *arachno* structure as that found in 9-NEt₃-*arachno*-6-SB₉H₁₁ (Hilty & Rudolf, 1979), [*arachno*-6-SB₉H₁₂]⁻ (Nestor *et al.*, 1991) and 9-MeCN-*arachno*-6-SB₉H₁₁ (Stibr *et al.*, 1996). Similar to the situation in all these three analogous compounds, the B—B distances in the title compound appear to fall into three fairly distinct groups: *ca* 1.90 Å for B2—B5, B2— B7, B8—B9 and B9—B10; *ca* 1.85 Å for the hydrogenbridged edges B5—B10 and B7—B8; *ca* 1.75–1.80 Å for all others. S—B distances in the title compound are comparable with those in the thiaboranes referenced above, but more spread than usual with S6—B2 being just significantly longer than S6—B5 and S6—B7.

Overall, the thiaborane cage has approximate C_s symmetry about the plane through S6, B2, B4 and B9, and the ¹¹B NMR spectrum is fully consistent with such symmetry in solution. However, it is clear from Fig. 1 that the PCy₂Ph substituent at B9 [which is clearly in an *exo* position, B9—P1 1.953 (6) Å] is not oriented so as to maintain overall C_s symmetry in the solid state. We assume that in solution at room temperature, the PCy₂Ph ligand is free to rotate about the B9—P1 bond.

Experimental

Under an atmosphere of dry nitrogen, a solution of $[RhCl(PCy_2Ph)_2(C_2H_4)]$ in Et₂O was prepared *in situ* by slow

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