Acta Crystallographica Section E

Structure Reports Online

ISSN 1600-5368

2-Diphenylarsino-1-(diphenylphosphino)ethane

Omar bin Shawkataly, a* Mei-Lin Chong, a Hoong-Kun Fun, b Claude Didierjean and André Aubry

^aChemical Sciences Programme, School of Distance Education, Universiti Sains Malaysia, Minden,11800, USM, Penang, Malaysia, ^bX-Ray Crystallography Unit, School of Physics, Universiti Sains Malaysia, Minden,11800, USM, Penang, Malaysia, and ^cLaboratoire de Cristallographie et Modélisation des Matériaux Minéraux, et Biologiques (LCM3B), UMR n° 7036, Université Henri Poincaré, Nancy I, Faculté des Sciences, BP 239, 54506 Vandoeuvre lès Nancy Cedex, France

Correspondence e-mail: omarsa@usm.my

Key indicators

Single-crystal X-ray study T = 293 KMean $\sigma(\text{C-C}) = 0.005 \text{ Å}$ Disorder in main residue R factor = 0.039 wR factor = 0.096 Data-to-parameter ratio = 18.9

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

The title compound, $C_{26}H_{24}PAs$, has inversion symmetry as a result of the orientational disorder of the molecule. The two possible positions of the As and P atoms are overlapped.

Received 14 September 2005 Accepted 16 September 2005 Online 21 September 2005

Comment

There have been numerous X-ray structural determinations of tertiary phosphines and also tertiary arsines which are widely used as ligands in organo-transition metal chemistry. However, X-ray structure determinations of ligands containing both P and As atoms are rare. As part of our study on the substitution of transition metal–carbonyl clusters with mixed-ligand complexes, we have published several structures of triruthenium–carbonyl clusters containing mixed P/As (Shawkataly *et al.*, 1998) or P/Sb ligands (Shawkataly *et al.*, 2004). A search of the November 2004 release of the Cambridge Structural Database (Allen, 2002) revealed only 20 structures of complexes containing the above ligands.

The title compound, (I), is commercially available and has been widely used, though the crystal structure has not been reported.

The molecular structure of (I), showing the atom-labeling scheme. Displacement ellipsoids are drawn at the 25% probability level. Unlabeled atoms are related to labeled atoms by (2 - x, -y, 2 - z).

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organic papers

The crystal structure of (I) is isomorphous with those of $Ph_2P(CH_2)_2PPh_2$ (Pelizzi & Pelizzi, 1979) and $Ph_2As(CH_2)_2AsPh_2$ (Hill *et al.*, 2001); all these compounds have a crystallographic inversion center at the mid-point of the Csp^3-Csp^3 bond. Positional disorder of the As and P atoms (Fig. 1). The positional disorder between the As/P atoms was also observed in $Ru_3(CO)_8(Ph_2AsCH_2AsPh_2)-(Ph_2PCH_2PPh_2)$ (Shawkataly *et al.*, 1998).

Experimental

The compound was supplied by Strem Chemicals. Single crystals of (I) were obtained by slow evaporation of an ethanol solution.

Crystal data

$C_{26}H_{24}AsP$	$D_x = 1.323 \text{ Mg m}^{-3}$	
$M_r = 442.34$	Mo $K\alpha$ radiation	
Monoclinic, $P2_{1}/n$	Cell parameters from 40	
a = 9.1011 (6) Å	reflections	
b = 5.7856 (8) Å	$\theta = 9.9 - 24.3^{\circ}$	
c = 21.4502 (10) Å	$\mu = 1.61 \text{ mm}^{-1}$	
$\beta = 100.557 (5)^{\circ}$	T = 293 (2) K	
$V = 1110.35 (18) \text{ Å}^3$	Prism, colorless	
Z = 2	$0.22 \times 0.16 \times 0.12 \text{ mm}$	

Data collection

$R_{\rm int} = 0.034$		
$\theta_{\text{max}} = 27.5^{\circ}$		
$h = -1 \rightarrow 11$		
$k = -1 \rightarrow 7$		
$t = -27 \rightarrow 27$		
3 standard reflections		
frequency: 60 min		
intensity decay: 0.2%		

Refinement

Refinement on F^2	
$R[F^2 > 2\sigma(F^2)] = 0.039$	
$wR(F^2) = 0.096$	
S = 0.95	
2520 reflections	
133 parameters	

H atoms treated by a mixture of independent and constrained refinement $w = 1/[\sigma^2(F_o^2) + (0.041P)^2]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\rm max} < 0.001$ $\Delta\rho_{\rm max} = 0.29$ e Å $^{-3}$

 $\Delta \rho_{\min} = -0.34 \text{ e Å}^{-3}$

Table 1 Selected geometric parameters (Å, °).

As/P-C13	1.910(3)	As/P-C7	1.915 (2)
As/P-C1	1.912 (3)	$C13 - C13^{i}$	1.520 (5)
C13-As/P-C1	100.51 (13)	C6-C1-As/P	115.8 (2)
C13-As/P-C7	98.01 (11)	C8-C7-C12	118.2 (2)
C1-As/P-C7	97.72 (10)	C8-C7-As/P	123.9 (2)
C2-C1-C6	118.7 (3)	C12-C7-As/P	117.9 (2)
C2-C1-As/P	125.3 (2)	$C13^{i} - C13 - As/P$	110.3 (3)

Symmetry code: (i) -x + 2, -y, -z + 2.

The site-occupancy factors of the As and P atoms were fixed at 0.5, and the same positional and atomic displacement parameters were assumed. The methylene H atoms were located in difference density maps and their coordinates were refined [C-H = 0.93 (3)-0.95 (3) Å]. The phenyl H atoms were placed at calculated positions and refined using a riding model, with C-H = 0.93 Å. All H-atom $U_{\rm iso}$ parameters were fixed at $1.2U_{\rm eq}(C)$.

Data collection: XSCANS (Siemens, 1994); cell refinement: XSCANS; data reduction: XSCANS; program(s) used to solve structure: SHELXS86 (Sheldrick, 1985); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP3 for Windows (Farrugia, 1997); software used to prepare material for publication: WinGX publication routines (Farrugia, 1999).

The authors thank the Malaysian Government and Universiti Sains Malaysia for support through IRPA grant Nos. 09–02-05–0008 and 190–9609-2801.

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