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Triorganophosphine-dithiomonometaphosphoryl halides

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The title compounds, ethyldiphenylphosphine–dithiomonometaphosphoryl chloride, EtPh₂P→PS₂Cl, C₁₄H₁₅ClP₂S₂, (I), and tris-*n*-propylphosphine–dithiomonometaphosphoryl chloride and bromide, "Pr₃P→PS₂Cl, C₉H₂₁ClP₂S₂, (II), and "Pr₃P→PS₂Br, C₉H₂₁BrP₂S₂, (III), respectively, are the first phosphine-stabilized dithiomonometaphosphoryl halides to be structurally characterized. In the tris-*n*-propylphosphine derivatives, the central P→P donor–acceptor bond becomes longer in the order bromo < chloro < fluoro. Substitution of the tris-*n*-propylphosphine group in (II) by the more bulky ethyldiphenylphosphine group also leads to a longer P→P bond. These structural features agree with the observed ³¹P NMR data. In (II) and (III), the central P–P bond coincides with the crystallographic threefold axis, entailing site-occupational disorder for the S₂Y group.

Comment

The first inner organylphosphonium dithiomonometaphosphonic salts were obtained by the reaction of perthioarylanhydrides, $(RPS_2)_2$, phosphonic with tris-*n*-butylphosphine (Fluck & Binder, 1967). The authors proposed a betaine structure for these compounds because of two doublets visible in the ³¹P NMR spectra. The observed doublets represent an AB spin system (see Scheme below), which should be typical for a betaine structure. At the same time, the first pyridine-stabilized dithiomonometaphosphoryl chloride, $py \rightarrow PS_2Cl$, was isolated (Meisel & Grunze, 1968). This compound was synthesized by nucleophilic degradation of the adamantane cage, P_4S_{10} , by pyridine in the presence of thiophosphoryl chloride. Initially, $py \rightarrow PS_2Cl$ was also described as a betaine. Later on, owing to the bonding data (especially the $N \rightarrow P$ bond distance) obtained from a crystal structure determination (Averbuch-Pouchot & Meisel, 1989), a donoracceptor bond between the pyridine N atom and the PS₂Cl group was formulated. The relevant $N \rightarrow P$ bond length of

1.849 (6) Å was the longest known at that time. The proposed bonding model corresponds with the thermal behaviour of this type of compound, which has been shown by photoelectron spectroscopic experiments on $py \rightarrow PX_2Y$ (X is O or S and Y is F or Cl; Meisel *et al.*, 1989; Bock *et al.*, 1992).

The present work is aimed at investigating the bonding situation of some analogous phosphine adducts, which are the two chlorides, ethyldiphenylphosphine–dithiomonometa-phosphoryl chloride, (I), and tris-*n*-propylphosphine–dithiomonometaphosphoryl chloride, (II), and the bromide, tris-*n*-propylphosphine–dithiomonometaphosphoryl bromide, (III). These compounds provide the first structure examples of phosphine-stabilized dithiomonometaphosphoryl derivatives.



Compound (I) crystallizes in a monoclinic space group, forming a well ordered structure. Compounds (II) and (III) both crystallize in a trigonal space group and are isotypic with one another. Because the $P \rightarrow P$ bond of the molecules in (II) and (III) lies on a threefold axis, the $S_2Y(Y \text{ is Cl or Br})$ ligands must be disordered. The same has been observed for the analogous fluoride compound, (IV), although the results for this compound are unpublished, because of another disorder problem restricting the results of additional structure determination (Rabis, 1999). In the tris-*n*-propylphosphine derivatives, the length of the $P \rightarrow P$ donor-acceptor bond is dependent on the halogen ligand and increases from Br *via* Cl



Figure 1

A view of the molecular structure of (I). Displacement ellipsoids are drawn at the 50% probability level and H atoms have been omitted for clarity.

to F, *i.e.* from 2.2035 (16) *via* 2.2253 (10) to 2.251 (4) Å. These experimental results correlate with the π -donating properties of the halides, which increase from Br to F.

The ³¹P NMR spectra of (I)–(III) reflect the bonding situation in the magnitudes of the indirect spin–spin coupling constant, ${}^{1}J_{PP}$. Usually, it is assumed that the shorter the bond length, the larger the J value. Our experiments show that the absolute magnitudes of ${}^{1}J_{PP}$ decrease from Br⁻ to F⁻. In the case of phosphorus compounds, the coupling constant ${}^{1}J_{PP}$ depends primarily on the valency of the P atoms involved and is related to the absence or presence of lone pairs. Thus, for Me₂P–PMe₂, ${}^{1}J_{PP}$ is –180 Hz, whereas for the corresponding disulfide, a value of 19 Hz has been found (Harris, 1983). This, in connection with the X-ray results, gives rise to the supposition that, in the present case, ${}^{1}J_{PP}$ has a positive sign.



Figure 2

A view of the molecular structure of (II). Displacement ellipsoids are drawn at the 50% probability level and H atoms have been omitted for clarity. Only one disordered component is shown, and the symmetry codes generate the three crystallographically equivalent positions for the atoms of the S₂Y group [symmetry codes: (i) 1 - y, x - y, z; (ii) 1 - x + y, 1 - x, z].



Figure 3

A view of the molecular structure of (III). Displacement ellipsoids are drawn at the 50% probability level and H atoms have been omitted for clarity. Only one disordered component is shown, and the symmetry codes generate the three crystallographically equivalent positions for the atoms of the S₂Y group [symmetry codes: (i) 1 - y, x - y, z; (ii) 1 - x + y, 1 - x, z].

A comparison of the structural data of the two chloride compounds, (I) and (II), shows that the $P \rightarrow P$ bond is shorter in the tris-*n*-propylphosphine adduct, (II) [2.253 (10) Å], than in the ethyldiphenylphosphine adduct, (I) [2.2522 (5) Å]. This can be explained by the higher steric demand of the ethyldiphenylphosphine group compared with the tris-*n*-propylphosphine group, which is also in agreement with the corresponding Tolman cone angles of 140 and 132°, respectively (Tolman, 1977).

Experimental

Using a method analogous to that of Fluck et al. (1973), a solution of tris-n-organylphosphine (9.5 mmol) was added dropwise to a suspension of $py \rightarrow PS_2Y$ (9.5 mmol) (Y is Br, Cl or F) in benzene with stirring. The product was filtered off after 1 h and washed with benzene. Yields: 81% for (I), 67% for (II), 78% for (III), 96% for Y =F, i.e. (IV). The ³¹P NMR solution spectra of both chlorides and of the bromide showed the two doublets characteristic for AB spin systems, as follows: for (I): 9.2 p.p.m. (d), 71.9 p.p.m. (d), ${}^{1}J_{PP} = 59.3$ Hz; for (II): 18.8 p.p.m. (d), 69.7 p.p.m. (d), ${}^{1}J_{PP} = 41.5$ Hz; for (III): 17.8 p.p.m. (d), 45.7 p.p.m. (d), ${}^{1}J_{PP} = 42.0$ Hz. In the ${}^{31}P$ NMR solution spectrum of $Pr_3P \rightarrow PS_2F$, (IV), there are two doublets of doublets because of the additional coupling to the ¹⁹F nucleus: 11.8 p.p.m. with ${}^{2}J_{PF} = 117.8$ Hz, and 102.2 p.p.m. with ${}^{1}J_{PF} =$ 1111.2 Hz and ${}^{1}J_{PP}$ = 11.4 Hz. Crystals of all four compounds suitable for X-ray analysis were obtained by recrystallization from solutions in benzene/acetonitrile (10:1).

Compound (I)

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Crystal data C14H15ClP2S2 $D_x = 1.472 \text{ Mg m}^{-3}$ $M_r = 344.77$ Mo $K\alpha$ radiation Cell parameters from 5000 Monoclinic, $P2_1/n$ a = 13.204(1) Å reflections b = 9.694(1) Å $\theta = 6.9 - 28.3^{\circ}$ $\mu = 0.70 \text{ mm}^{-1}$ c = 13.655(1) Å T = 190 (2) K $\beta = 117.11 \ (1)^{\circ}$ V = 1555.8 (3) Å³ Prismatic, colourless Z = 40.57 \times 0.38 \times 0.19 mm

Data collection

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toe IPDS diffractometer	3738 independent reflections
scans	3485 reflections with $I > 2\sigma(I)$
Absorption correction: refined from	$R_{\rm int} = 0.043$
ΔF^2 (ABSCOR; Stoe & Cie,	$\theta_{\rm max} = 28.1^{\circ}$
1996)	$h = -17 \rightarrow 15$
$T_{\min} = 0.659, T_{\max} = 0.875$	$k = -12 \rightarrow 12$
8 747 measured reflections	$l = -17 \rightarrow 17$

Table 1

Selected geometric parameters (Å, °) for (I).

P1-C12 P1-C13 P1-C11 P1-P2	1.7958 (13) 1.8023 (12) 1.8090 (13) 2.2522 (5)	P2-S12 P2-S22 P2-Cl2	1.9399 (5) 1.9454 (5) 2.0724 (5)
C12 - P1 - C13 $C12 - P1 - C11$ $C13 - P1 - C11$ $C12 - P1 - P2$ $C13 - P1 - P2$ $C13 - P1 - P2$ $C11 - P1 - P2$	109.64 (6) 110.54 (6) 109.31 (6) 110.95 (4) 109.08 (4) 107.27 (5)	S12-P2-S22 S12-P2-C12 S22-P2-C12 S12-P2-P1 S22-P2-P1 C12-P2-P1	122.76 (2) 110.64 (2) 109.07 (2) 108.87 (2) 105.08 (2) 97.25 (2)

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.028$ $wR(F^2) = 0.081$ S = 1.073738 reflections 172 parameters H-atom parameters constrained

Compound (II)

Crystal data

$C_9H_{21}ClP_2S_2$ M = 200.77	Mo $K\alpha$ radiation	Data collection	
$M_r = 290.77$ Trigonal, <i>P</i> 3c1 <i>a</i> = 11.413 (3) Å <i>b</i> = 11.413 (4) Å <i>c</i> = 13.566 (3) Å <i>V</i> = 1530.3 (8) Å ³ <i>Z</i> = 4 <i>D_x</i> = 1.262 Mg m ⁻³	reflections $\theta = 2.5-25.0^{\circ}$ $\mu = 0.70 \text{ mm}^{-1}$ T = 180 (2) K Block, colourless $0.68 \times 0.40 \times 0.28 \text{ mm}$	Stoe IPDS diffractometer φ scans Absorption correction: numerical (<i>XRED</i> ; Stoe & Cie, 1996) $T_{min} = 0.637, T_{max} = 0.707$ 9325 measured reflections 1024 independent reflections	722 reflec $R_{int} = 0.1$ $\theta_{max} = 26$ h = -14 k = -14 l = -16
-			

 $w = 1/[\sigma^2(F_o^2) + (0.0507P)^2]$

+ 0.4476P] where $P = (F_o^2 + 2F_c^2)/3$

 $(\Delta/\sigma)_{\rm max} = 0.002$ $\Delta \rho_{\rm max} = 0.41 \ {\rm e} \ {\rm \mathring{A}}^{-3}$

 $\Delta \rho_{\rm min} = -0.37 \ {\rm e} \ {\rm \AA}^{-3}$

Data collection

Stoe IPDS diffractometer	10 448 measured reflections
φ scans	1001 independent reflections
Absorption correction: numerical	987 reflections with $I > 2\sigma(I)$
[XRED (Stoe & Cie, 1996) and	$R_{\rm int} = 0.056$
XSHAPE (Stoe & Cie, 1996),	$\theta_{\rm max} = 26^{\circ}$
based on HABITUS (Herrendorf,	$h = -13 \rightarrow 14$
1993)]	$k = -14 \rightarrow 13$
$T_{\min} = 0.777, T_{\max} = 0.848$	$l = -16 \rightarrow 16$

Refinement

Refinement on F^2 $w = 1/[\sigma^2(F_o^2) + (0.0313P)^2]$ $R[F^2 > 2\sigma(F^2)] = 0.022$ + 0.2298P] $wR(F^2) = 0.054$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\rm max} = 0.002$ S = 1.10-3 $\Delta \rho_{\rm max} = 0.30 \ {\rm e} \ {\rm \AA}$ 1000 reflections $\Delta \rho_{\rm min} = -0.18 \ {\rm e} \ {\rm \AA}^{-3}$ 48 parameters H-atom parameters constrained

Table 2

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Selected	geometric	parameters	(Å.	°) for	(II)
	8	r	(,	()

C1-P1	1.805 (2)	P2-S1	1.964 (6)
P1-P2	2.2253 (10)	P2-Cl1	2.026 (12)
$C1 - P1 - C1^{1}$	108.84(5)	S1-P2-Cl1 ¹	114.8 (8)
C1-P1-P2	110.10 (5)	Cl1-P2-Cl1 ⁱ	116.1 (3)
S1-P2-S1i	112.2 (2)	S1-P2-P1	106.6 (2)
S1 ⁱ -P2-Cl1	113.9 (8)	Cl1-P2-P1	101.6 (4)

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

Compound (III)

Crystal data

C₉H₂₁BrP₂S₂ $M_r = 335.23$ Trigonal, P3c1 a = 11.522 (3) Å b = 11.522 (3) Å c = 13.450 (4) Å $V = 1546.3 (7) \text{ Å}^3$ Z = 4 $D_x = 1.440 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation Cell parameters from 1524 reflections $\theta = 2.5 - 25.0^{\circ}$ $\mu = 3.10 \text{ mm}^{-1}$ T = 180 (2) KTrigonal prism, colourless $0.16 \times 0.15 \times 0.12$ mm

Table 3

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Selected geometric parameters (Å, °) for (III).

C1-P1	1.801 (2)	P1-P2	2.2035 (16)
Br1-P2	2.128 (6)	P2-S1	1.954 (8)
$C1^{i}-P1-C1$	109.38 (8)	$S1^{i}-P2-Br1$	113.9 (12)
C1-P1-P2	109.57 (8)	$Br1^{i}-P2-Br1$	114.5 (4)
$S1^{i}-P2-S1$	113.1 (6)	S1-P2-P1	105.5 (7)
$S1-P2-Br1^{i}$	113.8 (12)	Br1-P2-P1	103.8 (5)

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

Stoe IPDS diffractometer	722 reflections with $I > 2\sigma(I)$
φ scans	$R_{\rm int} = 0.109$
Absorption correction: numerical	$\theta_{\rm max} = 26.2^{\circ}$
(XRED; Stoe & Cie, 1996)	$h = -14 \rightarrow 13$
$T_{\min} = 0.637, T_{\max} = 0.707$	$k = -14 \rightarrow 14$
9325 measured reflections	$l = -16 \rightarrow 16$
1024 independent reflections	
Refinement	
Refinement on F^2	H-atom parameters constraine
$R[F^2 > 2\sigma(F^2)] = 0.033$	$w = 1/[\sigma^2(F^2) + (0.0275P)^2]$
R[1 > 20(1)] = 0.055	$w = 1/[0 (1_0) + (0.02751)]$

Rennement on F	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.033$	$w = 1/[\sigma^2(F_o^2) + (0.0275P)^2]$
$wR(F^2) = 0.061$	where $P = (F_o^2 + 2F_c^2)/3$
S = 0.89	$(\Delta/\sigma)_{\rm max} = 0.004$
1024 reflections	$\Delta \rho_{\rm max} = 0.36 \text{ e } \text{\AA}^{-3}$
47 parameters	$\Delta \rho_{\rm min} = -0.45 \text{ e } \text{\AA}^{-3}$

All checked crystals of (II) were merohedrally twinned to [001]. In the crystal analysed here, the ratio of the two twin domains was 0.522 (2):0.478 (2). Disorder was observed in structures (II) and (III), so that two S atoms and one Cl atom in (II), and two S atoms and one Br atom in (III) partially occupy sites which are equivalent by threefold symmetry. However, the distances between the S-atom and halide sites are less than 0.20 Å in each structure. Therefore, the site separation is much lower than the atomic resolution achievable with the current data sets. Hence, the P-S and P-Cl distances were restrained to target values taken from the well ordered structure, (I) (Table 1), and the P–Br distance was restrained to 2.14 (2) Å (Vogt et al., 1993) throughout all the least-squares cycles. The U_{ii} values for the disordered atoms of (II) and (III) were constrained to be equal. For (I), a ΔF^2 -based absorption correction was applied, using ABSCOR (Stoe & Cie, 1996), a modification of DIFABS (Walker & Stuart, 1983). In contrast with DIFABS, ABSCOR loads F^2 values instead of F. For (II), a numerical absorption correction using a Gaussian algorithm was applied, using XRED (Stoe & Cie, 1996). Before the actual absorption correction, the program XSHAPE (Stoe & Cie, 1996) was applied to optimize the predetermined crystal shape. 1462 medium-strong symmetry-equivalent reflections, of which 150 were unique, were used for the optimization. XSHAPE is based on the program HABITUS (Herrendorf, 1993). For (III), all crystal faces were indexed by microscope on the diffractometer, and thereafter a Gaussian integration numerical absorption correction was applied. All H atoms were placed in geometrically idealized positions, with C-H = 0.95-0.99 Å, and constrained to ride on their parent atoms, with $U_{iso}(H) = 1.2U_{eq}(C)$. Additionally, the methyl H atoms were allowed to rotate about the C-C bonds.

For all three compounds, data collection: IPDS (Stoe & Cie, 1996); cell refinement: IPDS; data reduction: IPDS. For compounds (I) and (III), program(s) used to solve structure: SHELXS97 (Sheldrick, 1997). For compound (II), program(s) used to solve structure:

organic compounds

*SHELXS*86 (Sheldrick, 1990). For all three compounds, program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *DIAMOND* (Brandenburg, 1996); software used to prepare material for publication: *SHELXL*97.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: LN1136). Services for accessing these data are described at the back of the journal.

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