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## Triclinic polymorphs of triphenylphosphine and triphenylphosphine sulfide

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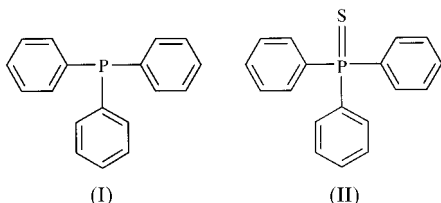
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New triclinic polymorphs of triphenylphosphine, (I), and triphenylphosphine sulfide, (II), were crystallized. For the structure of (I), it is obvious that the highest peaks in the final difference electron-density map are located at the 'lone-pair regions' of P atoms. We have observed this effect in two further phosphorous-containing structures.

### Comment

By reaction of triphenylphosphine, (I), with pyridinedithiomonometaphosphoryl chloride, an insertion of sulfur into triphenylphosphine takes place and a new triclinic polymorph of triphenylphosphine sulfide, (II), was isolated. The analogous reaction of triphenylphosphine with pyridinedithiomonometaphosphoryl fluoride yields in addition to the abovementioned sulfide a new crystallographic modification of triclinic triphenylphosphine. Both compounds form monoclinic structures, too, which have been described previously by Howells *et al.* (1954) and Codding & Kerr (1978), respectively.



In the unit cell of triphenylphosphine there are four independent molecules. Here, it is obvious that the four highest peaks in the final difference electron-density map [ $1.08(4)$ ,  $0.69(4)$ ,  $0.64(4)$  and  $0.31(4)$  e Å<sup>-3</sup>] are observed at the 'lone-pair regions' of P1–P4. The distances between those peaks and the corresponding P atoms are 1.25, 1.35, 1.35 and 1.22 Å, respectively.

### Experimental

A solution of triphenylphosphine in benzene was slowly dropped to a suspension of pyridinedithiomonometaphosphoryl chloride or fluoride in benzene. After 1 h, the solid products were filtrated and recrystallized from benzene.

### Compound (I)

#### Crystal data

C<sub>18</sub>H<sub>15</sub>P  
*M<sub>r</sub>* = 262.27  
Triclinic, *P* $\bar{1}$   
*a* = 11.0306 (19) Å  
*b* = 15.111 (3) Å  
*c* = 17.554 (3) Å  
 $\alpha$  = 84.38 (2)°  
 $\beta$  = 80.36 (2)°  
 $\gamma$  = 85.82 (2)°  
*V* = 2866.0 (8) Å<sup>3</sup>

*Z* = 8  
*D<sub>x</sub>* = 1.216 Mg m<sup>-3</sup>  
Mo *K*α radiation  
Cell parameters from 5000 reflections  
 $\theta$  = 2.5–25.0°  
 $\mu$  = 0.175 mm<sup>-1</sup>  
*T* = 180 (2) K  
Block, colorless  
0.48 × 0.48 × 0.28 mm

#### Data collection

Stoe IPDS diffractometer  
 $\varphi$ -oscill.,  $\varphi$ -incr. = 1.4°, 164 exposure  
Absorption correction: reldelf see publ section exptl prep  
*T*<sub>min</sub> = 0.921, *T*<sub>max</sub> = 0.953  
17 345 measured reflections  
9738 independent reflections

6818 reflections with *I* > 2σ(*I*)  
*R*<sub>int</sub> = 0.0324  
 $\theta$ <sub>max</sub> = 25.25°  
*h* = -13 → 13  
*k* = -18 → 18  
*l* = -20 → 20

#### Refinement

Refinement on *F*<sup>2</sup>  
*R* [*F*<sup>2</sup> > 2σ(*F*<sup>2</sup>)] = 0.0422  
*wR* (*F*<sup>2</sup>) = 0.1114  
*S* = 0.919  
9738 reflections  
685 parameters

H-atom parameters constrained  
*w* = 1/[σ<sup>2</sup>(*F<sub>o</sub>*<sup>2</sup>) + (0.0683*P*)<sup>2</sup>]  
where *P* = (*F<sub>o</sub>*<sup>2</sup> + 2*F<sub>c</sub>*<sup>2</sup>)/3  
(Δ/σ)<sub>max</sub> = 0.004  
Δρ<sub>max</sub> = 1.02 e Å<sup>-3</sup>  
Δρ<sub>min</sub> = -0.23 e Å<sup>-3</sup>

### Compound (II)

#### Crystal data

C<sub>18</sub>H<sub>15</sub>PS  
*M<sub>r</sub>* = 294.33  
Triclinic, *P* $\bar{1}$   
*a* = 9.542 (3) Å  
*b* = 11.207 (3) Å  
*c* = 14.537 (4) Å  
 $\alpha$  = 88.89 (4)°  
 $\beta$  = 80.39 (4)°  
 $\gamma$  = 78.28 (3)°  
*V* = 1500.5 (8) Å<sup>3</sup>

*Z* = 4  
*D<sub>x</sub>* = 1.303 Mg m<sup>-3</sup>  
Mo *K*α radiation  
Cell parameters from 5000 reflections  
 $\theta$  = 2.5–25.0°  
 $\mu$  = 0.309 mm<sup>-1</sup>  
*T* = 180 (2) K  
Prism, yellow  
0.72 × 0.38 × 0.11 mm

#### Data collection

Stoe IPDS diffractometer  
 $\varphi$ -oscill.,  $\varphi$ -incr. = 1.5°, 160 exposures  
Absorption correction: reldelf see publ section exptl prep  
*T*<sub>min</sub> = 0.808, *T*<sub>max</sub> = 0.967  
13 389 measured reflections

5111 independent reflections  
4117 reflections with *I* > 2σ(*I*)  
*R*<sub>int</sub> = 0.0502  
 $\theta$ <sub>max</sub> = 25.24°  
*h* = -11 → 11  
*k* = -13 → 13  
*l* = -17 → 17

#### Refinement

Refinement on *F*<sup>2</sup>  
*R* [*F*<sup>2</sup> > 2σ(*F*<sup>2</sup>)] = 0.0293  
*wR* (*F*<sup>2</sup>) = 0.0786  
*S* = 0.974  
5111 reflections  
481 parameters

All H-atom parameters refined  
*w* = 1/[σ<sup>2</sup>(*F<sub>o</sub>*<sup>2</sup>) + (0.0500*P*)<sup>2</sup>]  
where *P* = (*F<sub>o</sub>*<sup>2</sup> + 2*F<sub>c</sub>*<sup>2</sup>)/3  
Δρ<sub>max</sub> = 0.30 e Å<sup>-3</sup>  
Δρ<sub>min</sub> = -0.29 e Å<sup>-3</sup>

For the studied crystals, a Δ*F*<sup>2</sup>-based absorption correction was carried out. The calculation was performed with *ABSCOR* (Stoe & Cie, 1997), a modification of *DIFABS* (Walker & Stuart, 1983). In

contrary of *DIFABS*, *ABSCOR* loads and provides  $F^2$  values instead of  $F$  ones.

For both compounds, data collection: *IPDS-2.87* (Stoe & Cie, 1997); cell refinement: *IPDS-2.87* (Stoe & Cie, 1997); data reduction: *IPDS-2.87* (Stoe & Cie, 1997); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1990); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *XSTEP-2.18* (Stoe & Cie, 1997); software used to prepare material for publication: *SHELXL97* (Sheldrick, 1997).

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