

# 1,1,2,2-Tetraphenyl-1 $\lambda^5$ -diphosphane 1-sulfide

**Bhaskar R. Aluri,<sup>a\*</sup> Stephan Peitz,<sup>a</sup> Anina Wöhrl,<sup>b</sup> Normen Peulecke,<sup>a</sup> Bernd H. Müller,<sup>a</sup> Anke Spannenberg<sup>a</sup> and Uwe Rosenthal<sup>a</sup>**

<sup>a</sup>Leibniz-Institut für Katalyse e. V. an der Universität Rostock, Albert-Einstein-Strasse 29a, 18059 Rostock, Germany, and <sup>b</sup>Linde AG, Linde Engineering Division, Dr.-Carl-von-Linde-Strasse 6–14, 82049 Pullach, Germany

Correspondence e-mail: bhaskar-reddy.aluri@catalysis.de

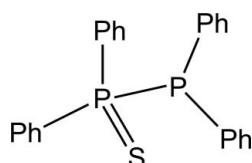
Received 20 January 2009; accepted 23 January 2009

Key indicators: single-crystal X-ray study;  $T = 200$  K; mean  $\sigma(\text{C}-\text{C}) = 0.002$  Å;  $R$  factor = 0.033;  $wR$  factor = 0.096; data-to-parameter ratio = 22.5.

In the title molecule,  $\text{C}_{24}\text{H}_{20}\text{P}_2\text{S}$ , the P–P bond length is 2.2263 (5) Å. The two phenyl rings attached to the three- and five-coordinated P atoms, respectively, form dihedral angles of 56.22 (5) and 71.74 (5)°.

## Related literature

For the literature on related compounds, see: Bhattacharyya *et al.* (1996); Gruber *et al.* (1990); Jones *et al.* (2002).



## Experimental

### Crystal data

$\text{C}_{24}\text{H}_{20}\text{P}_2\text{S}$   
 $M_r = 402.40$   
Monoclinic,  $P2_1/c$   
 $a = 9.32670$  (19) Å  
 $b = 13.6496$  (4) Å  
 $c = 16.0484$  (4) Å  
 $\beta = 91.7298$  (17)°

$V = 2042.12$  (9) Å<sup>3</sup>  
 $Z = 4$   
Mo  $K\alpha$  radiation  
 $\mu = 0.32$  mm<sup>-1</sup>  
 $T = 200$  (2) K  
 $0.45 \times 0.30 \times 0.28$  mm

### Data collection

Stoe IPDS II diffractometer  
Absorption correction: numerical (*X-SHAPE*; Stoe & Cie, 2005)  
 $T_{\min} = 0.892$ ,  $T_{\max} = 0.961$

38972 measured reflections  
5499 independent reflections  
4493 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.027$

### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.033$   
 $wR(F^2) = 0.096$   
 $S = 1.08$   
5499 reflections  
244 parameters

1 restraint  
H-atom parameters constrained  
 $\Delta\rho_{\max} = 0.29$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.28$  e Å<sup>-3</sup>

Data collection: *X-AREA* (Stoe & Cie, 2005); cell refinement: *X-AREA*; data reduction: *X-RED* (Stoe & Cie, 2005); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL*.

This work was supported by the Leibniz-Institut für Katalyse e. V. an der Universität Rostock.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: CV2512).

## References

- Bhattacharyya, P., Slawin, A. M. Z., Smith, M. B., Williams, D. J. & Woollins, J. D. (1996). *J. Chem. Soc. Dalton Trans.* pp. 3647–3651.
- Gruber, M., Jones, P. G. & Schmutzler, R. (1990). *Chem. Ber.* **123**, 1313–1317.
- Jones, P. G., Fischer, A. K., Farkens, M. & Schmutzler, R. (2002). *Acta Cryst. E* **58**, m478–m479.
- Sheldrick, G. M. (2008). *Acta Cryst. A* **64**, 112–122.
- Stoe & Cie (2005). *X-SHAPE*, *X-RED* and *X-AREA*. Stoe & Cie, Darmstadt, Germany.

# supporting information

*Acta Cryst.* (2009). E65, o404 [doi:10.1107/S1600536809002955]

## 1,1,2,2-Tetraphenyl-1 $\lambda^5$ -diphosphane 1-sulfide

**Bhaskar R. Aluri, Stephan Peitz, Anina Wöhl, Normen Peulecke, Bernd H. Müller, Anke Spannenberg and Uwe Rosenthal**

### S1. Comment

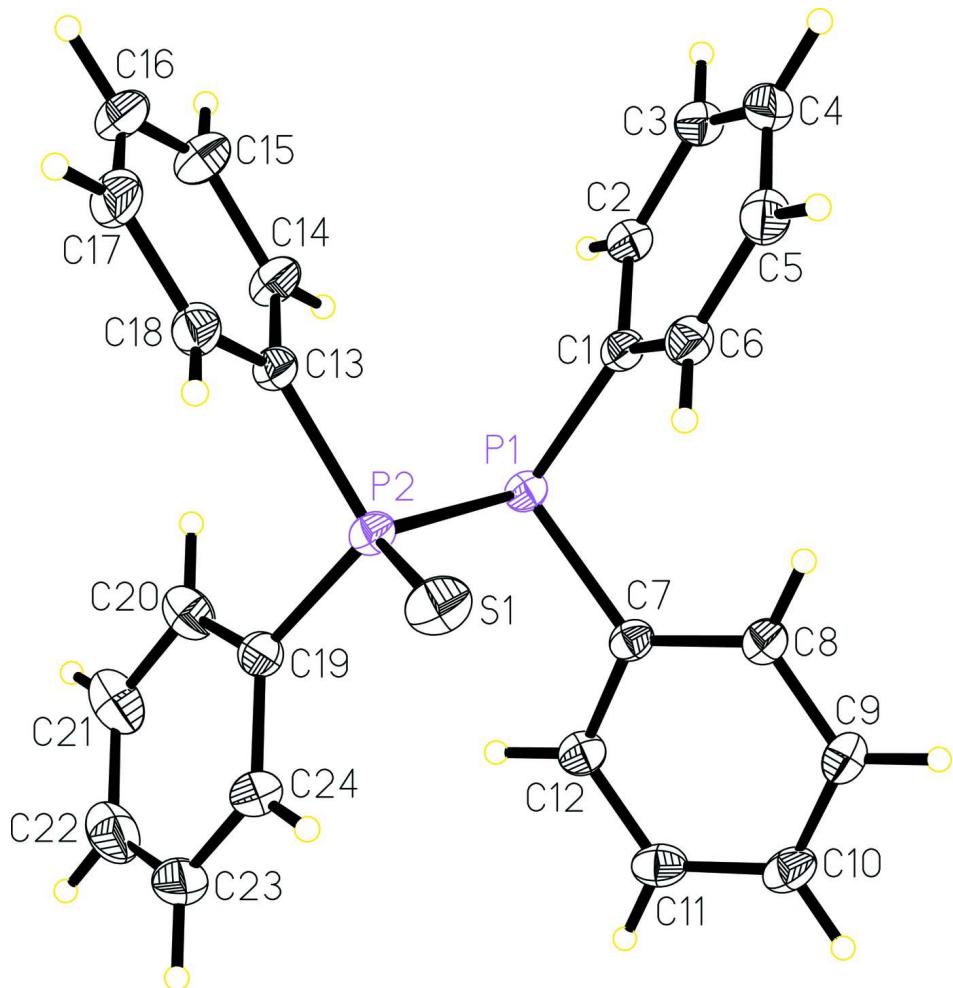
Earlier, Gruber *et al.* (1990) reported that the only phosphorus-containing product formed from the reaction between Ph<sub>2</sub>PCl and thiourea was tetraphenyldiphosphane monosulfide. Bhattacharyya *et al.* (1996) reported this monosulfide formed as the by-product from the same reaction using different reaction conditions. In the present publication, we report the formation of tetraphenyldiphosphane monosulfide, which was observed as a major by-product in the reaction of Ph<sub>2</sub>P(S)CH<sub>2</sub>N(Li)Ph and ClPPh<sub>2</sub>. Its molecular structure (Fig. 1) shows the bond lengths and bond angles are within the normal ranges and are in accordance with the corresponding values in tetramethyldiphosphane monosulfide (Gruber *et al.*, 1990). The molecular structure of pentacarbonyl(tetraphenyldiphosphinomonosulfide-P)chromium(0) was reported earlier by Jones *et al.* (2002).

### S2. Experimental

BuLi (0.77 ml, 1.94 mmol, 2.5 M in *n*-hexane) was added dropwise to a solution of Ph<sub>2</sub>P(S)CH<sub>2</sub>N(H)Ph (660 mg, 2.03 mmol) in THF (5 ml) at -78°C and this reaction mixture was stirred for 4 h while slowly warming up to -40°C. The resultant yellow solution was added to ClPPh<sub>2</sub> (0.38 ml, 2.03 mmol) at 0°C in small portions *via* a cannula over a period of 20 min and followed stirring at room temperature overnight. The major part of THF was removed from the reaction mixture and over-layered with *n*-hexane to get single crystals of the title compound, which were suitable for X-ray analysis. <sup>31</sup>P NMR (THF-*d*<sub>8</sub>): -14.11 (d, <sup>1</sup>J = 247.3 Hz), 44.11 (d, <sup>1</sup>J = 247.3 Hz).

### S3. Refinement

All H atoms were placed in idealized positions (C—H = 0.95 Å) and refined using a riding model with *U*<sub>iso</sub>(H) fixed at 1.2 *U*<sub>eq</sub>(C).

**Figure 1**

The molecular structure of the title compound showing the atom-labelling scheme and 30% probability displacement ellipsoids.

### 1,1,2,2-Tetraphenyl-1 $\lambda^5$ -diphosphane 1-sulfide

#### Crystal data

$C_{24}H_{20}P_2S$   
 $M_r = 402.40$   
Monoclinic,  $P2_1/c$   
Hall symbol: -P 2ybc  
 $a = 9.32670 (19)$  Å  
 $b = 13.6496 (4)$  Å  
 $c = 16.0484 (4)$  Å  
 $\beta = 91.7298 (17)^\circ$   
 $V = 2042.12 (9)$  Å<sup>3</sup>  
 $Z = 4$

$F(000) = 840$   
 $D_x = 1.309 \text{ Mg m}^{-3}$   
Mo  $K\alpha$  radiation,  $\lambda = 0.71073$  Å  
Cell parameters from 32981 reflections  
 $\theta = 2.0-29.6^\circ$   
 $\mu = 0.32 \text{ mm}^{-1}$   
 $T = 200$  K  
Prism, colourless  
 $0.45 \times 0.30 \times 0.28$  mm

*Data collection*

Stoe IPDS II  
 diffractometer  
 Radiation source: fine-focus sealed tube  
 Graphite monochromator  
 rotation method scans  
 Absorption correction: numerical  
 (*X-SHAPE*; Stoe & Cie, 2005)  
 $T_{\min} = 0.892$ ,  $T_{\max} = 0.961$   
 38972 measured reflections  
 5499 independent reflections  
 4493 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.027$   
 $\theta_{\max} = 29.1^\circ$ ,  $\theta_{\min} = 2.0^\circ$   
 $h = -12 \rightarrow 12$   
 $k = -18 \rightarrow 18$   
 $l = -21 \rightarrow 21$

*Refinement*

Refinement on  $F^2$   
 Least-squares matrix: full  
 $R[F^2 > 2\sigma(F^2)] = 0.033$   
 $wR(F^2) = 0.096$   
 $S = 1.08$   
 5499 reflections  
 244 parameters  
 1 restraint  
 Primary atom site location: structure-invariant  
 direct methods  
 Secondary atom site location: difference Fourier  
 map  
 Hydrogen site location: inferred from  
 neighbouring sites  
 H-atom parameters constrained  
 $w = 1/[\sigma^2(F_o^2) + (0.0564P)^2 + 0.1948P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} = 0.001$   
 $\Delta\rho_{\max} = 0.29 \text{ e } \text{\AA}^{-3}$   
 $\Delta\rho_{\min} = -0.28 \text{ e } \text{\AA}^{-3}$

*Special details*

**Geometry.** All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

**Refinement.** Refinement of  $F^2$  against ALL reflections. The weighted  $R$ -factor  $wR$  and goodness of fit  $S$  are based on  $F^2$ , conventional  $R$ -factors  $R$  are based on  $F$ , with  $F$  set to zero for negative  $F^2$ . The threshold expression of  $F^2 > \sigma(F^2)$  is used only for calculating  $R$ -factors(gt) etc. and is not relevant to the choice of reflections for refinement.  $R$ -factors based on  $F^2$  are statistically about twice as large as those based on  $F$ , and  $R$ -factors based on ALL data will be even larger.

*Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters ( $\text{\AA}^2$ )*

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.58305 (14)	0.77663 (9)	-0.01356 (8)	0.0323 (3)
C2	0.58628 (15)	0.78402 (11)	-0.10028 (8)	0.0380 (3)
H2A	0.5640	0.7283	-0.1337	0.046*
C3	0.62169 (16)	0.87176 (12)	-0.13813 (10)	0.0450 (3)
H3A	0.6232	0.8761	-0.1972	0.054*
C4	0.65480 (16)	0.95274 (11)	-0.08978 (11)	0.0469 (3)
H4A	0.6797	1.0128	-0.1156	0.056*
C5	0.65188 (16)	0.94677 (11)	-0.00372 (10)	0.0439 (3)
H5A	0.6751	1.0026	0.0293	0.053*
C6	0.61519 (15)	0.85951 (10)	0.03445 (9)	0.0376 (3)
H6A	0.6119	0.8561	0.0935	0.045*
C7	0.66830 (14)	0.62119 (9)	0.10362 (7)	0.0309 (2)
C8	0.78956 (15)	0.67684 (11)	0.12330 (9)	0.0399 (3)
H8A	0.7999	0.7401	0.0996	0.048*
C9	0.89570 (16)	0.64037 (12)	0.17754 (10)	0.0455 (3)
H9A	0.9782	0.6789	0.1906	0.055*

C10	0.88200 (16)	0.54883 (12)	0.21250 (9)	0.0436 (3)
H10A	0.9544	0.5244	0.2499	0.052*
C11	0.76235 (17)	0.49268 (11)	0.19287 (9)	0.0420 (3)
H11A	0.7528	0.4294	0.2167	0.050*
C12	0.65601 (15)	0.52821 (10)	0.13849 (8)	0.0360 (3)
H12A	0.5745	0.4889	0.1250	0.043*
C13	0.22346 (14)	0.75419 (10)	0.03664 (9)	0.0352 (3)
C14	0.21864 (17)	0.73376 (12)	-0.04835 (10)	0.0461 (3)
H14A	0.2841	0.6882	-0.0708	0.055*
C15	0.11782 (19)	0.78017 (15)	-0.10032 (11)	0.0589 (5)
H15A	0.1161	0.7675	-0.1586	0.071*
C16	0.02074 (18)	0.84419 (13)	-0.06782 (10)	0.0609 (5)
H16A	-0.0496	0.8743	-0.1034	0.073*
C17	0.02482 (17)	0.86493 (12)	0.01602 (12)	0.0557 (4)
H17A	-0.0422	0.9097	0.0380	0.067*
C18	0.12649 (16)	0.82073 (11)	0.06868 (11)	0.0430 (3)
H18A	0.1298	0.8359	0.1265	0.052*
C19	0.28040 (15)	0.57847 (10)	0.13544 (9)	0.0379 (3)
C20	0.21178 (18)	0.51924 (11)	0.07596 (10)	0.0459 (3)
H20A	0.1989	0.5421	0.0203	0.055*
C21	0.1620 (2)	0.42703 (13)	0.09741 (13)	0.0589 (5)
H21A	0.1140	0.3873	0.0568	0.071*
C22	0.1824 (2)	0.39341 (13)	0.17769 (15)	0.0642 (5)
H22A	0.1500	0.3299	0.1922	0.077*
C23	0.2495 (2)	0.45130 (16)	0.23683 (14)	0.0675 (6)
H23A	0.2627	0.4275	0.2922	0.081*
C24	0.29853 (18)	0.54423 (14)	0.21697 (11)	0.0532 (4)
H24A	0.3439	0.5841	0.2585	0.064*
P1	0.53134 (3)	0.65612 (2)	0.024981 (19)	0.03019 (9)
P2	0.35427 (4)	0.69731 (2)	0.10811 (2)	0.03240 (9)
S1	0.39983 (5)	0.77769 (3)	0.20569 (2)	0.04904 (11)

Atomic displacement parameters ( $\text{\AA}^2$ )

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
C1	0.0296 (6)	0.0325 (6)	0.0344 (6)	0.0026 (5)	-0.0032 (5)	0.0022 (5)
C2	0.0367 (7)	0.0421 (7)	0.0350 (6)	0.0007 (6)	-0.0036 (5)	0.0036 (5)
C3	0.0396 (8)	0.0531 (8)	0.0423 (7)	0.0007 (6)	0.0002 (6)	0.0132 (6)
C4	0.0374 (7)	0.0414 (8)	0.0619 (9)	0.0012 (6)	0.0019 (6)	0.0156 (7)
C5	0.0371 (7)	0.0336 (7)	0.0608 (9)	0.0017 (6)	-0.0018 (6)	-0.0012 (6)
C6	0.0363 (7)	0.0359 (7)	0.0404 (7)	0.0015 (5)	-0.0020 (5)	-0.0015 (5)
C7	0.0305 (6)	0.0328 (6)	0.0294 (5)	0.0062 (5)	-0.0001 (4)	-0.0012 (4)
C8	0.0347 (7)	0.0404 (7)	0.0441 (7)	-0.0002 (5)	-0.0050 (6)	0.0058 (6)
C9	0.0317 (7)	0.0548 (9)	0.0494 (8)	0.0009 (6)	-0.0065 (6)	0.0038 (7)
C10	0.0391 (7)	0.0533 (8)	0.0383 (7)	0.0160 (6)	-0.0034 (6)	0.0020 (6)
C11	0.0511 (8)	0.0361 (7)	0.0390 (7)	0.0121 (6)	0.0018 (6)	0.0043 (5)
C12	0.0392 (7)	0.0312 (6)	0.0373 (6)	0.0045 (5)	-0.0005 (5)	-0.0015 (5)
C13	0.0304 (6)	0.0321 (6)	0.0430 (7)	0.0011 (5)	-0.0030 (5)	0.0059 (5)

C14	0.0389 (7)	0.0547 (9)	0.0438 (7)	0.0065 (7)	-0.0107 (6)	0.0024 (6)
C15	0.0479 (9)	0.0735 (12)	0.0542 (9)	0.0031 (8)	-0.0182 (8)	0.0125 (8)
C16	0.0405 (8)	0.0593 (10)	0.0816 (13)	0.0032 (8)	-0.0181 (8)	0.0254 (9)
C17	0.0359 (8)	0.0426 (8)	0.0887 (13)	0.0083 (6)	0.0001 (8)	0.0139 (8)
C18	0.0343 (7)	0.0352 (7)	0.0595 (9)	0.0035 (5)	0.0032 (6)	0.0068 (6)
C19	0.0321 (6)	0.0387 (7)	0.0432 (7)	0.0083 (5)	0.0062 (5)	0.0076 (5)
C20	0.0526 (9)	0.0380 (7)	0.0481 (8)	0.0001 (6)	0.0162 (7)	-0.0033 (6)
C21	0.0642 (11)	0.0407 (8)	0.0733 (11)	-0.0032 (8)	0.0278 (9)	-0.0076 (8)
C22	0.0558 (11)	0.0419 (9)	0.0963 (15)	0.0086 (8)	0.0248 (10)	0.0192 (9)
C23	0.0510 (10)	0.0739 (13)	0.0776 (12)	0.0079 (9)	0.0020 (9)	0.0428 (11)
C24	0.0423 (8)	0.0632 (10)	0.0538 (9)	0.0029 (7)	-0.0049 (7)	0.0224 (8)
P1	0.03128 (16)	0.03040 (16)	0.02864 (15)	0.00166 (12)	-0.00294 (12)	-0.00140 (11)
P2	0.03193 (17)	0.03396 (17)	0.03117 (16)	0.00544 (13)	-0.00136 (12)	0.00023 (12)
S1	0.0586 (2)	0.0524 (2)	0.03591 (18)	0.00869 (18)	-0.00217 (16)	-0.01034 (15)

*Geometric parameters ( $\text{\AA}$ ,  $^{\circ}$ )*

C1—C6	1.3962 (18)	C13—C14	1.392 (2)
C1—C2	1.3966 (18)	C13—P2	1.8228 (13)
C1—P1	1.8273 (13)	C14—C15	1.391 (2)
C2—C3	1.387 (2)	C14—H14A	0.9500
C2—H2A	0.9500	C15—C16	1.3727 (19)
C3—C4	1.380 (2)	C15—H15A	0.9500
C3—H3A	0.9500	C16—C17	1.3743 (19)
C4—C5	1.385 (2)	C16—H16A	0.9500
C4—H4A	0.9500	C17—C18	1.389 (2)
C5—C6	1.387 (2)	C17—H17A	0.9500
C5—H5A	0.9500	C18—H18A	0.9500
C6—H6A	0.9500	C19—C20	1.392 (2)
C7—C8	1.3907 (19)	C19—C24	1.395 (2)
C7—C12	1.3931 (18)	C19—P2	1.8210 (15)
C7—P1	1.8316 (12)	C20—C21	1.388 (2)
C8—C9	1.3905 (19)	C20—H20A	0.9500
C8—H8A	0.9500	C21—C22	1.375 (3)
C9—C10	1.377 (2)	C21—H21A	0.9500
C9—H9A	0.9500	C22—C23	1.372 (3)
C10—C11	1.382 (2)	C22—H22A	0.9500
C10—H10A	0.9500	C23—C24	1.389 (3)
C11—C12	1.3887 (19)	C23—H23A	0.9500
C11—H11A	0.9500	C24—H24A	0.9500
C12—H12A	0.9500	P1—P2	2.2263 (5)
C13—C18	1.391 (2)	P2—S1	1.9486 (5)
C6—C1—C2	118.73 (13)	C13—C14—H14A	120.1
C6—C1—P1	126.67 (10)	C16—C15—C14	120.26 (16)
C2—C1—P1	114.59 (10)	C16—C15—H15A	119.9
C3—C2—C1	120.75 (14)	C14—C15—H15A	119.9
C3—C2—H2A	119.6	C15—C16—C17	120.27 (15)

C1—C2—H2A	119.6	C15—C16—H16A	119.9
C4—C3—C2	119.83 (14)	C17—C16—H16A	119.9
C4—C3—H3A	120.1	C16—C17—C18	120.30 (15)
C2—C3—H3A	120.1	C16—C17—H17A	119.9
C3—C4—C5	120.19 (14)	C18—C17—H17A	119.9
C3—C4—H4A	119.9	C17—C18—C13	119.88 (16)
C5—C4—H4A	119.9	C17—C18—H18A	120.1
C4—C5—C6	120.24 (14)	C13—C18—H18A	120.1
C4—C5—H5A	119.9	C20—C19—C24	119.26 (14)
C6—C5—H5A	119.9	C20—C19—P2	121.44 (11)
C5—C6—C1	120.26 (13)	C24—C19—P2	119.23 (12)
C5—C6—H6A	119.9	C21—C20—C19	120.44 (16)
C1—C6—H6A	119.9	C21—C20—H20A	119.8
C8—C7—C12	118.85 (12)	C19—C20—H20A	119.8
C8—C7—P1	124.04 (10)	C22—C21—C20	119.80 (19)
C12—C7—P1	116.73 (10)	C22—C21—H21A	120.1
C9—C8—C7	120.35 (14)	C20—C21—H21A	120.1
C9—C8—H8A	119.8	C23—C22—C21	120.22 (17)
C7—C8—H8A	119.8	C23—C22—H22A	119.9
C10—C9—C8	120.45 (14)	C21—C22—H22A	119.9
C10—C9—H9A	119.8	C22—C23—C24	120.92 (17)
C8—C9—H9A	119.8	C22—C23—H23A	119.5
C9—C10—C11	119.63 (13)	C24—C23—H23A	119.5
C9—C10—H10A	120.2	C23—C24—C19	119.34 (18)
C11—C10—H10A	120.2	C23—C24—H24A	120.3
C10—C11—C12	120.40 (13)	C19—C24—H24A	120.3
C10—C11—H11A	119.8	C1—P1—C7	106.37 (6)
C12—C11—H11A	119.8	C1—P1—P2	100.58 (5)
C11—C12—C7	120.31 (13)	C7—P1—P2	99.63 (4)
C11—C12—H12A	119.8	C19—P2—C13	106.26 (6)
C7—C12—H12A	119.8	C19—P2—S1	112.51 (5)
C18—C13—C14	119.41 (13)	C13—P2—S1	113.05 (5)
C18—C13—P2	118.41 (11)	C19—P2—P1	102.26 (5)
C14—C13—P2	122.18 (11)	C13—P2—P1	103.00 (5)
C15—C14—C13	119.85 (16)	S1—P2—P1	118.43 (2)
C15—C14—H14A	120.1		
C6—C1—C2—C3	0.4 (2)	C22—C23—C24—C19	-0.7 (3)
P1—C1—C2—C3	179.04 (11)	C20—C19—C24—C23	0.9 (2)
C1—C2—C3—C4	0.3 (2)	P2—C19—C24—C23	-176.34 (14)
C2—C3—C4—C5	-0.4 (2)	C6—C1—P1—C7	-52.24 (13)
C3—C4—C5—C6	-0.2 (2)	C2—C1—P1—C7	129.30 (10)
C4—C5—C6—C1	1.0 (2)	C6—C1—P1—P2	51.19 (12)
C2—C1—C6—C5	-1.1 (2)	C2—C1—P1—P2	-127.26 (10)
P1—C1—C6—C5	-179.48 (11)	C8—C7—P1—C1	-3.34 (14)
C12—C7—C8—C9	-0.6 (2)	C12—C7—P1—C1	-176.18 (10)
P1—C7—C8—C9	-173.34 (12)	C8—C7—P1—P2	-107.46 (12)
C7—C8—C9—C10	-0.1 (2)	C12—C7—P1—P2	79.70 (10)

C8—C9—C10—C11	0.5 (2)	C20—C19—P2—C13	41.29 (14)
C9—C10—C11—C12	-0.3 (2)	C24—C19—P2—C13	-141.56 (12)
C10—C11—C12—C7	-0.5 (2)	C20—C19—P2—S1	165.51 (11)
C8—C7—C12—C11	0.9 (2)	C24—C19—P2—S1	-17.34 (14)
P1—C7—C12—C11	174.15 (11)	C20—C19—P2—P1	-66.36 (12)
C18—C13—C14—C15	0.3 (2)	C24—C19—P2—P1	110.79 (12)
P2—C13—C14—C15	-179.17 (13)	C18—C13—P2—C19	97.09 (12)
C13—C14—C15—C16	-1.6 (3)	C14—C13—P2—C19	-83.43 (14)
C14—C15—C16—C17	1.7 (3)	C18—C13—P2—S1	-26.79 (13)
C15—C16—C17—C18	-0.5 (3)	C14—C13—P2—S1	152.68 (11)
C16—C17—C18—C13	-0.8 (2)	C18—C13—P2—P1	-155.79 (10)
C14—C13—C18—C17	0.9 (2)	C14—C13—P2—P1	23.68 (13)
P2—C13—C18—C17	-179.60 (12)	C1—P1—P2—C19	171.48 (6)
C24—C19—C20—C21	-0.1 (2)	C7—P1—P2—C19	-79.70 (6)
P2—C19—C20—C21	177.07 (13)	C1—P1—P2—C13	61.35 (6)
C19—C20—C21—C22	-0.9 (3)	C7—P1—P2—C13	170.16 (6)
C20—C21—C22—C23	1.1 (3)	C1—P1—P2—S1	-64.24 (5)
C21—C22—C23—C24	-0.3 (3)	C7—P1—P2—S1	44.58 (5)