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Bis(μ -diisopropylphosphanido- $\kappa^2P:P$)bis-[hydrido(triisopropylphosphane- κP)-platinum(II)]

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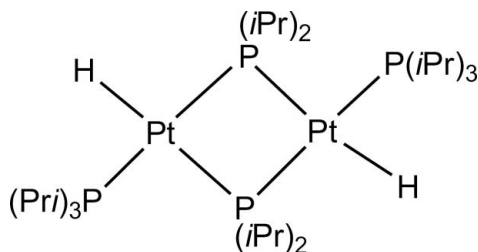
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Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(C-C) = 0.004$ Å; R factor = 0.018; wR factor = 0.038; data-to-parameter ratio = 22.3.

In the centrosymmetric molecular structure of the title compound $[Pt_2(C_6H_{14}P)_2H_2(C_9H_{21}P)_2]$, each Pt^{II} atom is bound on one side to a phosphane ligand ($PiPr_3$) and a hydrido ligand. On the other side, it is bound to two phosphanido ligands (μ - $PiPr_2$), which engage a bridging position between the two Pt^{II} atoms, forming a distorted square-planar structure motif. The $Pt \cdots Pt$ distance is 3.6755 (2) Å. A comparable molecular structure was observed for bis(μ -di-*tert*-butylphosphanido)-bis[hydrido(triethylphosphane)platinum(II)] [Itazaki *et al.* (2004). *Organometallics*, **23**, 1610–1621].

Related literature

For the syntheses of similar phosphido-bridged complexes of platinum(II) with phosphine ligands, see: Itazaki *et al.* (2004) or with other ligands such as carbonyl, see: Albinati *et al.* (2008). For $Pt-H$ bond lengths in related structures, see: Chiang *et al.* (1984); Knobler *et al.* (1983).



Experimental

Crystal data

$[Pt_2(C_6H_{14}P)_2H_2(C_9H_{21}P)_2]$
 $M_r = 946.94$
 Monoclinic, $P2_1/n$
 $a = 8.8301$ (3) Å
 $b = 14.8153$ (5) Å
 $c = 14.1688$ (5) Å
 $\beta = 90.097$ (2)°

$V = 1853.57$ (11) Å³
 $Z = 2$
 Mo $K\alpha$ radiation
 $\mu = 7.73$ mm⁻¹
 $T = 100$ K
 $0.53 \times 0.13 \times 0.11$ mm

Data collection

Bruker X8 APEXII diffractometer
 Absorption correction: multi-scan
 (SADABS; Bruker, 2008)
 $T_{min} = 0.360$, $T_{max} = 0.745$

38282 measured reflections
 3943 independent reflections
 3478 reflections with $I > 2\sigma(I)$
 $R_{int} = 0.051$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.018$
 $wR(F^2) = 0.038$
 $S = 1.03$
 3943 reflections
 177 parameters

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{max} = 0.72$ e Å⁻³
 $\Delta\rho_{min} = -0.66$ e Å⁻³

Data collection: APEX2 (Bruker, 2010); cell refinement: SAINT-Plus (Bruker, 2010); data reduction: SAINT-Plus; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: XP in SHELXTL (Sheldrick, 2008); software used to prepare material for publication: SHELXL97.

Financial support by the DFG is gratefully acknowledged.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HP2036).

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supporting information

Acta Cryst. (2012). E68, m808 [doi:10.1107/S1600536812022829]

Bis(μ -diisopropylphosphanido- $\kappa^2P:P$)bis[hydrido(triisopropylphosphane- κP)platinum(II)]

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S1. Comment

Bis[μ -di(*isopropyl*)phosphino]-di(hydrido)-bis[tri(*isopropyl*)phosphine]-di(platinum), bridged by the μ -*PtPr*₂ ligands, displays a slightly distorted square-planar geometry. The two platinum centers show a Pt(1)–Pt(1ⁱ) distance of 3.6755 (2) Å. The Pt–Pt distance is comparable to that in bis[μ -di(*tert*-butyl)phosphino]-di(hydrido)-bis[tri(ethyl)phosphine]-di(platinum) [Pt₂H₂(μ -P^tBu₂)₂(PEt₃)₂] (3.646 Å).

The bond angles P(13)–Pt(1)–P(13ⁱ) [77.47 (3)°] and Pt(1)–P(13)–Pt(1ⁱ) [102.53 (3)°] are slightly out of range of the structural parameters of the complexes without Pt–Pt bonding from Itazaki *et al.* (2004) [P–Pt–P 74.6–77.2° and Pt–P–Pt 102.8–105.4°]. This could be due to the less sterical hindrance of the *iso*-propyl groups by contrast with the *tert*-butyl groups in the reference substance [Pt₂H₂(μ -P^tBu₂)₂(PEt₃)₂].

Chiang *et al.* (1984) reported the bond length of a terminal Pt–H bond determined by neutron diffraction method. They found for the Pt–H bond on a five coordinate platinum centre a bond length of 1.610 (2) Å in the compound [Pt₂H₃(Ph₂PCH₂CH₂PPh₂)₂]⁺[BPh₄]⁻. In the title compound [Pt₂H₂(μ -*PtPr*₂)₂(*PtPr*₃)₂] [1.57 (3) Å] the bonding distance of Pt–H is 2.5% shorter than in the neutron experiment of Chiang *et al.*, due to the smaller coordination number of four in the former species.

The group of Knobler *et al.* (1983) also determined the Pt–H bond length by X-Ray diffraction in [Pt₂H₃(Ph₂PCH₂CH₂PPh₂)₂]⁺[BPh₄]⁻ to be 1.527 Å, however without further refinement.

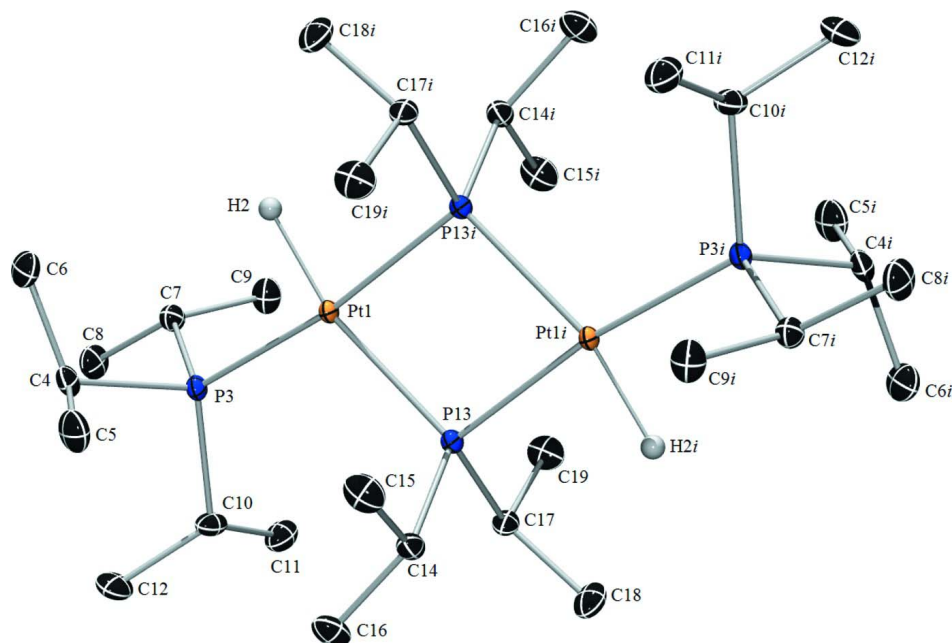
The bonding distances Pt–P in *trans*-position to the hydrido ligand are with 2.3773 (7) Å longer than the bonding distances in *trans*-position to the phosphine ligand 2.3343 (7) Å.

S2. Experimental

Bis(tri-*iso*-propylphosphine)platinum (50.0 mg, 0.09 mmol) dissolved in 1 ml benzene was added to a solution of dichloro(2,3,5,6-tetramethylphenyl)borane (29.5 mg, 0.09 mmol) in 1 ml benzene. The solvent was removed under reduced pressure and the obtained dark brown residue was dissolved in hexanes. The title compound was obtained as a off-white solid. Colourless crystals suitable for X-ray analysis were grown from a hexanes solution at 238 K.

S3. Refinement

The H atoms were placed at idealized positions and treated as riding atoms: C–H = 0.98 Å (CH₃), 1.00 Å (aliphatic H-atoms). $U_{\text{iso}}(\text{H})$ values were fixed at 1.5 times (for primary H atoms) and 1.2 times (tertiary H atoms) U_{eq} of the attached C atoms.

**Figure 1**

The molecular structure of the title compound showing the atom numbering scheme and displacement ellipsoids for the non-H atoms at the 50% probability level. Hydrogen atoms are omitted for clarity.

Bis(μ -diisopropylphosphanido- $\kappa^2P:P$)bis[hydrido(triisopropylphosphane- κP)platinum(II)]

Crystal data

[Pt₂(C₆H₁₄P)₂H₂(C₉H₂₁P)₂]

$M_r = 946.94$

Monoclinic, $P2_1/n$

Hall symbol: -P 2yn

$a = 8.8301$ (3) Å

$b = 14.8153$ (5) Å

$c = 14.1688$ (5) Å

$\beta = 90.097$ (2)°

$V = 1853.57$ (11) Å³

$Z = 2$

$F(000) = 936$

$D_x = 1.697$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 8162 reflections

$\theta = 2.7$ – 26.7 °

$\mu = 7.73$ mm⁻¹

$T = 100$ K

Needle, colourless

$0.53 \times 0.13 \times 0.11$ mm

Data collection

Bruker X8 APEXII
diffractometer

Radiation source: rotating anode

Multi-layer mirror monochromator

Detector resolution: 8.333 pixels mm⁻¹

φ and ω scans

Absorption correction: multi-scan
(*SADABS*; Bruker, 2008)

$T_{\min} = 0.360$, $T_{\max} = 0.745$

38282 measured reflections

3943 independent reflections

3478 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.051$

$\theta_{\text{max}} = 26.8$ °, $\theta_{\text{min}} = 2.0$ °

$h = -11 \rightarrow 11$

$k = -18 \rightarrow 18$

$l = -17 \rightarrow 17$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.018$
 $wR(F^2) = 0.038$
 $S = 1.03$
 3943 reflections
 177 parameters
 0 restraints
 Primary atom site location: structure-invariant
 direct methods

Secondary atom site location: difference Fourier
 map
 Hydrogen site location: inferred from
 neighbouring sites
 H atoms treated by a mixture of independent
 and constrained refinement
 $w = 1/[\sigma^2(F_o^2) + (0.0141P)^2 + 0.9947P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.009$
 $\Delta\rho_{\max} = 0.72 \text{ e } \text{Å}^{-3}$
 $\Delta\rho_{\min} = -0.66 \text{ e } \text{Å}^{-3}$

Special details

Experimental. The crystal was immersed in a film of perfluoropolyether oil, mounted on a polyimide microloop (MicroMounts of MiTeGen) and transferred to stream of cold nitrogen (Oxford Cryostream 700).

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 (Å^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
Pt1	0.068246 (11)	1.032840 (7)	0.882448 (6)	0.00916 (4)
P3	0.15611 (8)	0.99092 (5)	0.73662 (5)	0.01104 (15)
C4	0.0188 (3)	1.01911 (19)	0.64139 (19)	0.0155 (6)
H4	0.0570	0.9925	0.5811	0.019*
C5	-0.1344 (3)	0.9764 (2)	0.6632 (2)	0.0230 (7)
H5A	-0.1717	0.9993	0.7238	0.034*
H5B	-0.1233	0.9107	0.6667	0.034*
H5C	-0.2067	0.9918	0.6132	0.034*
C6	0.0005 (3)	1.1205 (2)	0.6271 (2)	0.0218 (7)
H6A	-0.0818	1.1318	0.5822	0.033*
H6B	0.0951	1.1457	0.6023	0.033*
H6C	-0.0234	1.1491	0.6876	0.033*
C7	0.3257 (3)	1.05748 (19)	0.70284 (18)	0.0141 (6)
H7	0.2888	1.1209	0.6953	0.017*
C8	0.3981 (3)	1.0330 (2)	0.60752 (19)	0.0195 (7)
H8A	0.4768	1.0773	0.5921	0.029*
H8B	0.3202	1.0334	0.5581	0.029*
H8C	0.4434	0.9727	0.6116	0.029*
C9	0.4449 (3)	1.0616 (2)	0.7817 (2)	0.0199 (7)
H9A	0.4974	1.0034	0.7859	0.030*
H9B	0.3951	1.0747	0.8419	0.030*
H9C	0.5184	1.1093	0.7675	0.030*

C10	0.1983 (3)	0.86919 (18)	0.71777 (19)	0.0156 (6)
H10	0.1209	0.8361	0.7558	0.019*
C11	0.3510 (3)	0.8423 (2)	0.7605 (2)	0.0212 (7)
H11A	0.3587	0.7763	0.7627	0.032*
H11B	0.3591	0.8667	0.8246	0.032*
H11C	0.4331	0.8666	0.7215	0.032*
C12	0.1825 (4)	0.8318 (2)	0.6177 (2)	0.0235 (7)
H12A	0.2552	0.8620	0.5761	0.035*
H12B	0.0794	0.8427	0.5946	0.035*
H12C	0.2028	0.7668	0.6181	0.035*
P13	0.00800 (7)	0.90365 (5)	0.97455 (5)	0.01004 (14)
C14	-0.1496 (3)	0.82851 (18)	0.93640 (18)	0.0141 (6)
H14	-0.1686	0.7858	0.9897	0.017*
C15	-0.2950 (3)	0.8827 (2)	0.9230 (2)	0.0220 (7)
H15A	-0.2809	0.9267	0.8721	0.033*
H15B	-0.3194	0.9145	0.9817	0.033*
H15C	-0.3782	0.8418	0.9066	0.033*
C16	-0.1193 (3)	0.7706 (2)	0.8494 (2)	0.0197 (7)
H16A	-0.2015	0.7268	0.8415	0.030*
H16B	-0.0231	0.7385	0.8573	0.030*
H16C	-0.1139	0.8094	0.7934	0.030*
C17	0.1654 (3)	0.82156 (18)	0.99243 (18)	0.0129 (6)
H17	0.1924	0.7960	0.9293	0.015*
C18	0.1266 (3)	0.7430 (2)	1.0573 (2)	0.0222 (7)
H18A	0.2147	0.7032	1.0633	0.033*
H18B	0.0414	0.7090	1.0307	0.033*
H18C	0.0987	0.7662	1.1197	0.033*
C19	0.3042 (3)	0.8713 (2)	1.0303 (2)	0.0216 (7)
H19A	0.2810	0.8971	1.0923	0.032*
H19B	0.3318	0.9199	0.9866	0.032*
H19C	0.3890	0.8290	1.0362	0.032*
H2	0.099 (3)	1.131 (2)	0.845 (2)	0.040 (9)*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Pt1	0.01140 (6)	0.00952 (6)	0.00656 (6)	0.00047 (4)	0.00126 (4)	-0.00011 (4)
P3	0.0133 (4)	0.0122 (4)	0.0076 (3)	0.0013 (3)	0.0004 (3)	-0.0003 (3)
C4	0.0205 (16)	0.0180 (16)	0.0081 (13)	0.0010 (12)	-0.0003 (11)	-0.0021 (11)
C5	0.0195 (17)	0.0312 (19)	0.0182 (15)	-0.0033 (14)	-0.0068 (13)	-0.0003 (13)
C6	0.0242 (17)	0.0225 (18)	0.0187 (15)	0.0058 (14)	-0.0075 (12)	0.0020 (13)
C7	0.0161 (15)	0.0142 (15)	0.0118 (14)	-0.0016 (12)	0.0010 (11)	0.0002 (11)
C8	0.0206 (16)	0.0240 (18)	0.0140 (15)	-0.0027 (13)	0.0056 (12)	0.0024 (12)
C9	0.0155 (16)	0.0264 (17)	0.0179 (15)	-0.0036 (13)	-0.0011 (12)	0.0028 (13)
C10	0.0203 (16)	0.0105 (14)	0.0160 (14)	0.0000 (12)	0.0059 (12)	-0.0014 (12)
C11	0.0253 (17)	0.0175 (17)	0.0208 (16)	0.0049 (13)	0.0050 (13)	0.0028 (13)
C12	0.0338 (19)	0.0163 (16)	0.0204 (16)	0.0022 (14)	0.0037 (13)	-0.0076 (13)
P13	0.0118 (4)	0.0099 (3)	0.0085 (3)	0.0007 (3)	0.0010 (3)	-0.0005 (3)

C14	0.0178 (15)	0.0125 (15)	0.0121 (14)	-0.0047 (12)	0.0017 (11)	-0.0003 (11)
C15	0.0154 (16)	0.0255 (18)	0.0250 (16)	-0.0012 (13)	-0.0018 (12)	-0.0051 (14)
C16	0.0186 (16)	0.0195 (17)	0.0209 (15)	-0.0076 (13)	0.0024 (12)	-0.0062 (13)
C17	0.0150 (15)	0.0111 (15)	0.0125 (13)	0.0039 (11)	0.0023 (11)	-0.0006 (11)
C18	0.0283 (18)	0.0182 (17)	0.0201 (16)	0.0104 (14)	0.0030 (13)	0.0050 (13)
C19	0.0155 (16)	0.0229 (17)	0.0265 (17)	0.0036 (13)	-0.0028 (12)	-0.0021 (14)

Geometric parameters (Å, °)

Pt1—P3	2.2940 (7)	C11—H11A	0.9800
Pt1—P13 ⁱ	2.3343 (7)	C11—H11B	0.9800
Pt1—P13	2.3773 (7)	C11—H11C	0.9800
Pt1—H2	1.57 (3)	C12—H12A	0.9800
P3—C7	1.857 (3)	C12—H12B	0.9800
P3—C4	1.860 (3)	C12—H12C	0.9800
P3—C10	1.861 (3)	P13—C14	1.862 (3)
C4—C6	1.524 (4)	P13—C17	1.864 (3)
C4—C5	1.526 (4)	P13—Pt1 ⁱ	2.3343 (7)
C4—H4	1.0000	C14—C16	1.526 (4)
C5—H5A	0.9800	C14—C15	1.526 (4)
C5—H5B	0.9800	C14—H14	1.0000
C5—H5C	0.9800	C15—H15A	0.9800
C6—H6A	0.9800	C15—H15B	0.9800
C6—H6B	0.9800	C15—H15C	0.9800
C6—H6C	0.9800	C16—H16A	0.9800
C7—C9	1.534 (4)	C16—H16B	0.9800
C7—C8	1.538 (4)	C16—H16C	0.9800
C7—H7	1.0000	C17—C18	1.523 (4)
C8—H8A	0.9800	C17—C19	1.527 (4)
C8—H8B	0.9800	C17—H17	1.0000
C8—H8C	0.9800	C18—H18A	0.9800
C9—H9A	0.9800	C18—H18B	0.9800
C9—H9B	0.9800	C18—H18C	0.9800
C9—H9C	0.9800	C19—H19A	0.9800
C10—C12	1.529 (4)	C19—H19B	0.9800
C10—C11	1.530 (4)	C19—H19C	0.9800
C10—H10	1.0000		
P3—Pt1—P13 ⁱ	171.67 (3)	C10—C11—H11A	109.5
P3—Pt1—P13	110.66 (2)	C10—C11—H11B	109.5
P13 ⁱ —Pt1—P13	77.47 (3)	H11A—C11—H11B	109.5
P3—Pt1—H2	83.5 (12)	C10—C11—H11C	109.5
P13 ⁱ —Pt1—H2	88.4 (12)	H11A—C11—H11C	109.5
P13—Pt1—H2	165.8 (12)	H11B—C11—H11C	109.5
C7—P3—C4	102.65 (13)	C10—C12—H12A	109.5
C7—P3—C10	108.41 (13)	C10—C12—H12B	109.5
C4—P3—C10	104.12 (13)	H12A—C12—H12B	109.5
C7—P3—Pt1	111.26 (9)	C10—C12—H12C	109.5

C4—P3—Pt1	111.81 (9)	H12A—C12—H12C	109.5
C10—P3—Pt1	117.37 (9)	H12B—C12—H12C	109.5
C6—C4—C5	110.0 (2)	C14—P13—C17	101.90 (12)
C6—C4—P3	112.7 (2)	C14—P13—Pt1 ⁱ	106.05 (9)
C5—C4—P3	109.69 (19)	C17—P13—Pt1 ⁱ	111.17 (9)
C6—C4—H4	108.1	C14—P13—Pt1	119.34 (9)
C5—C4—H4	108.1	C17—P13—Pt1	115.63 (9)
P3—C4—H4	108.1	Pt1 ⁱ —P13—Pt1	102.53 (3)
C4—C5—H5A	109.5	C16—C14—C15	110.1 (2)
C4—C5—H5B	109.5	C16—C14—P13	116.01 (19)
H5A—C5—H5B	109.5	C15—C14—P13	110.48 (19)
C4—C5—H5C	109.5	C16—C14—H14	106.6
H5A—C5—H5C	109.5	C15—C14—H14	106.6
H5B—C5—H5C	109.5	P13—C14—H14	106.6
C4—C6—H6A	109.5	C14—C15—H15A	109.5
C4—C6—H6B	109.5	C14—C15—H15B	109.5
H6A—C6—H6B	109.5	H15A—C15—H15B	109.5
C4—C6—H6C	109.5	C14—C15—H15C	109.5
H6A—C6—H6C	109.5	H15A—C15—H15C	109.5
H6B—C6—H6C	109.5	H15B—C15—H15C	109.5
C9—C7—C8	111.3 (2)	C14—C16—H16A	109.5
C9—C7—P3	112.72 (19)	C14—C16—H16B	109.5
C8—C7—P3	115.95 (19)	H16A—C16—H16B	109.5
C9—C7—H7	105.3	C14—C16—H16C	109.5
C8—C7—H7	105.3	H16A—C16—H16C	109.5
P3—C7—H7	105.3	H16B—C16—H16C	109.5
C7—C8—H8A	109.5	C18—C17—C19	109.8 (2)
C7—C8—H8B	109.5	C18—C17—P13	114.37 (19)
H8A—C8—H8B	109.5	C19—C17—P13	109.31 (19)
C7—C8—H8C	109.5	C18—C17—H17	107.7
H8A—C8—H8C	109.5	C19—C17—H17	107.7
H8B—C8—H8C	109.5	P13—C17—H17	107.7
C7—C9—H9A	109.5	C17—C18—H18A	109.5
C7—C9—H9B	109.5	C17—C18—H18B	109.5
H9A—C9—H9B	109.5	H18A—C18—H18B	109.5
C7—C9—H9C	109.5	C17—C18—H18C	109.5
H9A—C9—H9C	109.5	H18A—C18—H18C	109.5
H9B—C9—H9C	109.5	H18B—C18—H18C	109.5
C12—C10—C11	110.6 (2)	C17—C19—H19A	109.5
C12—C10—P3	117.8 (2)	C17—C19—H19B	109.5
C11—C10—P3	111.9 (2)	H19A—C19—H19B	109.5
C12—C10—H10	105.1	C17—C19—H19C	109.5
C11—C10—H10	105.1	H19A—C19—H19C	109.5
P3—C10—H10	105.1	H19B—C19—H19C	109.5
P13 ⁱ —Pt1—P3—C7	31.4 (2)	C7—P3—C10—C11	48.1 (2)
P13—Pt1—P3—C7	-135.55 (10)	C4—P3—C10—C11	156.87 (19)
P13 ⁱ —Pt1—P3—C4	-82.7 (2)	Pt1—P3—C10—C11	-79.0 (2)

P13—Pt1—P3—C4	110.32 (10)	P3—Pt1—P13—C14	-65.19 (10)
P13 ⁱ —Pt1—P3—C10	157.11 (18)	P13 ⁱ —Pt1—P13—C14	116.73 (10)
P13—Pt1—P3—C10	-9.88 (11)	P3—Pt1—P13—C17	56.96 (10)
C7—P3—C4—C6	-51.8 (2)	P13 ⁱ —Pt1—P13—C17	-121.13 (10)
C10—P3—C4—C6	-164.7 (2)	P3—Pt1—P13—Pt1 ⁱ	178.09 (2)
Pt1—P3—C4—C6	67.6 (2)	P13 ⁱ —Pt1—P13—Pt1 ⁱ	0.0
C7—P3—C4—C5	-174.7 (2)	C17—P13—C14—C16	-57.0 (2)
C10—P3—C4—C5	72.3 (2)	Pt1 ⁱ —P13—C14—C16	-173.41 (19)
Pt1—P3—C4—C5	-55.3 (2)	Pt1—P13—C14—C16	71.7 (2)
C4—P3—C7—C9	167.9 (2)	C17—P13—C14—C15	176.81 (19)
C10—P3—C7—C9	-82.4 (2)	Pt1 ⁱ —P13—C14—C15	60.41 (19)
Pt1—P3—C7—C9	48.1 (2)	Pt1—P13—C14—C15	-54.5 (2)
C4—P3—C7—C8	-62.2 (2)	C14—P13—C17—C18	-53.1 (2)
C10—P3—C7—C8	47.5 (2)	Pt1 ⁱ —P13—C17—C18	59.6 (2)
Pt1—P3—C7—C8	178.02 (18)	Pt1—P13—C17—C18	175.91 (17)
C7—P3—C10—C12	-81.7 (2)	C14—P13—C17—C19	-176.60 (19)
C4—P3—C10—C12	27.1 (3)	Pt1 ⁱ —P13—C17—C19	-63.98 (19)
Pt1—P3—C10—C12	151.27 (19)	Pt1—P13—C17—C19	52.4 (2)

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