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Crystal structures of three complexes of zinc chloride with tri-*tert*-butylphosphane

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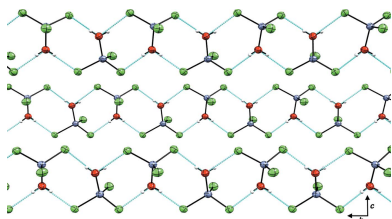
Under anhydrous conditions and in the absence of a Lewis-base solvent, a zinc chloride complex with tri-*tert*-butylphosphane as the μ -bridged dimer is formed, *viz.* di- μ -chlorido-bis[chloridobis(tri-*tert*-butylphosphane)zinc], $[\text{ZnCl}_4(\text{C}_{12}\text{H}_{27}\text{P})_2]$, (**1**), which features a nearly square-shaped $(\text{ZnCl})_2$ cyclic core and whose Cl atoms interact weakly with C–H groups on the phosphane ligand. In the presence of THF, monomeric dichlorido(tetrahydrofuran- κO)-(tri-*tert*-butylphosphane- κP)zinc, $[\text{ZnCl}_2(\text{C}_4\text{H}_8\text{O})(\text{C}_{12}\text{H}_{27}\text{P})]$ or $[\text{P}(\text{tBu}_3)(\text{THF})\text{ZnCl}_2]$, (**2**), is formed. This slightly distorted tetrahedral Zn complex has weak C–H \cdots Cl interactions between the Cl atoms and phosphane and THF C–H groups. Under ambient conditions, the hydrolysed complex tri-*tert*-butylphosphonium aquatrachloridozincate 1,2-dichloroethane monosolvate, $(\text{C}_{12}\text{H}_{28}\text{P})[\text{ZnCl}_3(\text{H}_2\text{O})]\cdot\text{C}_2\text{H}_4\text{Cl}_2$ or $[\text{HPtBu}_3]^+[(\text{H}_2\text{O})\text{ZnCl}_3]^- \cdot \text{C}_2\text{H}_4\text{Cl}_2$, (**3**), is formed. This complex forms chains of $[(\text{H}_2\text{O})\text{ZnCl}_3]^-$ anions from hydrogen-bonding interactions between the water H atoms and Cl atoms that propagate along the *b* axis.

1. Chemical context

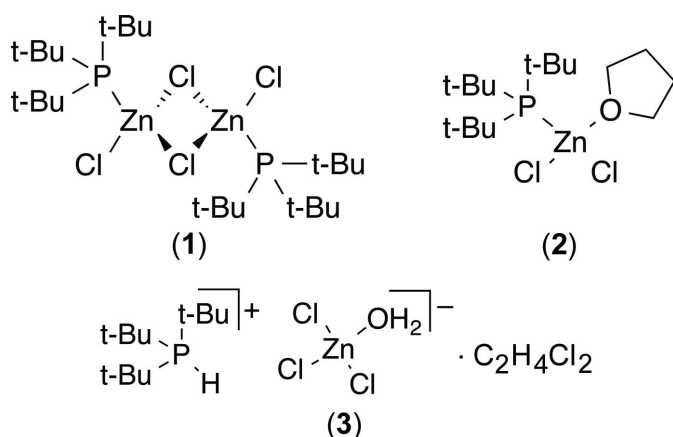
Tri-*tert*-butylphosphane PtBu_3 is a bulky, weak Lewis base. It has found considerable utility as a ligand for Pd-catalysed cross couplings (Fu, 2008). More recently, its reactivity with bulky Lewis acids to form the so-called ‘frustrated Lewis pairs’ has opened up new avenues of chemical reactivity (Stephan & Erker, 2015). Lewis acidic complexes containing zinc have been used as ring-opening polymerization catalysts (Wu *et al.*, 2006). The reactivity of PtBu_3 with weak transition metal Lewis acids has been less well explored. The reaction of PtBu_3 with ZnCl_2 has been reported (Goel & Ogini, 1977), but without structural characterization. Therein, $[(\text{PtBu}_3)(\text{ZnCl}_2)]$ (**1**) was proposed to exist as the di- μ -chlorido-bridged dimer based on molecular weight measurements. We describe the structure of two complexes of $[(\text{PtBu}_3)(\text{ZnCl}_2)]$: the aforementioned μ -bridged dimer (**1**), and the monomeric THF complex (**2**). The complex is sensitive to ambient moisture, and hydrolyses to form the hydrolysis product $[\text{HPtBu}_3]^+[(\text{H}_2\text{O})\text{ZnCl}_3]^- \cdot \text{C}_2\text{H}_4\text{Cl}_2$ (**3**) under ambient conditions from a 1,2-dichloroethane solution. The related compound $[\text{HPtBu}_3]^+[(\text{H}_2\text{O})\text{ZnI}_3]^-$ was reported from the preparation of PtBu_3 and ZnI_2 in benzene under ambient conditions (Goel & Ogini, 1977), but no structural data were reported.

2. Structural commentary

Compound (**1**) is a neutral μ -bridged dimer with one PtBu_3 per zinc atom. The asymmetric unit is one half of (**1**) with the



other half related by inversion symmetry (Fig. 1). The coordination sphere of the Zn is filled with two Cl atoms, one of which, Cl1, is μ -bound to both Zn atoms of (1) [$\text{Zn1}-\text{Cl1} = 2.3703(13) \text{ \AA}$] and the other, Cl2, is bound to only one Zn [$\text{Zn1}-\text{Cl2} = 2.2133(14) \text{ \AA}$]. The four-membered ring consisting of two Zn1 and two Cl1 is planar. The bond angles are only slightly distorted from the ideal values of 90° [$\text{Cl1}-\text{Zn1}-\text{Cl1}^i = 90.98(4)$, $\text{Zn1}-\text{Cl1}-\text{Zn1}^i = 89.02(4)^\circ$; symmetry code: (i) $-x, -y + 1, -z + 1$]. The Zn atom is a distorted tetrahedron [$\text{P1}-\text{Zn1}-\text{Cl2} = 117.30(5)$, $\text{P1}-\text{Zn1}-\text{Cl1} = 112.62(5)^\circ$; $\tau_4 = 0.92$ (Yang *et al.*, 2007)]. The $\text{Zn}\cdots\text{Zn}^i$ distance is $3.3189(10) \text{ \AA}$ [symmetry code: (i) $-x, -y + 1, -z + 1$]. The Zn–P bond [$\text{Zn1}-\text{P1} = 2.3859(13) \text{ \AA}$] for (1) is in line with other Zn–trialkylphosphane complexes.



Compound (2) is a neutral tetrahedral Zn complex with two Cl ligands, one PrBu_3 ligand, and one THF ligand (Fig. 2). The Zn–P bond length [$\text{Zn1}-\text{P1} = 2.4167(9) \text{ \AA}$] is in line with other Zn–trialkylphosphane complexes. The Zn–Cl bond lengths are very similar [$\text{Zn1}-\text{Cl1} = 2.2370(13)$, $\text{Zn1}-\text{Cl2} = 2.2301(13) \text{ \AA}$]. The Zn environment is slightly distorted

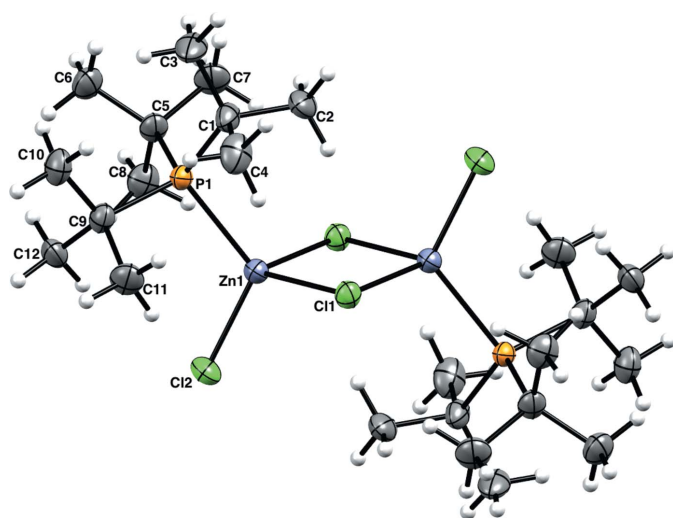


Figure 1
The molecular structure of (1), showing 50% probability ellipsoids for non-H atoms and spheres of arbitrary size for H atoms. The unlabeled atoms are related by the symmetry operator ($-x, 1 - y, 1 - z$).

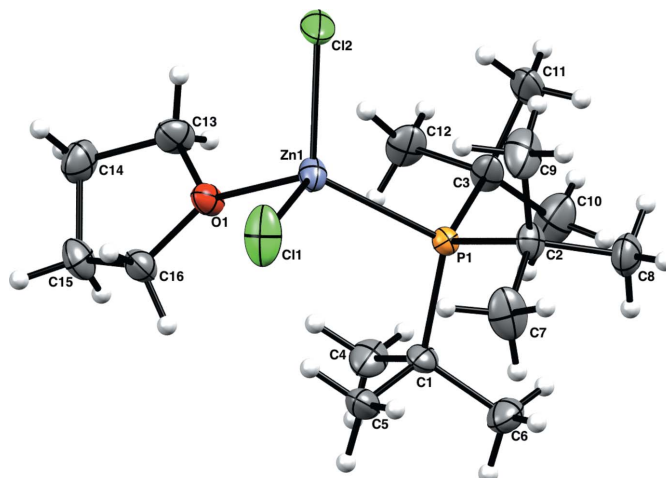


Figure 2
The molecular structure of (2), showing 50% probability ellipsoids for non-H atoms and spheres of arbitrary size for H atoms.

tetrahedral due to the steric influence of the PrBu_3 ligand ($\tau_4 = 0.94$) (Yang *et al.*, 2007).

The asymmetric unit of compound (3) (Fig. 3) comprises three $[\text{HPtBu}_3]^+ [(\text{H}_2\text{O})\text{ZnCl}_3]^-$ ion pairs, along with three 1,2-dichloroethane molecules not related by symmetry ($Z' = 3$). The three groups are similar in structure, connectivity, and supramolecular interactions; despite this, no additional crystallographic symmetry or twinning was found using *PLATON* (Spek, 2009). The $[(\text{H}_2\text{O})\text{ZnCl}_3]^-$ ion has some interesting properties. Two of the three Cl atoms in each $[(\text{H}_2\text{O})\text{ZnCl}_3]^-$ ion are involved in hydrogen bonding with nearby water ligands. The Zn–Cl bonds [$\text{Zn1}-\text{Cl1} = 2.2690(10)$, $\text{Zn1}-\text{Cl2} = 2.2666(10)$, $\text{Zn1}-\text{Cl3} = 2.2219(11)$, $\text{Zn2}-\text{Cl4} = 2.2203(11) \text{ \AA}$, $\text{Zn2}-\text{Cl5} = 2.2666(10) \text{ \AA}$, $\text{Zn2}-\text{Cl6} = 2.2699(10)$, $\text{Zn3}-\text{Cl7} = 2.2199(11) \text{ \AA}$, $\text{Zn3}-\text{Cl8} = 2.2695(10)$, $\text{Zn3}-\text{Cl9} = 2.2686(10) \text{ \AA}$] are affected significantly by the hydrogen bonding. The Zn–Cl bonds involved in hydrogen bonding are significantly longer (by *ca.* 0.04 \AA) than the Zn–Cl bonds not involved in hydrogen bonding. The Zn–OH₂ bonds [$\text{Zn1}-\text{O1} = 2.024(3)$, $\text{Zn2}-\text{O2} = 2.025(3)$,

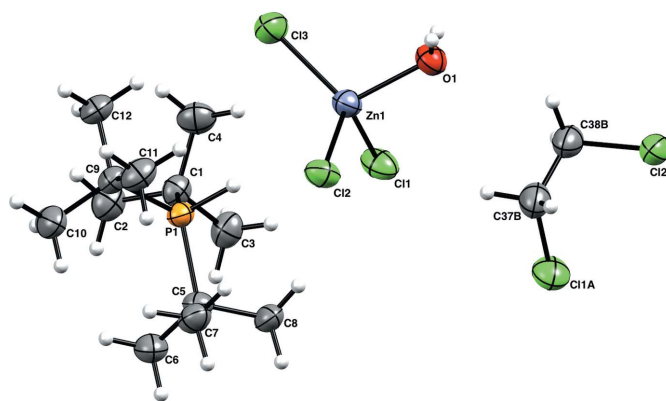


Figure 3
The molecular structure of (3), showing one of the three molecules of the asymmetric unit ($Z' = 3$) showing 50% probability ellipsoids for non-H atoms and spheres of arbitrary size for H atoms.

Table 1
 Hydrogen-bond geometry (Å, °) for (1).

| $D-H\cdots A$ | $D-H$ | $H\cdots A$ | $D\cdots A$ | $D-H\cdots A$ |
|----------------------|-------|-------------|-------------|---------------|
| $C3-H3C\cdots Cl1^i$ | 0.98 | 2.88 | 3.479 (6) | 120 |

 Symmetry code: (i) $-x - \frac{1}{2}, y - \frac{1}{2}, z$.

Table 2
 Hydrogen-bond geometry (Å, °) for (2).

| $D-H\cdots A$ | $D-H$ | $H\cdots A$ | $D\cdots A$ | $D-H\cdots A$ |
|---------------------------|-------|-------------|-------------|---------------|
| $C12-H12A\cdots Cl2^i$ | 0.98 | 2.90 | 3.819 (5) | 157 |
| $C15-H15A\cdots Cl1^{ii}$ | 0.99 | 2.94 | 3.747 (5) | 140 |

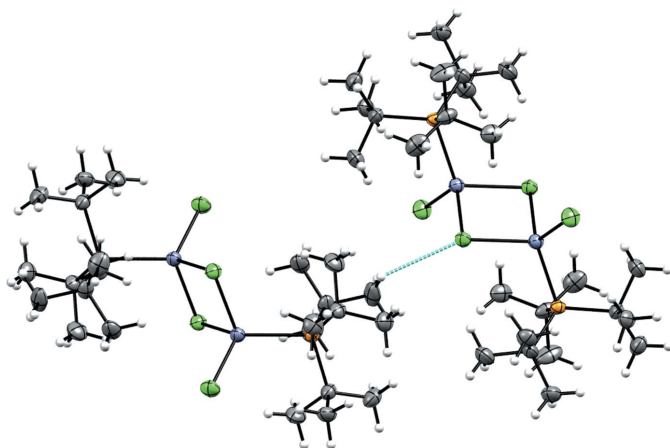
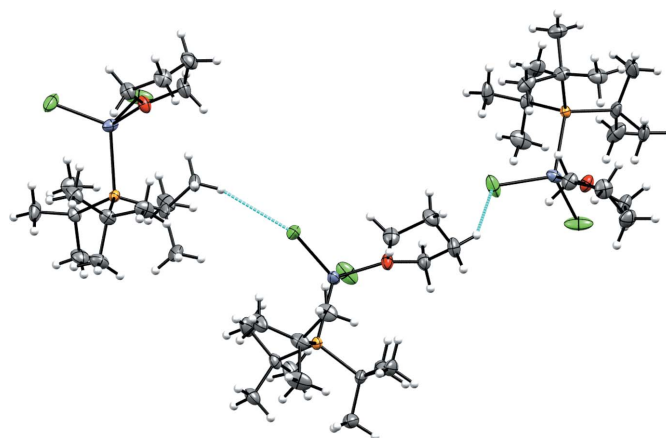
 Symmetry codes: (i) $-x + \frac{3}{2}, y + \frac{1}{2}, z - \frac{1}{2}$; (ii) $-x + 2, -y, z - \frac{1}{2}$.

Table 3
 Hydrogen-bond geometry (Å, °) for (3).

| $D-H\cdots A$ | $D-H$ | $H\cdots A$ | $D\cdots A$ | $D-H\cdots A$ |
|---------------------------|----------|-------------|-------------|---------------|
| $O1-H1OA\cdots Cl2^i$ | 0.82 (2) | 2.37 (3) | 3.107 (3) | 150 (4) |
| $O1-H1OB\cdots Cl1^{ii}$ | 0.82 (2) | 2.27 (2) | 3.086 (3) | 173 (4) |
| $O2-H2OA\cdots Cl9^{iii}$ | 0.82 (2) | 2.33 (2) | 3.120 (3) | 161 (4) |
| $O2-H2OB\cdots Cl8^{iv}$ | 0.82 (2) | 2.28 (2) | 3.095 (3) | 177 (4) |
| $O3-H3OA\cdots Cl5^{iii}$ | 0.85 (2) | 2.30 (2) | 3.100 (3) | 158 (4) |
| $O3-H3OB\cdots Cl6^{iv}$ | 0.83 (2) | 2.28 (2) | 3.106 (3) | 173 (4) |

 Symmetry codes: (i) $-x + \frac{1}{2}, y + \frac{1}{2}, -z + \frac{3}{2}$; (ii) $-x + \frac{1}{2}, y - \frac{1}{2}, -z + \frac{3}{2}$; (iii) $-x + \frac{3}{2}, y + \frac{1}{2}, -z + \frac{3}{2}$; (iv) $-x + \frac{3}{2}, y - \frac{1}{2}, -z + \frac{3}{2}$.

$Zn3-O3 = 2.028$ (3) Å] are all within one s.u. of the average tetrahedral $Zn-OH_2$ bond length of 2.00 (4) Å. The coordination environments of the Zn atoms in the $[(H_2O)ZnCl_3]^-$ anions are all slightly distorted tetrahedral [$\tau_4(Zn1) = 0.92$, $\tau_4(Zn2) = 0.93$, $\tau_4(Zn3) = 0.93$] (Yang *et al.*, 2007). The phosphonium hydrogen atoms were found in a difference map and restrained to be similar to each other; the average P-H bond length is 1.31 (3) Å. The 1,2-dichloroethane solvent has significantly larger displacement parameters than the other two moieties, indicating disorder. Thus, each solvent molecule


Figure 4
 The weak $H\cdots Cl$ interaction in (1) with short contact shown in cyan. The second molecule (left) is related to the first by the symmetry operation $(\frac{3}{2} - x, -\frac{1}{2} + y, \frac{1}{2} + z)$.

Figure 5
 The weak $H\cdots Cl$ interactions in (2) with short contacts shown in cyan. The left molecule is related to the middle one by the symmetry operation $(2 - x, -y, \frac{1}{2} + z)$, and the right molecule is related to the middle one by the symmetry operation $(\frac{3}{2} - x, -\frac{1}{2} + y, \frac{1}{2} + z)$.

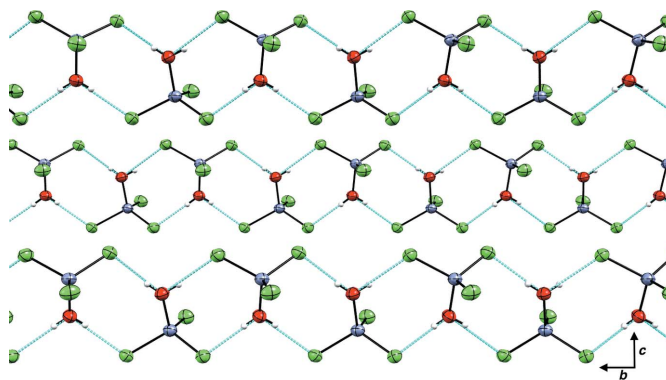
was modeled over two discrete positions (see *Refinement* section).

3. Supramolecular features

Supramolecular features of (1) form from weak $C1-H3C\cdots Cl1^i$ interactions (Fig. 4 and Table 1), which creates layers in the ab plane that stack along the c axis.

The supramolecular features of (2) are also based on weak interactions. There are weak $C15-H15A\cdots Cl1^{ii}$ interactions as well as weak $C12-H12A\cdots Cl2^i$ interactions (Fig. 5 and Table 2). Together the weak interactions, where each Cl atom is an acceptor, create a three-dimensional packing structure.

The hydrogen atoms of the water ligands in (3) undergo hydrogen-bonding interactions with nearby chloride ligands of the $[(H_2O)ZnCl_3]^-$ anion, forming chains that propagate along the b -axis direction (Fig. 6 and Table 3). The chains in each layer are staggered by half a unit cell along the b axis. The orientation of the P-H bond relative to the $[(H_2O)ZnCl_3]^-$


Figure 6
 Chains of the three $[(H_2O)ZnCl_3]^-$ ions formed from the hydrogen bonds between $Zn-Cl$ and water ligands in (3), viewed along the a axis, with hydrogen-bond interactions shown in cyan. $[HPtBu_3]^+$ ions and 1,2-DCE solvent molecules are not shown.

ion is optimized for steric interactions; that is, the P–H hydrogen atom is oriented toward the center of the Zn tetrahedron surrounded by three Cl atoms, suggesting a nucleophilic-type protonation of the phosphane, with the water ligand pointing away from the P–H bond. Each *tert*-butyl group is staggered slightly relative to the positions of the Cl atoms. In this arrangement, there are no hydrogen-bonding interactions involving the phosphonium hydrogen. This arrangement also optimizes the ion contact between the phosphonium cations and $[(\text{H}_2\text{O})\text{ZnCl}_3]^-$ anions. The disorder of the solvent molecules suggests no or at best weak interactions between the solvent and hosts; indeed, none can be found.

4. Database survey

Zinc chloride-bulky phosphane compounds tend to form ZnCl_2 -monophosphane complexes. Two closely related compounds include a ZnCl_2 -phosphane dimer (LUZVEI; Liang *et al.*, 2010), and a ZnCl_2 -bulky NHC dimer (XONKUI; Flidel *et al.*, 2014). In both cases, the molecular geometries are very similar to that of **(1)**. A search of the Cambridge Structural Database (CSD; Groom & Allen, 2014) returned 110 dimeric complexes with the general formula $[\text{TM}(\text{PR}_3\text{Cl}(\mu\text{-Cl}))_2]$ (TM = transition metal). Most of these entries are complexes of group 10 metals (Ni, Pd, Pt), but, due to their different electron configuration to Zn, these tend to be mostly planar complexes. As expected, another Group 11 transition metal, Hg, forms similar complexes as Zn; there are 14 entries in the CSD with the formula $[\text{Hg}(\text{PR}_3\text{Cl}(\mu\text{-Cl}))_2]$. Notably similar complexes to **(1)** include $[\text{Hg}(\text{P}(\text{cyclohexyl})_3\text{Cl}(\mu\text{-Cl}))_2]$ (BULSOQ; Bell *et al.*, 1983) and $[\text{Hg}(\text{P}(2,5\text{-}(\text{OMe})_2\text{Ph})_3\text{Cl}(\mu\text{-Cl}))_2]$ (WONKEP; Bell *et al.*, 2000). Interestingly, there are no similar entries in the CSD that contain Cd.

There are three compounds in the CSD with the general formula $[(\text{thf})\text{TMCl}_2]$. There is a compound closely related to **(2)**, $[\text{ZnCl}_2(\text{THF})(\text{P}(\text{SnMe}_3)_3)]$ (ASEBUV; Fuhr & Fenske, 2004). Like **(2)**, it forms from the reaction of ZnCl_2 with $\text{P}(\text{SnMe}_3)_3$ in THF. The other two compounds are complexes of Pd (FIRDAN, Cohen *et al.*, 2014; UHUDAC, Kim & Verkade, 2003).

Besides Goel's report on the hydrolysis of $[(\text{PtBu}_3)(\text{ZnI}_2)]$, there are no other reports on the hydrolysis of zinc-phosphane complexes to form phosphonium salts. The $[(\text{H}_2\text{O})\text{ZnCl}_3]^-$ ion is relatively uncommon in the CSD: there are 19 entries containing such an ion. However, there is one report of the hydrolysis of a triphenylphosphinomethyl– ZnCl_2 dimer (CORRAD; Pattacini *et al.*, 2009) with water to form $[\text{Ph}_3\text{PMe}]^+[(\text{H}_2\text{O})\text{ZnCl}_3]^-$ (CORQEG; Pattacini *et al.*, 2009). The $[(\text{H}_2\text{O})\text{ZnCl}_3]^-$ ions also form chains similar to **(3)** arising from hydrogen-bonding interactions between the two H atoms of the water ligand with two of the three Cl atoms of the ion. Likewise, the lengthening of the Zn–Cl bond as a result of hydrogen bonding as seen in **(3)** is also observed here. There are 67 entries in the CSD containing the moiety $[\text{HPtBu}_3]^+$, none with Zn-containing counter-ions. Most of the counter-

ions of $[\text{HPtBu}_3]^+$ reported therein are anionic tetrahedral borates arising from frustrated Lewis pair reactivity.

5. Synthesis and crystallization

The synthesis of **(1)** has been reported (Goel & Ogini, 1977); the methods reported here are modified from the original report. Crystals of **(1)** were grown from slow diffusion of pentane into an equimolar solution of ZnCl_2 and PtBu_3 in $(\text{CH}_2\text{Cl})_2$ at 243 K under an atmosphere of Ar gas. Crystals of **(2)** were grown from slow diffusion of pentane into an equimolar solution of ZnCl_2 and PtBu_3 in THF at 243 K under an atmosphere of Ar gas. Crystals of **(3)** were grown from slow diffusion of pentane into an equimolar solution of ZnCl_2 and PtBu_3 in 1,2-dichloroethane (1,2-DCE) at room temperature under ambient conditions.

6. Refinement

Compound **(1)**: A structural model consisting of one-half of **(1)** was developed. Methyl H atom positions, $R\text{--CH}_3$, were optimized by rotation about $R\text{--C}$ bonds with idealized C–H, $R\text{--H}$ and $\text{H}\cdots\text{H}$ distances. For all H atoms, $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{carrier})$.

Compound **(2)**: A structural model consisting of the host molecule was developed. The coordinating Cl atoms had elongated anisotropic displacement parameters in one direction; however, splitting the Cl positions did not significantly improve the model so it was removed from the final model. Methyl H atom positions, $R\text{--CH}_3$, were optimized by rotation about $R\text{--C}$ bonds with idealized C–H, $R\text{--H}$ and $\text{H}\cdots\text{H}$ distances. Remaining H atoms were included as riding idealized contributors. $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$ for methyl atoms and $1.2U_{\text{eq}}(\text{carrier})$ for remaining H atoms. On the basis of 1704 unmerged Friedel opposites, the minor component occupancy of the inversion twin was 0.206 (13) (Flack & Bernardinelli, 2000).

Compound **(3)**: A structural model consisting of three ion pairs and three 1,2-DCE solvent molecules per asymmetric unit was developed. Methyl H atom positions, $R\text{--CH}_3$, were optimized by rotation about $R\text{--C}$ bonds with idealized C–H, $R\text{--H}$ and $\text{H}\cdots\text{H}$ distances. Water H atoms and phosphonium H atoms were identified in a difference Fourier map and refined. Water atom H atoms were restrained (s.u. 0.02) to a bond length of 0.84 Å. Phosphonium H atoms were restrained to be similar (s.u. 0.01). Remaining H atoms were included as riding idealized contributors. $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$ for methyl atoms and $1.2U_{\text{eq}}(\text{carrier})$ for remaining H atoms. The 1,2-DCE molecules had significantly larger displacement parameters; thus, these moieties were modeled as disordered over two discrete positions. Enhanced rigid-bond restraints (s.u. 0.004) (Thorn *et al.*, 2012) were imposed on displacement parameters for all disordered sites and similar displacement amplitudes (s.u. 0.01) were imposed on disordered sites overlapping by less than the sum of van der Waals radii. In addition, the C–Cl bonds in the 1,2-DCE molecules and the C–C bonds were restrained to be similar (s.u. 0.01). The

Table 4
Experimental details.

| | (1) | (2) | (3) |
|---|---|--|---|
| Crystal data | | | |
| Chemical formula | [ZnCl ₄ (C ₁₂ H ₂₇ P) ₂] | [ZnCl ₂ (C ₄ H ₈ O)(C ₁₂ H ₂₇ P)] | (C ₁₂ H ₂₈ P)[ZnCl ₃ (H ₂ O)]·C ₂ H ₄ Cl ₂ |
| <i>M_r</i> | 677.15 | 410.68 | 492.00 |
| Crystal system, space group | Orthorhombic, <i>Pbca</i> | Orthorhombic, <i>Pna</i> 2 ₁ | Monoclinic, <i>P</i> 2 ₁ / <i>n</i> |
| Temperature (K) | 193 | 193 | 123 |
| <i>a</i> , <i>b</i> , <i>c</i> (Å) | 14.6408 (16), 12.9891 (14), 16.8190 (18) | 26.4580 (11), 8.9281 (4), 8.5790 (4) | 25.3722 (5), 8.5841 (2), 32.912 (2) |
| α , β , γ (°) | 90, 90, 90 | 90, 90, 90 | 90, 98.909 (7), 90 |
| <i>V</i> (Å ³) | 3198.5 (6) | 2026.53 (16) | 7081.7 (6) |
| <i>Z</i> | 4 | 4 | 12 |
| Radiation type | Mo <i>K</i> α | Mo <i>K</i> α | Cu <i>K</i> α |
| μ (mm ⁻¹) | 1.95 | 1.55 | 7.28 |
| Crystal size (mm) | 0.22 × 0.18 × 0.09 | 0.47 × 0.46 × 0.27 | 0.18 × 0.05 × 0.02 |
| Data collection | | | |
| Diffractometer | Bruker APEXII CCD area detector | Bruker APEXII CCD area detector | Rigaku CCD area detector |
| Absorption correction | Integration (<i>SADABS</i> ; Bruker, 2008) | Integration (<i>SADABS</i> ; Bruker, 2008) | Multi-scan (<i>CrystalClear</i> ; Rigaku, 2010) |
| <i>T</i> _{min} , <i>T</i> _{max} | 0.846, 0.999 | 0.580, 0.754 | 0.354, 0.868 |
| No. of measured, independent and observed [<i>I</i> > 2 σ (<i>I</i>)] reflections | 32092, 2942, 1836 | 23933, 3698, 3581 | 67698, 11604, 7131 |
| <i>R</i> _{int} (<i>sin</i> θ / λ) _{max} (Å ⁻¹) | 0.169 0.604 | 0.038 0.603 | 0.083 0.581 |
| Refinement | | | |
| <i>R</i> [<i>F</i> ² > 2 σ (<i>F</i> ²)], <i>wR</i> (<i>F</i> ²), <i>S</i> | 0.045, 0.099, 1.00 | 0.031, 0.075, 1.14 | 0.049, 0.142, 0.98 |
| No. of reflections | 2942 | 3698 | 11604 |
| No. of parameters | 154 | 200 | 760 |
| No. of restraints | 0 | 1 | 462 |
| H-atom treatment | H-atom parameters constrained | H-atom parameters constrained | H atoms treated by a mixture of independent and constrained refinement |
| $\Delta\rho_{\max}$, $\Delta\rho_{\min}$ (e Å ⁻³) | 0.44, -0.42 | 0.69, -0.30 | 0.51, -0.70 |
| Absolute structure | – | Refined as an inversion twin | – |
| Absolute structure parameter | – | 0.206 (18) | – |

Computer programs: *APEX2* (Bruker, 2004), *CrystalClear* (Rigaku, 2010), *SAINT* and *XPREP* (Bruker, 2005), *SADABS* (Bruker, 2008) and *TWINABS* (Bruker, 2008), *SHELXS97* (Sheldrick, 2008), *SHELXL2014* (Sheldrick, 2015) and *OLEX2* (Bourhis *et al.*, 2015).

major:minor occupancy factor ratios for the three 1,2-DCE molecules are 0.52 (3):0.48 (3), 0.119 (7):0.881 (7), and 0.38 (3):0.62 (3). Crystal data, data collection and structure refinement details are summarized in Table 4.

Acknowledgements

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supporting information

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Crystal structures of three complexes of zinc chloride with tri-*tert*-butylphosphane

Aaron D. Finke, Danielle L. Gray and Jeffrey S. Moore

Computing details

Data collection: *APEX2* (Bruker, 2004) for (1), (2); *CrystalClear* (Rigaku, 2010) for (3). Cell refinement: *SAINT* (Bruker, 2005) for (1), (2); *CrystalClear* (Rigaku, 2010) for (3). Data reduction: *SAINT* (Bruker, 2005) for (1); *SAINT* (Bruker, 2005), *XPREP* (Bruker, 2005), *SADABS* (Bruker, 2008) and *TWINABS* (Bruker, 2008) for (2); *CrystalClear* (Rigaku, 2010) for (3). For all compounds, program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL2014* (Sheldrick, 2015); molecular graphics: *OLEX2* (Bourhis *et al.*, 2015); software used to prepare material for publication: *OLEX2* (Bourhis *et al.*, 2015).

(1) Di- μ -chlorido-bis[chloridobis(tri-*tert*-butylphosphane)zinc]

Crystal data

[ZnCl₄(C₁₂H₂₇P)₂]

$M_r = 677.15$

Orthorhombic, *Pbca*

$a = 14.6408$ (16) Å

$b = 12.9891$ (14) Å

$c = 16.8190$ (18) Å

$V = 3198.5$ (6) Å³

$Z = 4$

$F(000) = 1424$

$D_x = 1.406$ Mg m⁻³

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

Cell parameters from 1388 reflections

$\theta = 2.4$ – 18.9°

$\mu = 1.95$ mm⁻¹

$T = 193$ K

Block, colourless

$0.22 \times 0.18 \times 0.09$ mm

Data collection

Bruker APEXII CCD area-detector diffractometer

Radiation source: Sealed Tube

Graphite monochromator

Detector resolution: 8.33 pixels mm⁻¹

φ and ω scans

Absorption correction: integration (*SADABS*; Bruker, 2008)

$T_{\min} = 0.846$, $T_{\max} = 0.999$

32092 measured reflections

2942 independent reflections

1836 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.169$

$\theta_{\max} = 25.4^\circ$, $\theta_{\min} = 2.4^\circ$

$h = -16 \rightarrow 17$

$k = -15 \rightarrow 15$

$l = -20 \rightarrow 20$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.045$

$wR(F^2) = 0.099$

$S = 1.00$

2942 reflections

154 parameters

0 restraints

Hydrogen site location: inferred from neighbouring sites

H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0282P)^2 + 3.8822P]$$

where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$

$$\Delta\rho_{\max} = 0.44 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\min} = -0.42 \text{ e } \text{\AA}^{-3}$$

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.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

| | x | y | z | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|------|--------------|--------------|-------------|----------------------------------|
| Zn1 | 0.01390 (4) | 0.41919 (4) | 0.42454 (3) | 0.02543 (17) |
| Cl1 | -0.07300 (8) | 0.57038 (9) | 0.44450 (7) | 0.0287 (3) |
| Cl2 | 0.12470 (9) | 0.45042 (10) | 0.33832 (8) | 0.0372 (3) |
| P1 | -0.07908 (9) | 0.27135 (9) | 0.40137 (7) | 0.0229 (3) |
| C1 | -0.1263 (4) | 0.2197 (4) | 0.4984 (3) | 0.0340 (13) |
| C2 | -0.1597 (4) | 0.3112 (4) | 0.5491 (3) | 0.0412 (14) |
| H2A | -0.1840 | 0.2855 | 0.5997 | 0.062* |
| H2B | -0.2078 | 0.3482 | 0.5204 | 0.062* |
| H2C | -0.1086 | 0.3579 | 0.5595 | 0.062* |
| C3 | -0.2037 (4) | 0.1424 (4) | 0.4882 (3) | 0.0505 (17) |
| H3A | -0.2239 | 0.1183 | 0.5405 | 0.076* |
| H3B | -0.1821 | 0.0837 | 0.4568 | 0.076* |
| H3C | -0.2549 | 0.1754 | 0.4606 | 0.076* |
| C4 | -0.0504 (4) | 0.1693 (4) | 0.5459 (3) | 0.0464 (16) |
| H4A | -0.0733 | 0.1509 | 0.5988 | 0.070* |
| H4B | 0.0008 | 0.2175 | 0.5513 | 0.070* |
| H4C | -0.0297 | 0.1071 | 0.5184 | 0.070* |
| C5 | -0.1755 (3) | 0.3087 (4) | 0.3323 (3) | 0.0322 (13) |
| C6 | -0.2210 (4) | 0.2184 (4) | 0.2886 (3) | 0.0429 (15) |
| H6A | -0.2687 | 0.2448 | 0.2532 | 0.064* |
| H6B | -0.2483 | 0.1713 | 0.3275 | 0.064* |
| H6C | -0.1750 | 0.1815 | 0.2573 | 0.064* |
| C7 | -0.2506 (4) | 0.3675 (4) | 0.3768 (3) | 0.0457 (15) |
| H7A | -0.2951 | 0.3944 | 0.3386 | 0.069* |
| H7B | -0.2233 | 0.4248 | 0.4064 | 0.069* |
| H7C | -0.2814 | 0.3209 | 0.4140 | 0.069* |
| C8 | -0.1373 (4) | 0.3861 (4) | 0.2711 (3) | 0.0390 (14) |
| H8A | -0.1847 | 0.4026 | 0.2320 | 0.059* |
| H8B | -0.0845 | 0.3557 | 0.2440 | 0.059* |
| H8C | -0.1184 | 0.4492 | 0.2985 | 0.059* |
| C9 | -0.0043 (3) | 0.1705 (3) | 0.3523 (3) | 0.0301 (12) |
| C10 | -0.0439 (4) | 0.0616 (4) | 0.3532 (3) | 0.0405 (14) |
| H10A | -0.0025 | 0.0150 | 0.3250 | 0.061* |
| H10B | -0.1036 | 0.0617 | 0.3268 | 0.061* |
| H10C | -0.0510 | 0.0385 | 0.4083 | 0.061* |
| C11 | 0.0904 (3) | 0.1705 (4) | 0.3942 (3) | 0.0424 (14) |

| | | | | |
|------|------------|------------|------------|-------------|
| H11A | 0.1301 | 0.1193 | 0.3690 | 0.064* |
| H11B | 0.0825 | 0.1533 | 0.4505 | 0.064* |
| H11C | 0.1181 | 0.2389 | 0.3895 | 0.064* |
| C12 | 0.0166 (4) | 0.2009 (4) | 0.2653 (3) | 0.0393 (14) |
| H12A | 0.0631 | 0.1546 | 0.2436 | 0.059* |
| H12B | 0.0392 | 0.2719 | 0.2637 | 0.059* |
| H12C | -0.0393 | 0.1956 | 0.2335 | 0.059* |

Atomic displacement parameters (Å²)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|-----|------------|------------|------------|-------------|-------------|-------------|
| Zn1 | 0.0231 (3) | 0.0292 (3) | 0.0240 (3) | -0.0022 (3) | 0.0020 (3) | -0.0028 (2) |
| C11 | 0.0262 (7) | 0.0331 (6) | 0.0267 (7) | 0.0034 (6) | -0.0036 (5) | -0.0036 (5) |
| C12 | 0.0343 (8) | 0.0456 (8) | 0.0318 (8) | -0.0039 (6) | 0.0110 (6) | -0.0018 (6) |
| P1 | 0.0229 (7) | 0.0272 (7) | 0.0185 (7) | -0.0020 (6) | -0.0010 (6) | -0.0017 (5) |
| C1 | 0.043 (3) | 0.039 (3) | 0.020 (3) | -0.013 (3) | 0.004 (3) | -0.004 (2) |
| C2 | 0.038 (3) | 0.059 (4) | 0.027 (3) | -0.011 (3) | 0.010 (3) | -0.007 (3) |
| C3 | 0.060 (4) | 0.054 (4) | 0.038 (4) | -0.032 (3) | 0.017 (3) | -0.005 (3) |
| C4 | 0.070 (4) | 0.043 (3) | 0.026 (3) | -0.004 (3) | -0.004 (3) | 0.008 (2) |
| C5 | 0.024 (3) | 0.039 (3) | 0.034 (3) | 0.006 (2) | -0.008 (3) | -0.007 (2) |
| C6 | 0.035 (3) | 0.054 (4) | 0.040 (3) | -0.001 (3) | -0.009 (3) | -0.012 (3) |
| C7 | 0.028 (3) | 0.058 (4) | 0.051 (4) | 0.007 (3) | -0.009 (3) | -0.005 (3) |
| C8 | 0.042 (4) | 0.044 (3) | 0.032 (3) | 0.003 (3) | -0.010 (3) | 0.004 (2) |
| C9 | 0.031 (3) | 0.027 (2) | 0.032 (3) | 0.003 (2) | -0.005 (3) | -0.006 (2) |
| C10 | 0.049 (4) | 0.032 (3) | 0.040 (3) | 0.005 (3) | -0.004 (3) | -0.008 (2) |
| C11 | 0.028 (3) | 0.047 (3) | 0.052 (4) | 0.010 (3) | -0.006 (3) | -0.006 (3) |
| C12 | 0.029 (3) | 0.054 (3) | 0.035 (3) | 0.003 (3) | 0.005 (3) | -0.015 (3) |

Geometric parameters (Å, °)

| | | | |
|----------------------|-------------|----------|-----------|
| Zn1—C11 ⁱ | 2.3703 (13) | C5—C8 | 1.544 (7) |
| Zn1—C11 | 2.3639 (13) | C6—H6A | 0.9800 |
| Zn1—C12 | 2.2133 (14) | C6—H6B | 0.9800 |
| Zn1—P1 | 2.3859 (13) | C6—H6C | 0.9800 |
| C11—Zn1 ⁱ | 2.3704 (13) | C7—H7A | 0.9800 |
| P1—C1 | 1.894 (5) | C7—H7B | 0.9800 |
| P1—C5 | 1.891 (5) | C7—H7C | 0.9800 |
| P1—C9 | 1.896 (5) | C8—H8A | 0.9800 |
| C1—C2 | 1.543 (7) | C8—H8B | 0.9800 |
| C1—C3 | 1.524 (7) | C8—H8C | 0.9800 |
| C1—C4 | 1.517 (7) | C9—C10 | 1.528 (6) |
| C2—H2A | 0.9800 | C9—C11 | 1.555 (7) |
| C2—H2B | 0.9800 | C9—C12 | 1.546 (7) |
| C2—H2C | 0.9800 | C10—H10A | 0.9800 |
| C3—H3A | 0.9800 | C10—H10B | 0.9800 |
| C3—H3B | 0.9800 | C10—H10C | 0.9800 |
| C3—H3C | 0.9800 | C11—H11A | 0.9800 |
| C4—H4A | 0.9800 | C11—H11B | 0.9800 |

| | | | |
|--------------------------|-------------|---------------|-----------|
| C4—H4B | 0.9800 | C11—H11C | 0.9800 |
| C4—H4C | 0.9800 | C12—H12A | 0.9800 |
| C5—C6 | 1.536 (7) | C12—H12B | 0.9800 |
| C5—C7 | 1.534 (7) | C12—H12C | 0.9800 |
| | | | |
| C11—Zn1—C11 ⁱ | 90.98 (4) | C5—C6—H6A | 109.5 |
| C11—Zn1—P1 | 112.62 (5) | C5—C6—H6B | 109.5 |
| C11 ⁱ —Zn1—P1 | 113.96 (5) | C5—C6—H6C | 109.5 |
| C12—Zn1—C11 ⁱ | 109.32 (5) | H6A—C6—H6B | 109.5 |
| C12—Zn1—C11 | 109.59 (5) | H6A—C6—H6C | 109.5 |
| C12—Zn1—P1 | 117.30 (5) | H6B—C6—H6C | 109.5 |
| Zn1—C11—Zn1 ⁱ | 89.02 (4) | C5—C7—H7A | 109.5 |
| C1—P1—Zn1 | 110.64 (16) | C5—C7—H7B | 109.5 |
| C1—P1—C9 | 109.9 (2) | C5—C7—H7C | 109.5 |
| C5—P1—Zn1 | 108.67 (16) | H7A—C7—H7B | 109.5 |
| C5—P1—C1 | 110.3 (2) | H7A—C7—H7C | 109.5 |
| C5—P1—C9 | 109.9 (2) | H7B—C7—H7C | 109.5 |
| C9—P1—Zn1 | 107.32 (16) | C5—C8—H8A | 109.5 |
| C2—C1—P1 | 108.6 (3) | C5—C8—H8B | 109.5 |
| C3—C1—P1 | 114.1 (4) | C5—C8—H8C | 109.5 |
| C3—C1—C2 | 109.5 (4) | H8A—C8—H8B | 109.5 |
| C4—C1—P1 | 109.9 (4) | H8A—C8—H8C | 109.5 |
| C4—C1—C2 | 105.8 (4) | H8B—C8—H8C | 109.5 |
| C4—C1—C3 | 108.7 (4) | C10—C9—P1 | 114.6 (4) |
| C1—C2—H2A | 109.5 | C10—C9—C11 | 109.5 (4) |
| C1—C2—H2B | 109.5 | C10—C9—C12 | 108.7 (4) |
| C1—C2—H2C | 109.5 | C11—C9—P1 | 108.5 (3) |
| H2A—C2—H2B | 109.5 | C12—C9—P1 | 110.5 (3) |
| H2A—C2—H2C | 109.5 | C12—C9—C11 | 104.6 (4) |
| H2B—C2—H2C | 109.5 | C9—C10—H10A | 109.5 |
| C1—C3—H3A | 109.5 | C9—C10—H10B | 109.5 |
| C1—C3—H3B | 109.5 | C9—C10—H10C | 109.5 |
| C1—C3—H3C | 109.5 | H10A—C10—H10B | 109.5 |
| H3A—C3—H3B | 109.5 | H10A—C10—H10C | 109.5 |
| H3A—C3—H3C | 109.5 | H10B—C10—H10C | 109.5 |
| H3B—C3—H3C | 109.5 | C9—C11—H11A | 109.5 |
| C1—C4—H4A | 109.5 | C9—C11—H11B | 109.5 |
| C1—C4—H4B | 109.5 | C9—C11—H11C | 109.5 |
| C1—C4—H4C | 109.5 | H11A—C11—H11B | 109.5 |
| H4A—C4—H4B | 109.5 | H11A—C11—H11C | 109.5 |
| H4A—C4—H4C | 109.5 | H11B—C11—H11C | 109.5 |
| H4B—C4—H4C | 109.5 | C9—C12—H12A | 109.5 |
| C6—C5—P1 | 115.0 (3) | C9—C12—H12B | 109.5 |
| C6—C5—C8 | 109.6 (4) | C9—C12—H12C | 109.5 |
| C7—C5—P1 | 111.3 (4) | H12A—C12—H12B | 109.5 |
| C7—C5—C6 | 107.7 (4) | H12A—C12—H12C | 109.5 |
| C7—C5—C8 | 105.1 (4) | H12B—C12—H12C | 109.5 |
| C8—C5—P1 | 107.8 (3) | | |

| | | | |
|---------------|-----------|--------------|-----------|
| Zn1—P1—C1—C2 | 43.7 (4) | C1—P1—C9—C12 | 167.1 (3) |
| Zn1—P1—C1—C3 | 166.1 (4) | C5—P1—C1—C2 | -76.6 (4) |
| Zn1—P1—C1—C4 | -71.7 (4) | C5—P1—C1—C3 | 45.8 (5) |
| Zn1—P1—C5—C6 | 159.6 (3) | C5—P1—C1—C4 | 168.0 (3) |
| Zn1—P1—C5—C7 | -77.7 (4) | C5—P1—C9—C10 | -77.7 (4) |
| Zn1—P1—C5—C8 | 37.0 (4) | C5—P1—C9—C11 | 159.6 (3) |
| Zn1—P1—C9—C10 | 164.3 (3) | C5—P1—C9—C12 | 45.5 (4) |
| Zn1—P1—C9—C11 | 41.6 (4) | C9—P1—C1—C2 | 162.0 (3) |
| Zn1—P1—C9—C12 | -72.6 (3) | C9—P1—C1—C3 | -75.6 (4) |
| C1—P1—C5—C6 | -79.0 (4) | C9—P1—C1—C4 | 46.7 (4) |
| C1—P1—C5—C7 | 43.8 (4) | C9—P1—C5—C6 | 42.4 (4) |
| C1—P1—C5—C8 | 158.5 (3) | C9—P1—C5—C7 | 165.1 (3) |
| C1—P1—C9—C10 | 43.9 (4) | C9—P1—C5—C8 | -80.1 (4) |
| C1—P1—C9—C11 | -78.8 (4) | | |

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

Hydrogen-bond geometry ($\text{\AA}, ^\circ$)

| $D-H\cdots A$ | $D-H$ | $H\cdots A$ | $D\cdots A$ | $D-H\cdots A$ |
|-------------------------|-------|-------------|-------------|---------------|
| $C3-H3C\cdots C11^{ii}$ | 0.98 | 2.88 | 3.479 (6) | 120 |

Symmetry code: (ii) $-x-1/2, y-1/2, z$.

(2) Dichlorido(tetrahydrofuran- κO)(tri-*tert*-butylphosphane- κP)zinc

Crystal data

$[\text{ZnCl}_2(\text{C}_4\text{H}_8\text{O})(\text{C}_{12}\text{H}_{27}\text{P})]$

$M_r = 410.68$

Orthorhombic, $Pna2_1$

$a = 26.4580$ (11) \AA

$b = 8.9281$ (4) \AA

$c = 8.5790$ (4) \AA

$V = 2026.53$ (16) \AA^3

$Z = 4$

$F(000) = 872$

$D_x = 1.346$ Mg m^{-3}

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

Cell parameters from 9968 reflections

$\theta = 2.8\text{--}27.6^\circ$

$\mu = 1.55$ mm^{-1}

$T = 193$ K

Block, colourless

$0.47 \times 0.46 \times 0.27$ mm

Data collection

Bruker APEXII CCD area-detector
diffractometer

Radiation source: sealed tube

Graphite monochromator

Detector resolution: 8.33 pixels mm^{-1}

ω and φ scans

Absorption correction: integration

(*SADABS*; Bruker, 2008)

$T_{\min} = 0.580, T_{\max} = 0.754$

23933 measured reflections

3698 independent reflections

3581 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.038$

$\theta_{\max} = 25.4^\circ, \theta_{\min} = 2.4^\circ$

$h = -31 \rightarrow 31$

$k = -10 \rightarrow 10$

$l = -10 \rightarrow 10$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.031$

$wR(F^2) = 0.075$

$S = 1.14$

3698 reflections

200 parameters
 1 restraint
 Hydrogen site location: inferred from
 neighbouring sites
 H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0318P)^2 + 1.5653P]$
 where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.69 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.30 \text{ e } \text{\AA}^{-3}$
 Absolute structure: Refined as an inversion
 twin.
 Absolute structure parameter: 0.206 (18)

Special details

Experimental. One distinct cell was identified using *APEX2* (Bruker, 2004). Six frame series were integrated and filtered for statistical outliers using *SAINT* (Bruker, 2005) then corrected for absorption by integration using *SHELXTL/XPREF* V2005/2 (Bruker, 2005) before using *SAINT/SADABS* (Bruker, 2005) to sort, merge, and scale the combined data. No decay correction was applied.

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. Refined as a 2-component inversion twin.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

| | <i>x</i> | <i>y</i> | <i>z</i> | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|-----|--------------|--------------|--------------|----------------------------------|
| Zn1 | 0.88274 (2) | 0.23668 (4) | 0.83834 (8) | 0.02419 (14) |
| Cl1 | 0.95342 (5) | 0.17357 (16) | 0.96546 (16) | 0.0477 (4) |
| Cl2 | 0.81608 (5) | 0.08915 (12) | 0.88009 (17) | 0.0494 (4) |
| P1 | 0.86195 (3) | 0.49987 (10) | 0.85327 (14) | 0.0168 (2) |
| O1 | 0.90182 (12) | 0.1814 (4) | 0.6087 (4) | 0.0266 (7) |
| C1 | 0.91147 (16) | 0.6144 (5) | 0.7468 (5) | 0.0257 (9) |
| C2 | 0.96425 (16) | 0.5523 (5) | 0.7851 (6) | 0.0342 (11) |
| H2A | 0.9897 | 0.6048 | 0.7224 | 0.051* |
| H2B | 0.9653 | 0.4450 | 0.7612 | 0.051* |
| H2C | 0.9714 | 0.5675 | 0.8960 | 0.051* |
| C3 | 0.9103 (2) | 0.7834 (5) | 0.7839 (6) | 0.0357 (12) |
| H3A | 0.9352 | 0.8355 | 0.7190 | 0.054* |
| H3B | 0.9185 | 0.7990 | 0.8941 | 0.054* |
| H3C | 0.8765 | 0.8232 | 0.7621 | 0.054* |
| C4 | 0.9053 (2) | 0.5948 (6) | 0.5701 (6) | 0.0369 (12) |
| H4A | 0.9325 | 0.6482 | 0.5161 | 0.055* |
| H4B | 0.8726 | 0.6358 | 0.5376 | 0.055* |
| H4C | 0.9068 | 0.4881 | 0.5439 | 0.055* |
| C5 | 0.86075 (17) | 0.5543 (5) | 1.0665 (5) | 0.0256 (9) |
| C6 | 0.9152 (2) | 0.5661 (7) | 1.1274 (6) | 0.0414 (15) |
| H6A | 0.9147 | 0.5778 | 1.2410 | 0.062* |
| H6B | 0.9317 | 0.6531 | 1.0801 | 0.062* |
| H6C | 0.9338 | 0.4750 | 1.1000 | 0.062* |
| C7 | 0.8333 (2) | 0.7020 (6) | 1.1031 (7) | 0.0370 (11) |
| H7A | 0.8381 | 0.7274 | 1.2132 | 0.055* |
| H7B | 0.7971 | 0.6905 | 1.0815 | 0.055* |
| H7C | 0.8471 | 0.7823 | 1.0378 | 0.055* |
| C8 | 0.8363 (3) | 0.4261 (6) | 1.1571 (6) | 0.0464 (14) |

| | | | | |
|------|--------------|------------|------------|-------------|
| H8A | 0.8371 | 0.4488 | 1.2689 | 0.070* |
| H8B | 0.8550 | 0.3332 | 1.1374 | 0.070* |
| H8C | 0.8012 | 0.4140 | 1.1233 | 0.070* |
| C9 | 0.79715 (16) | 0.5310 (5) | 0.7633 (6) | 0.0292 (10) |
| C10 | 0.7853 (2) | 0.6937 (6) | 0.7234 (8) | 0.0479 (14) |
| H10A | 0.7507 | 0.7011 | 0.6841 | 0.072* |
| H10B | 0.8089 | 0.7289 | 0.6432 | 0.072* |
| H10C | 0.7889 | 0.7556 | 0.8170 | 0.072* |
| C11 | 0.75568 (15) | 0.4718 (6) | 0.8714 (7) | 0.0433 (14) |
| H11A | 0.7229 | 0.4780 | 0.8184 | 0.065* |
| H11B | 0.7547 | 0.5323 | 0.9667 | 0.065* |
| H11C | 0.7628 | 0.3672 | 0.8983 | 0.065* |
| C12 | 0.79414 (19) | 0.4325 (6) | 0.6168 (6) | 0.0356 (12) |
| H12A | 0.7611 | 0.4456 | 0.5672 | 0.053* |
| H12B | 0.7986 | 0.3272 | 0.6460 | 0.053* |
| H12C | 0.8208 | 0.4619 | 0.5437 | 0.053* |
| C13 | 0.86717 (17) | 0.1097 (6) | 0.4985 (6) | 0.0344 (11) |
| H13A | 0.8461 | 0.1856 | 0.4448 | 0.041* |
| H13B | 0.8446 | 0.0384 | 0.5530 | 0.041* |
| C14 | 0.90056 (19) | 0.0286 (6) | 0.3834 (6) | 0.0390 (12) |
| H14A | 0.8853 | 0.0281 | 0.2780 | 0.047* |
| H14B | 0.9069 | -0.0759 | 0.4167 | 0.047* |
| C15 | 0.94860 (18) | 0.1199 (6) | 0.3865 (6) | 0.0360 (11) |
| H15A | 0.9779 | 0.0601 | 0.3505 | 0.043* |
| H15B | 0.9455 | 0.2107 | 0.3209 | 0.043* |
| C16 | 0.95347 (17) | 0.1599 (6) | 0.5564 (6) | 0.0299 (10) |
| H16A | 0.9700 | 0.0781 | 0.6153 | 0.036* |
| H16B | 0.9734 | 0.2528 | 0.5697 | 0.036* |

Atomic displacement parameters (\AA^2)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|-----|-------------|-------------|-------------|--------------|--------------|--------------|
| Zn1 | 0.0309 (2) | 0.0208 (2) | 0.0208 (2) | 0.00455 (17) | 0.0058 (3) | 0.0024 (3) |
| Cl1 | 0.0617 (8) | 0.0524 (7) | 0.0288 (6) | 0.0304 (6) | -0.0150 (6) | -0.0030 (6) |
| Cl2 | 0.0564 (7) | 0.0252 (5) | 0.0666 (11) | -0.0062 (5) | 0.0398 (7) | -0.0007 (6) |
| P1 | 0.0177 (4) | 0.0191 (4) | 0.0138 (5) | 0.0009 (3) | -0.0005 (5) | 0.0008 (5) |
| O1 | 0.0250 (15) | 0.0374 (17) | 0.0175 (15) | -0.0053 (13) | 0.0032 (13) | -0.0063 (13) |
| C1 | 0.028 (2) | 0.027 (2) | 0.022 (2) | -0.0059 (18) | 0.0090 (18) | -0.0005 (18) |
| C2 | 0.025 (2) | 0.039 (3) | 0.039 (3) | -0.0069 (19) | 0.0081 (19) | -0.009 (2) |
| C3 | 0.046 (3) | 0.027 (2) | 0.035 (3) | -0.007 (2) | 0.007 (2) | -0.0012 (19) |
| C4 | 0.055 (3) | 0.033 (3) | 0.023 (3) | -0.006 (2) | 0.011 (2) | 0.004 (2) |
| C5 | 0.032 (2) | 0.030 (2) | 0.014 (2) | 0.0020 (19) | 0.0022 (17) | -0.0040 (18) |
| C6 | 0.044 (3) | 0.061 (4) | 0.019 (3) | 0.013 (3) | -0.012 (2) | -0.008 (2) |
| C7 | 0.042 (3) | 0.035 (3) | 0.034 (3) | 0.005 (2) | 0.004 (2) | -0.013 (2) |
| C8 | 0.076 (4) | 0.043 (3) | 0.020 (2) | 0.005 (3) | 0.015 (3) | 0.003 (2) |
| C9 | 0.018 (2) | 0.031 (2) | 0.039 (3) | 0.0066 (17) | -0.0133 (19) | -0.008 (2) |
| C10 | 0.049 (3) | 0.038 (3) | 0.056 (4) | 0.016 (2) | -0.028 (3) | -0.003 (3) |
| C11 | 0.020 (2) | 0.046 (3) | 0.063 (4) | 0.0013 (18) | 0.002 (2) | -0.021 (3) |

| | | | | | | |
|-----|-----------|-----------|-----------|--------------|--------------|------------|
| C12 | 0.032 (2) | 0.040 (3) | 0.036 (3) | 0.003 (2) | -0.018 (2) | -0.007 (2) |
| C13 | 0.029 (2) | 0.047 (3) | 0.028 (3) | -0.008 (2) | -0.0043 (19) | -0.009 (2) |
| C14 | 0.049 (3) | 0.039 (3) | 0.029 (3) | -0.002 (2) | -0.001 (2) | -0.011 (2) |
| C15 | 0.037 (2) | 0.046 (3) | 0.025 (3) | 0.004 (2) | 0.0101 (19) | -0.004 (2) |
| C16 | 0.023 (2) | 0.037 (2) | 0.029 (3) | -0.0011 (19) | 0.0047 (18) | -0.004 (2) |

Geometric parameters (Å, °)

| | | | |
|-------------|-------------|------------|-----------|
| Zn1—C11 | 2.2370 (13) | C7—H7B | 0.9800 |
| Zn1—C12 | 2.2301 (13) | C7—H7C | 0.9800 |
| Zn1—P1 | 2.4167 (9) | C8—H8A | 0.9800 |
| Zn1—O1 | 2.093 (3) | C8—H8B | 0.9800 |
| P1—C1 | 1.896 (4) | C8—H8C | 0.9800 |
| P1—C5 | 1.893 (5) | C9—C10 | 1.525 (7) |
| P1—C9 | 1.900 (4) | C9—C11 | 1.531 (7) |
| O1—C13 | 1.464 (5) | C9—C12 | 1.536 (7) |
| O1—C16 | 1.451 (5) | C10—H10A | 0.9800 |
| C1—C2 | 1.538 (6) | C10—H10B | 0.9800 |
| C1—C3 | 1.543 (6) | C10—H10C | 0.9800 |
| C1—C4 | 1.535 (7) | C11—H11A | 0.9800 |
| C2—H2A | 0.9800 | C11—H11B | 0.9800 |
| C2—H2B | 0.9800 | C11—H11C | 0.9800 |
| C2—H2C | 0.9800 | C12—H12A | 0.9800 |
| C3—H3A | 0.9800 | C12—H12B | 0.9800 |
| C3—H3B | 0.9800 | C12—H12C | 0.9800 |
| C3—H3C | 0.9800 | C13—H13A | 0.9900 |
| C4—H4A | 0.9800 | C13—H13B | 0.9900 |
| C4—H4B | 0.9800 | C13—C14 | 1.510 (7) |
| C4—H4C | 0.9800 | C14—H14A | 0.9900 |
| C5—C6 | 1.535 (7) | C14—H14B | 0.9900 |
| C5—C7 | 1.538 (6) | C14—C15 | 1.510 (7) |
| C5—C8 | 1.528 (7) | C15—H15A | 0.9900 |
| C6—H6A | 0.9800 | C15—H15B | 0.9900 |
| C6—H6B | 0.9800 | C15—C16 | 1.507 (6) |
| C6—H6C | 0.9800 | C16—H16A | 0.9900 |
| C7—H7A | 0.9800 | C16—H16B | 0.9900 |
| C11—Zn1—P1 | 114.16 (5) | H7B—C7—H7C | 109.5 |
| C12—Zn1—C11 | 115.72 (6) | C5—C8—H8A | 109.5 |
| C12—Zn1—P1 | 112.69 (4) | C5—C8—H8B | 109.5 |
| O1—Zn1—C11 | 101.41 (10) | C5—C8—H8C | 109.5 |
| O1—Zn1—C12 | 101.70 (9) | H8A—C8—H8B | 109.5 |
| O1—Zn1—P1 | 109.51 (10) | H8A—C8—H8C | 109.5 |
| C1—P1—Zn1 | 109.97 (14) | H8B—C8—H8C | 109.5 |
| C1—P1—C9 | 110.4 (2) | C10—C9—P1 | 114.5 (3) |
| C5—P1—Zn1 | 107.75 (15) | C10—C9—C11 | 108.6 (4) |
| C5—P1—C1 | 109.8 (2) | C10—C9—C12 | 110.5 (4) |
| C5—P1—C9 | 109.9 (2) | C11—C9—P1 | 110.5 (3) |

| | | | |
|--------------|-------------|---------------|------------|
| C9—P1—Zn1 | 109.02 (14) | C11—C9—C12 | 105.1 (4) |
| C13—O1—Zn1 | 124.0 (3) | C12—C9—P1 | 107.2 (3) |
| C16—O1—Zn1 | 123.3 (3) | C9—C10—H10A | 109.5 |
| C16—O1—C13 | 109.4 (3) | C9—C10—H10B | 109.5 |
| C2—C1—P1 | 109.3 (3) | C9—C10—H10C | 109.5 |
| C2—C1—C3 | 109.1 (4) | H10A—C10—H10B | 109.5 |
| C3—C1—P1 | 114.5 (3) | H10A—C10—H10C | 109.5 |
| C4—C1—P1 | 109.9 (3) | H10B—C10—H10C | 109.5 |
| C4—C1—C2 | 105.5 (4) | C9—C11—H11A | 109.5 |
| C4—C1—C3 | 108.2 (4) | C9—C11—H11B | 109.5 |
| C1—C2—H2A | 109.5 | C9—C11—H11C | 109.5 |
| C1—C2—H2B | 109.5 | H11A—C11—H11B | 109.5 |
| C1—C2—H2C | 109.5 | H11A—C11—H11C | 109.5 |
| H2A—C2—H2B | 109.5 | H11B—C11—H11C | 109.5 |
| H2A—C2—H2C | 109.5 | C9—C12—H12A | 109.5 |
| H2B—C2—H2C | 109.5 | C9—C12—H12B | 109.5 |
| C1—C3—H3A | 109.5 | C9—C12—H12C | 109.5 |
| C1—C3—H3B | 109.5 | H12A—C12—H12B | 109.5 |
| C1—C3—H3C | 109.5 | H12A—C12—H12C | 109.5 |
| H3A—C3—H3B | 109.5 | H12B—C12—H12C | 109.5 |
| H3A—C3—H3C | 109.5 | O1—C13—H13A | 110.7 |
| H3B—C3—H3C | 109.5 | O1—C13—H13B | 110.7 |
| C1—C4—H4A | 109.5 | O1—C13—C14 | 105.4 (4) |
| C1—C4—H4B | 109.5 | H13A—C13—H13B | 108.8 |
| C1—C4—H4C | 109.5 | C14—C13—H13A | 110.7 |
| H4A—C4—H4B | 109.5 | C14—C13—H13B | 110.7 |
| H4A—C4—H4C | 109.5 | C13—C14—H14A | 111.2 |
| H4B—C4—H4C | 109.5 | C13—C14—H14B | 111.2 |
| C6—C5—P1 | 109.3 (3) | C13—C14—C15 | 102.8 (4) |
| C6—C5—C7 | 108.4 (4) | H14A—C14—H14B | 109.1 |
| C7—C5—P1 | 115.2 (3) | C15—C14—H14A | 111.2 |
| C8—C5—P1 | 107.8 (3) | C15—C14—H14B | 111.2 |
| C8—C5—C6 | 106.0 (4) | C14—C15—H15A | 111.3 |
| C8—C5—C7 | 109.8 (4) | C14—C15—H15B | 111.3 |
| C5—C6—H6A | 109.5 | H15A—C15—H15B | 109.2 |
| C5—C6—H6B | 109.5 | C16—C15—C14 | 102.5 (4) |
| C5—C6—H6C | 109.5 | C16—C15—H15A | 111.3 |
| H6A—C6—H6B | 109.5 | C16—C15—H15B | 111.3 |
| H6A—C6—H6C | 109.5 | O1—C16—C15 | 104.5 (4) |
| H6B—C6—H6C | 109.5 | O1—C16—H16A | 110.9 |
| C5—C7—H7A | 109.5 | O1—C16—H16B | 110.9 |
| C5—C7—H7B | 109.5 | C15—C16—H16A | 110.9 |
| C5—C7—H7C | 109.5 | C15—C16—H16B | 110.9 |
| H7A—C7—H7B | 109.5 | H16A—C16—H16B | 108.9 |
| H7A—C7—H7C | 109.5 | | |
| Zn1—P1—C1—C2 | -42.1 (3) | C5—P1—C1—C3 | -46.3 (4) |
| Zn1—P1—C1—C3 | -164.7 (3) | C5—P1—C1—C4 | -168.4 (3) |

| | | | |
|----------------|------------|-----------------|------------|
| Zn1—P1—C1—C4 | 73.2 (3) | C9—P1—C1—C2 | -162.4 (3) |
| Zn1—P1—C5—C6 | 75.2 (4) | C9—P1—C1—C3 | 74.9 (4) |
| Zn1—P1—C5—C7 | -162.5 (3) | C9—P1—C1—C4 | -47.1 (4) |
| Zn1—P1—C5—C8 | -39.6 (4) | C9—P1—C5—C6 | -166.2 (3) |
| Zn1—O1—C13—C14 | -154.5 (3) | C9—P1—C5—C7 | -43.9 (4) |
| Zn1—O1—C16—C15 | 179.1 (3) | C9—P1—C5—C8 | 79.1 (4) |
| O1—C13—C14—C15 | -27.6 (5) | C13—O1—C16—C15 | 18.7 (5) |
| C1—P1—C5—C6 | -44.6 (4) | C13—C14—C15—C16 | 38.5 (5) |
| C1—P1—C5—C7 | 77.7 (4) | C14—C15—C16—O1 | -35.4 (5) |
| C1—P1—C5—C8 | -159.3 (3) | C16—O1—C13—C14 | 5.7 (5) |
| C5—P1—C1—C2 | 76.3 (3) | | |

Hydrogen-bond geometry (Å, °)

| <i>D</i> —H... <i>A</i> | <i>D</i> —H | H... <i>A</i> | <i>D</i> ... <i>A</i> | <i>D</i> —H... <i>A</i> |
|---------------------------------------|-------------|---------------|-----------------------|-------------------------|
| C12—H12 <i>A</i> ...C12 ⁱ | 0.98 | 2.90 | 3.819 (5) | 157 |
| C15—H15 <i>A</i> ...C11 ⁱⁱ | 0.99 | 2.94 | 3.747 (5) | 140 |

Symmetry codes: (i) $-x+3/2, y+1/2, z-1/2$; (ii) $-x+2, -y, z-1/2$.(3) Tri-*tert*-butylphosphonium aquatrichloridozincate 1,2-dichloroethane monosolvate

Crystal data

(C₁₂H₂₈P)[ZnCl₃(H₂O)]·C₂H₄Cl₂*M_r* = 492.00Monoclinic, *P*2₁/*n**a* = 25.3722 (5) Å*b* = 8.5841 (2) Å*c* = 32.912 (2) Å β = 98.909 (7)°*V* = 7081.7 (6) Å³*Z* = 12*F*(000) = 3072*D_x* = 1.384 Mg m⁻³Cu *K* α radiation, λ = 1.54187 Å

Cell parameters from 2316 reflections

 θ = 23.1–67.3° μ = 7.28 mm⁻¹*T* = 123 K

Needle, colourless

0.18 × 0.05 × 0.02 mm

Data collection

Rigaku CCD area-detector
diffractometer

Radiation source: sealed tube

Focusing graphite monochromator

Detector resolution: 22.2222 pixels mm⁻¹ ω and ϕ scans

Absorption correction: multi-scan

(CrystalClear; Rigaku, 2010)

T_{min} = 0.354, *T_{max}* = 0.868

67698 measured reflections

11604 independent reflections

7131 reflections with *I* > 2 σ (*I*)*R_{int}* = 0.083 θ_{\max} = 63.7°, θ_{\min} = 6.5°*h* = -29→29*k* = -9→9*l* = -37→38

Refinement

Refinement on *F*²

Least-squares matrix: full

R[*F*² > 2 σ (*F*²)] = 0.049*wR*(*F*²) = 0.142*S* = 0.98

11604 reflections

760 parameters

462 restraints

Hydrogen site location: mixed

H atoms treated by a mixture of independent
and constrained refinement*w* = 1/[$\sigma^2(F_o^2) + (0.0718P)^2$]where *P* = (*F_o*² + 2*F_c*²)/3(Δ/σ)_{max} = 0.002 $\Delta\rho_{\max}$ = 0.51 e Å⁻³ $\Delta\rho_{\min}$ = -0.70 e Å⁻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.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

| | <i>x</i> | <i>y</i> | <i>z</i> | $U_{\text{iso}}^*/U_{\text{eq}}$ | Occ. (<1) |
|------|--------------|---------------|--------------|----------------------------------|-----------|
| Zn1 | 0.29184 (2) | -0.04314 (6) | 0.78814 (2) | 0.03885 (16) | |
| Cl1 | 0.25307 (4) | 0.14800 (11) | 0.82087 (3) | 0.0522 (3) | |
| Cl2 | 0.28227 (4) | -0.27034 (11) | 0.82132 (3) | 0.0457 (3) | |
| Cl3 | 0.37414 (4) | 0.00389 (13) | 0.77610 (3) | 0.0548 (3) | |
| O1 | 0.24204 (11) | -0.0627 (3) | 0.73407 (8) | 0.0445 (7) | |
| H1OA | 0.2483 (16) | 0.017 (3) | 0.7219 (11) | 0.053* | |
| H1OB | 0.2458 (16) | -0.137 (3) | 0.7191 (10) | 0.053* | |
| Zn2 | 0.71198 (2) | -0.04606 (6) | 0.87883 (2) | 0.03871 (16) | |
| Cl4 | 0.63093 (4) | 0.01180 (13) | 0.89180 (3) | 0.0547 (3) | |
| Cl5 | 0.71685 (4) | -0.27780 (11) | 0.84662 (3) | 0.0485 (3) | |
| Cl6 | 0.75106 (4) | 0.13977 (11) | 0.84448 (3) | 0.0516 (3) | |
| O2 | 0.76256 (11) | -0.0670 (3) | 0.93254 (8) | 0.0437 (7) | |
| H2OA | 0.7606 (16) | 0.016 (3) | 0.9446 (11) | 0.052* | |
| H2OB | 0.7594 (15) | -0.146 (3) | 0.9460 (10) | 0.052* | |
| Zn3 | 0.71002 (2) | -0.04753 (6) | 0.54563 (2) | 0.03781 (16) | |
| Cl7 | 0.62855 (4) | 0.00691 (12) | 0.55823 (3) | 0.0513 (3) | |
| Cl8 | 0.75010 (4) | 0.14341 (11) | 0.51378 (3) | 0.0474 (3) | |
| Cl9 | 0.71658 (4) | -0.27393 (11) | 0.51121 (3) | 0.0453 (3) | |
| O3 | 0.75974 (11) | -0.0739 (3) | 0.59969 (8) | 0.0426 (7) | |
| H3OA | 0.7567 (16) | 0.010 (3) | 0.6128 (11) | 0.051* | |
| H3OB | 0.7559 (15) | -0.155 (3) | 0.6128 (10) | 0.051* | |
| Cl1B | 0.0853 (6) | -0.0049 (19) | 0.8119 (4) | 0.059 (2) | 0.52 (3) |
| Cl2B | 0.0469 (4) | 0.0155 (16) | 0.7101 (4) | 0.0494 (17) | 0.52 (3) |
| C37B | 0.1183 (7) | -0.091 (2) | 0.7729 (4) | 0.057 (3) | 0.52 (3) |
| H37A | 0.1563 | -0.1081 | 0.7842 | 0.068* | 0.52 (3) |
| H37B | 0.1019 | -0.1930 | 0.7649 | 0.068* | 0.52 (3) |
| C38B | 0.1147 (5) | 0.011 (2) | 0.7355 (4) | 0.054 (3) | 0.52 (3) |
| H38A | 0.1380 | -0.0311 | 0.7166 | 0.065* | 0.52 (3) |
| H38B | 0.1268 | 0.1174 | 0.7436 | 0.065* | 0.52 (3) |
| Cl1A | 0.0777 (6) | -0.033 (2) | 0.8115 (4) | 0.057 (2) | 0.48 (3) |
| Cl2A | 0.0519 (5) | 0.058 (2) | 0.7128 (5) | 0.058 (2) | 0.48 (3) |
| C37A | 0.1256 (7) | -0.057 (2) | 0.7772 (5) | 0.058 (3) | 0.48 (3) |
| H37C | 0.1622 | -0.0472 | 0.7925 | 0.070* | 0.48 (3) |
| H37D | 0.1220 | -0.1624 | 0.7645 | 0.070* | 0.48 (3) |
| C38A | 0.1159 (5) | 0.065 (2) | 0.7445 (5) | 0.056 (3) | 0.48 (3) |
| H38C | 0.1437 | 0.0548 | 0.7265 | 0.067* | 0.48 (3) |
| H38D | 0.1203 | 0.1682 | 0.7578 | 0.067* | 0.48 (3) |
| Cl3B | 0.95522 (14) | 0.0663 (5) | 0.95863 (12) | 0.0614 (8) | 0.881 (7) |
| Cl4B | 0.9273 (2) | -0.0185 (7) | 0.85820 (13) | 0.0625 (11) | 0.881 (7) |

| | | | | | |
|------|--------------|--------------|--------------|-------------|-----------|
| C39B | 0.89081 (19) | 0.0740 (7) | 0.92710 (18) | 0.0607 (16) | 0.881 (7) |
| H39A | 0.8629 | 0.0644 | 0.9449 | 0.073* | 0.881 (7) |
| H39B | 0.8864 | 0.1767 | 0.9133 | 0.073* | 0.881 (7) |
| C40B | 0.88298 (18) | -0.0527 (7) | 0.89497 (14) | 0.0564 (15) | 0.881 (7) |
| H40A | 0.8456 | -0.0526 | 0.8809 | 0.068* | 0.881 (7) |
| H40B | 0.8906 | -0.1556 | 0.9082 | 0.068* | 0.881 (7) |
| Cl4A | 0.9330 (16) | -0.044 (5) | 0.8654 (10) | 0.051 (4) | 0.119 (7) |
| Cl3A | 0.9579 (9) | 0.017 (3) | 0.9553 (7) | 0.035 (4) | 0.119 (7) |
| C39A | 0.8883 (10) | 0.007 (6) | 0.9349 (10) | 0.055 (3) | 0.119 (7) |
| H39C | 0.8746 | -0.1005 | 0.9353 | 0.066* | 0.119 (7) |
| H39D | 0.8669 | 0.0762 | 0.9502 | 0.066* | 0.119 (7) |
| C40A | 0.8882 (14) | 0.064 (5) | 0.8920 (9) | 0.057 (3) | 0.119 (7) |
| H40C | 0.8985 | 0.1751 | 0.8928 | 0.068* | 0.119 (7) |
| H40D | 0.8516 | 0.0555 | 0.8766 | 0.068* | 0.119 (7) |
| Cl5A | 0.9498 (6) | 0.027 (2) | 0.6225 (4) | 0.052 (2) | 0.38 (3) |
| Cl6A | 0.9263 (6) | -0.038 (2) | 0.5236 (4) | 0.060 (3) | 0.38 (3) |
| C41A | 0.8869 (6) | 0.053 (3) | 0.5893 (6) | 0.057 (3) | 0.38 (3) |
| H41A | 0.8578 | 0.0477 | 0.6061 | 0.069* | 0.38 (3) |
| H41B | 0.8862 | 0.1578 | 0.5768 | 0.069* | 0.38 (3) |
| C42A | 0.8763 (10) | -0.064 (3) | 0.5558 (6) | 0.060 (3) | 0.38 (3) |
| H42A | 0.8403 | -0.0486 | 0.5398 | 0.072* | 0.38 (3) |
| H42B | 0.8782 | -0.1711 | 0.5673 | 0.072* | 0.38 (3) |
| Cl5B | 0.9580 (3) | -0.0244 (15) | 0.6235 (3) | 0.0596 (16) | 0.62 (3) |
| Cl6B | 0.9182 (4) | -0.0052 (12) | 0.5223 (3) | 0.0643 (18) | 0.62 (3) |
| C41B | 0.8897 (3) | -0.0012 (19) | 0.5993 (4) | 0.058 (2) | 0.62 (3) |
| H41C | 0.8654 | -0.0432 | 0.6174 | 0.069* | 0.62 (3) |
| H41D | 0.8815 | 0.1108 | 0.5946 | 0.069* | 0.62 (3) |
| C42B | 0.8817 (6) | -0.0858 (19) | 0.5595 (4) | 0.062 (3) | 0.62 (3) |
| H42C | 0.8432 | -0.0850 | 0.5481 | 0.075* | 0.62 (3) |
| H42D | 0.8926 | -0.1957 | 0.5646 | 0.075* | 0.62 (3) |
| P1 | 0.38352 (4) | 0.01636 (11) | 0.90951 (3) | 0.0318 (2) | |
| H1P | 0.3525 (12) | -0.005 (3) | 0.8743 (8) | 0.038* | |
| C1 | 0.39955 (14) | 0.2288 (4) | 0.91048 (11) | 0.0388 (9) | |
| C2 | 0.44827 (15) | 0.2657 (5) | 0.94268 (12) | 0.0529 (11) | |
| H2A | 0.4796 | 0.2118 | 0.9355 | 0.063* | |
| H2B | 0.4417 | 0.2308 | 0.9698 | 0.063* | |
| H2C | 0.4547 | 0.3784 | 0.9433 | 0.063* | |
| C3 | 0.35169 (14) | 0.3250 (4) | 0.92003 (12) | 0.0477 (10) | |
| H3A | 0.3449 | 0.3006 | 0.9479 | 0.057* | |
| H3B | 0.3200 | 0.2994 | 0.9001 | 0.057* | |
| H3C | 0.3597 | 0.4363 | 0.9182 | 0.057* | |
| C4 | 0.40893 (16) | 0.2786 (5) | 0.86721 (11) | 0.0533 (11) | |
| H4A | 0.4111 | 0.3924 | 0.8660 | 0.064* | |
| H4B | 0.3793 | 0.2419 | 0.8468 | 0.064* | |
| H4C | 0.4424 | 0.2331 | 0.8614 | 0.064* | |
| C5 | 0.34024 (14) | -0.0466 (4) | 0.94810 (10) | 0.0366 (9) | |
| C6 | 0.36314 (16) | 0.0051 (5) | 0.99169 (12) | 0.0508 (11) | |
| H6A | 0.3400 | -0.0318 | 1.0109 | 0.061* | |

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|------|--------------|--------------|--------------|-------------|
| H6B | 0.3652 | 0.1191 | 0.9927 | 0.061* |
| H6C | 0.3990 | -0.0388 | 0.9994 | 0.061* |
| C7 | 0.33375 (14) | -0.2249 (4) | 0.94671 (11) | 0.0464 (10) |
| H7A | 0.3685 | -0.2742 | 0.9556 | 0.056* |
| H7B | 0.3198 | -0.2575 | 0.9186 | 0.056* |
| H7C | 0.3088 | -0.2565 | 0.9651 | 0.056* |
| C8 | 0.28426 (14) | 0.0216 (4) | 0.93549 (12) | 0.0439 (10) |
| H8A | 0.2597 | -0.0254 | 0.9522 | 0.053* |
| H8B | 0.2719 | -0.0008 | 0.9064 | 0.053* |
| H8C | 0.2854 | 0.1346 | 0.9398 | 0.053* |
| C9 | 0.44243 (13) | -0.1135 (4) | 0.90654 (11) | 0.0362 (9) |
| C10 | 0.47453 (15) | -0.1455 (5) | 0.94889 (11) | 0.0546 (12) |
| H10A | 0.5052 | -0.2119 | 0.9459 | 0.065* |
| H10B | 0.4518 | -0.1985 | 0.9661 | 0.065* |
| H10C | 0.4872 | -0.0468 | 0.9618 | 0.065* |
| C11 | 0.42058 (15) | -0.2665 (4) | 0.88554 (12) | 0.0520 (11) |
| H11A | 0.4504 | -0.3327 | 0.8806 | 0.062* |
| H11B | 0.3982 | -0.2424 | 0.8593 | 0.062* |
| H11C | 0.3993 | -0.3215 | 0.9034 | 0.062* |
| C12 | 0.47862 (14) | -0.0434 (5) | 0.87794 (11) | 0.0480 (11) |
| H12A | 0.4948 | 0.0528 | 0.8901 | 0.058* |
| H12B | 0.4574 | -0.0202 | 0.8511 | 0.058* |
| H12C | 0.5067 | -0.1181 | 0.8743 | 0.058* |
| P2 | 0.61653 (4) | 0.00937 (11) | 0.75750 (3) | 0.0357 (3) |
| H2P | 0.6500 (12) | -0.002 (4) | 0.7919 (9) | 0.043* |
| C13 | 0.56962 (15) | 0.1673 (5) | 0.76841 (12) | 0.0480 (11) |
| C14 | 0.52888 (15) | 0.1050 (5) | 0.79467 (12) | 0.0599 (12) |
| H14A | 0.5067 | 0.1911 | 0.8019 | 0.072* |
| H14B | 0.5479 | 0.0576 | 0.8198 | 0.072* |
| H14C | 0.5061 | 0.0266 | 0.7790 | 0.072* |
| C15 | 0.53920 (16) | 0.2341 (5) | 0.72847 (12) | 0.0589 (12) |
| H15A | 0.5169 | 0.1526 | 0.7137 | 0.071* |
| H15B | 0.5647 | 0.2723 | 0.7112 | 0.071* |
| H15C | 0.5165 | 0.3204 | 0.7349 | 0.071* |
| C16 | 0.60288 (16) | 0.2926 (4) | 0.79426 (12) | 0.0555 (12) |
| H16A | 0.5790 | 0.3716 | 0.8029 | 0.067* |
| H16B | 0.6272 | 0.3418 | 0.7777 | 0.067* |
| H16C | 0.6235 | 0.2440 | 0.8186 | 0.067* |
| C17 | 0.58579 (17) | -0.1869 (5) | 0.74906 (12) | 0.0550 (11) |
| C18 | 0.57213 (17) | -0.2461 (5) | 0.79071 (12) | 0.0656 (13) |
| H18A | 0.5615 | -0.3559 | 0.7881 | 0.079* |
| H18B | 0.5427 | -0.1844 | 0.7984 | 0.079* |
| H18C | 0.6035 | -0.2355 | 0.8120 | 0.079* |
| C19 | 0.53501 (17) | -0.1839 (5) | 0.71654 (13) | 0.0711 (14) |
| H19A | 0.5440 | -0.1466 | 0.6903 | 0.085* |
| H19B | 0.5087 | -0.1139 | 0.7257 | 0.085* |
| H19C | 0.5201 | -0.2892 | 0.7130 | 0.085* |
| C20 | 0.62687 (18) | -0.3035 (4) | 0.73614 (12) | 0.0641 (13) |

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|------|--------------|--------------|--------------|-------------|
| H20A | 0.6590 | -0.3043 | 0.7570 | 0.077* |
| H20B | 0.6364 | -0.2717 | 0.7096 | 0.077* |
| H20C | 0.6112 | -0.4081 | 0.7337 | 0.077* |
| C21 | 0.66109 (15) | 0.0665 (4) | 0.71933 (11) | 0.0443 (10) |
| C22 | 0.71302 (15) | -0.0295 (5) | 0.72902 (12) | 0.0494 (11) |
| H22A | 0.7049 | -0.1403 | 0.7243 | 0.059* |
| H22B | 0.7289 | -0.0135 | 0.7578 | 0.059* |
| H22C | 0.7383 | 0.0042 | 0.7111 | 0.059* |
| C23 | 0.67727 (15) | 0.2390 (4) | 0.72459 (11) | 0.0487 (11) |
| H23A | 0.7020 | 0.2650 | 0.7055 | 0.058* |
| H23B | 0.6947 | 0.2571 | 0.7529 | 0.058* |
| H23C | 0.6454 | 0.3046 | 0.7187 | 0.058* |
| C24 | 0.63430 (17) | 0.0377 (5) | 0.67480 (12) | 0.0558 (12) |
| H24A | 0.6005 | 0.0948 | 0.6695 | 0.067* |
| H24B | 0.6275 | -0.0740 | 0.6706 | 0.067* |
| H24C | 0.6579 | 0.0738 | 0.6558 | 0.067* |
| P3 | 0.61839 (4) | 0.01954 (11) | 0.42420 (3) | 0.0311 (2) |
| H3P | 0.6475 (12) | -0.009 (3) | 0.4597 (8) | 0.037* |
| C25 | 0.55825 (14) | -0.1067 (4) | 0.42541 (11) | 0.0376 (9) |
| C26 | 0.57715 (15) | -0.2653 (4) | 0.44388 (12) | 0.0525 (11) |
| H26A | 0.5990 | -0.2495 | 0.4708 | 0.063* |
| H26B | 0.5984 | -0.3181 | 0.4255 | 0.063* |
| H26C | 0.5461 | -0.3295 | 0.4470 | 0.063* |
| C27 | 0.52264 (14) | -0.0382 (5) | 0.45445 (11) | 0.0474 (11) |
| H27A | 0.4927 | -0.1089 | 0.4561 | 0.057* |
| H27B | 0.5089 | 0.0632 | 0.4440 | 0.057* |
| H27C | 0.5435 | -0.0250 | 0.4819 | 0.057* |
| C28 | 0.52641 (15) | -0.1280 (5) | 0.38235 (11) | 0.0515 (11) |
| H28A | 0.4946 | -0.1907 | 0.3841 | 0.062* |
| H28B | 0.5486 | -0.1811 | 0.3648 | 0.062* |
| H28C | 0.5157 | -0.0258 | 0.3706 | 0.062* |
| C29 | 0.60386 (14) | 0.2337 (4) | 0.42467 (11) | 0.0370 (9) |
| C30 | 0.65241 (14) | 0.3281 (4) | 0.41584 (12) | 0.0459 (10) |
| H30A | 0.6589 | 0.3063 | 0.3878 | 0.055* |
| H30B | 0.6839 | 0.2985 | 0.4355 | 0.055* |
| H30C | 0.6454 | 0.4395 | 0.4186 | 0.055* |
| C31 | 0.55516 (15) | 0.2753 (4) | 0.39264 (12) | 0.0497 (11) |
| H31A | 0.5235 | 0.2229 | 0.3997 | 0.060* |
| H31B | 0.5614 | 0.2412 | 0.3654 | 0.060* |
| H31C | 0.5496 | 0.3883 | 0.3925 | 0.060* |
| C32 | 0.59471 (15) | 0.2798 (4) | 0.46828 (11) | 0.0493 (11) |
| H32A | 0.5905 | 0.3930 | 0.4697 | 0.059* |
| H32B | 0.6254 | 0.2471 | 0.4883 | 0.059* |
| H32C | 0.5624 | 0.2285 | 0.4746 | 0.059* |
| C33 | 0.66206 (14) | -0.0426 (4) | 0.38595 (11) | 0.0371 (9) |
| C34 | 0.66708 (15) | -0.2203 (4) | 0.38602 (11) | 0.0465 (10) |
| H34A | 0.6323 | -0.2666 | 0.3756 | 0.056* |
| H34B | 0.6792 | -0.2567 | 0.4141 | 0.056* |

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|------|--------------|------------|--------------|-------------|
| H34C | 0.6930 | -0.2513 | 0.3683 | 0.056* |
| C35 | 0.71849 (14) | 0.0222 (4) | 0.39901 (12) | 0.0415 (10) |
| H35A | 0.7300 | 0.0018 | 0.4283 | 0.050* |
| H35B | 0.7184 | 0.1348 | 0.3940 | 0.050* |
| H35C | 0.7431 | -0.0285 | 0.3830 | 0.050* |
| C36 | 0.63950 (16) | 0.0148 (5) | 0.34197 (11) | 0.0517 (11) |
| H36A | 0.6623 | -0.0226 | 0.3226 | 0.062* |
| H36B | 0.6386 | 0.1289 | 0.3416 | 0.062* |
| H36C | 0.6033 | -0.0257 | 0.3340 | 0.062* |

Atomic displacement parameters (Å²)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|-------|-------------|-------------|-------------|--------------|-------------|--------------|
| Zn1 | 0.0415 (3) | 0.0430 (3) | 0.0322 (3) | -0.0021 (2) | 0.0062 (2) | 0.0010 (2) |
| Cl1 | 0.0715 (7) | 0.0485 (6) | 0.0376 (6) | 0.0113 (5) | 0.0117 (5) | -0.0014 (4) |
| Cl2 | 0.0556 (6) | 0.0456 (6) | 0.0373 (5) | -0.0081 (5) | 0.0119 (5) | 0.0038 (4) |
| Cl3 | 0.0462 (6) | 0.0697 (7) | 0.0492 (7) | -0.0127 (5) | 0.0097 (5) | 0.0043 (5) |
| O1 | 0.0489 (17) | 0.048 (2) | 0.0355 (17) | 0.0021 (15) | 0.0033 (14) | -0.0030 (13) |
| Zn2 | 0.0400 (3) | 0.0443 (3) | 0.0320 (3) | 0.0022 (2) | 0.0059 (2) | 0.0003 (2) |
| Cl4 | 0.0430 (6) | 0.0744 (8) | 0.0478 (7) | 0.0118 (5) | 0.0102 (5) | -0.0053 (5) |
| Cl5 | 0.0595 (7) | 0.0472 (6) | 0.0399 (6) | 0.0085 (5) | 0.0115 (5) | -0.0028 (4) |
| Cl6 | 0.0661 (7) | 0.0491 (6) | 0.0405 (6) | -0.0072 (5) | 0.0112 (5) | 0.0030 (5) |
| O2 | 0.0476 (16) | 0.0423 (19) | 0.0396 (18) | -0.0021 (15) | 0.0012 (13) | 0.0041 (13) |
| Zn3 | 0.0372 (3) | 0.0435 (3) | 0.0331 (3) | 0.0026 (2) | 0.0067 (2) | -0.0004 (2) |
| Cl7 | 0.0412 (6) | 0.0630 (7) | 0.0513 (7) | 0.0091 (5) | 0.0119 (5) | 0.0004 (5) |
| Cl8 | 0.0590 (7) | 0.0461 (6) | 0.0384 (5) | -0.0053 (5) | 0.0120 (5) | 0.0014 (4) |
| Cl9 | 0.0551 (6) | 0.0455 (6) | 0.0376 (5) | 0.0071 (5) | 0.0142 (5) | -0.0028 (4) |
| O3 | 0.0454 (16) | 0.0445 (19) | 0.0369 (17) | -0.0014 (15) | 0.0029 (13) | 0.0024 (12) |
| Cl11B | 0.065 (4) | 0.068 (4) | 0.043 (3) | 0.003 (3) | 0.009 (2) | -0.011 (3) |
| Cl2B | 0.039 (2) | 0.067 (5) | 0.043 (2) | -0.005 (2) | 0.0073 (17) | 0.000 (3) |
| C37B | 0.045 (4) | 0.073 (6) | 0.051 (4) | 0.006 (4) | 0.004 (4) | -0.008 (4) |
| C38B | 0.040 (4) | 0.078 (7) | 0.045 (4) | -0.002 (4) | 0.006 (3) | -0.009 (4) |
| Cl1A | 0.056 (3) | 0.073 (5) | 0.041 (3) | 0.004 (3) | 0.006 (2) | 0.014 (3) |
| Cl2A | 0.049 (3) | 0.080 (6) | 0.044 (3) | -0.009 (3) | 0.003 (2) | 0.012 (4) |
| C37A | 0.046 (4) | 0.075 (6) | 0.052 (4) | 0.003 (4) | 0.004 (4) | -0.004 (4) |
| C38A | 0.043 (4) | 0.072 (6) | 0.052 (5) | -0.002 (4) | 0.010 (4) | 0.001 (4) |
| Cl3B | 0.0523 (11) | 0.089 (2) | 0.0457 (12) | -0.0071 (14) | 0.0169 (9) | -0.0158 (14) |
| Cl4B | 0.067 (2) | 0.085 (2) | 0.0352 (16) | 0.0038 (14) | 0.0082 (15) | -0.0016 (12) |
| C39B | 0.043 (3) | 0.083 (4) | 0.059 (3) | -0.005 (3) | 0.015 (2) | -0.014 (3) |
| C40B | 0.045 (3) | 0.070 (4) | 0.054 (3) | -0.005 (3) | 0.009 (2) | -0.007 (2) |
| Cl4A | 0.051 (7) | 0.061 (9) | 0.040 (9) | 0.017 (6) | 0.000 (7) | -0.007 (7) |
| Cl3A | 0.022 (5) | 0.060 (9) | 0.021 (5) | 0.015 (5) | -0.004 (4) | -0.027 (5) |
| C39A | 0.037 (5) | 0.076 (6) | 0.051 (5) | 0.002 (5) | 0.003 (5) | -0.009 (5) |
| C40A | 0.044 (5) | 0.074 (6) | 0.051 (5) | 0.003 (5) | 0.003 (5) | -0.011 (5) |
| Cl5A | 0.044 (4) | 0.078 (6) | 0.034 (2) | -0.016 (3) | 0.009 (3) | -0.003 (3) |
| Cl6A | 0.049 (3) | 0.093 (6) | 0.038 (4) | -0.005 (4) | 0.010 (3) | -0.025 (4) |
| C41A | 0.040 (4) | 0.085 (7) | 0.049 (5) | -0.001 (5) | 0.016 (4) | -0.007 (5) |
| C42A | 0.042 (5) | 0.087 (6) | 0.049 (5) | 0.001 (4) | 0.004 (4) | 0.003 (4) |

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|------|-------------|------------|-------------|--------------|-------------|--------------|
| C15B | 0.0352 (17) | 0.096 (5) | 0.0475 (15) | 0.000 (2) | 0.0064 (12) | 0.006 (3) |
| C16B | 0.069 (4) | 0.079 (3) | 0.045 (2) | -0.012 (3) | 0.0077 (18) | 0.016 (2) |
| C41B | 0.040 (3) | 0.091 (6) | 0.042 (4) | 0.003 (4) | 0.006 (3) | 0.005 (4) |
| C42B | 0.044 (4) | 0.090 (5) | 0.052 (4) | -0.007 (4) | 0.002 (3) | 0.011 (3) |
| P1 | 0.0305 (6) | 0.0356 (6) | 0.0302 (6) | -0.0023 (4) | 0.0080 (5) | -0.0020 (4) |
| C1 | 0.037 (2) | 0.037 (2) | 0.043 (2) | -0.0040 (18) | 0.0054 (18) | -0.0059 (18) |
| C2 | 0.050 (3) | 0.048 (3) | 0.060 (3) | -0.005 (2) | 0.008 (2) | -0.008 (2) |
| C3 | 0.044 (2) | 0.038 (2) | 0.062 (3) | 0.0019 (19) | 0.009 (2) | -0.001 (2) |
| C4 | 0.062 (3) | 0.046 (3) | 0.055 (3) | -0.005 (2) | 0.018 (2) | 0.006 (2) |
| C5 | 0.042 (2) | 0.041 (2) | 0.030 (2) | -0.0003 (19) | 0.0141 (18) | 0.0025 (17) |
| C6 | 0.049 (3) | 0.066 (3) | 0.040 (3) | 0.005 (2) | 0.013 (2) | 0.002 (2) |
| C7 | 0.042 (2) | 0.047 (3) | 0.053 (3) | 0.000 (2) | 0.017 (2) | 0.0093 (19) |
| C8 | 0.034 (2) | 0.055 (3) | 0.045 (3) | 0.0057 (19) | 0.014 (2) | 0.0081 (19) |
| C9 | 0.030 (2) | 0.040 (2) | 0.041 (2) | 0.0006 (17) | 0.0104 (18) | -0.0001 (18) |
| C10 | 0.046 (3) | 0.067 (3) | 0.051 (3) | 0.017 (2) | 0.012 (2) | 0.007 (2) |
| C11 | 0.047 (3) | 0.051 (3) | 0.062 (3) | 0.002 (2) | 0.020 (2) | -0.007 (2) |
| C12 | 0.037 (2) | 0.061 (3) | 0.050 (3) | 0.002 (2) | 0.018 (2) | 0.001 (2) |
| P2 | 0.0354 (6) | 0.0385 (6) | 0.0332 (6) | -0.0001 (4) | 0.0051 (5) | 0.0025 (4) |
| C13 | 0.043 (2) | 0.056 (3) | 0.047 (3) | 0.009 (2) | 0.014 (2) | 0.013 (2) |
| C14 | 0.048 (3) | 0.075 (3) | 0.061 (3) | 0.014 (2) | 0.024 (2) | 0.021 (2) |
| C15 | 0.053 (3) | 0.070 (3) | 0.056 (3) | 0.016 (2) | 0.014 (2) | 0.016 (2) |
| C16 | 0.070 (3) | 0.049 (3) | 0.051 (3) | 0.015 (2) | 0.023 (2) | 0.002 (2) |
| C17 | 0.062 (3) | 0.047 (3) | 0.051 (3) | -0.012 (2) | -0.007 (2) | 0.007 (2) |
| C18 | 0.068 (3) | 0.062 (3) | 0.063 (3) | -0.021 (3) | 0.000 (3) | 0.012 (2) |
| C19 | 0.071 (3) | 0.071 (3) | 0.064 (3) | -0.025 (3) | -0.013 (3) | 0.004 (3) |
| C20 | 0.090 (4) | 0.039 (3) | 0.059 (3) | -0.002 (2) | -0.003 (3) | -0.004 (2) |
| C21 | 0.050 (3) | 0.048 (3) | 0.037 (2) | 0.003 (2) | 0.013 (2) | 0.0036 (18) |
| C22 | 0.044 (3) | 0.061 (3) | 0.045 (3) | 0.013 (2) | 0.014 (2) | 0.000 (2) |
| C23 | 0.048 (3) | 0.052 (3) | 0.050 (3) | -0.004 (2) | 0.019 (2) | 0.003 (2) |
| C24 | 0.061 (3) | 0.067 (3) | 0.038 (3) | 0.003 (2) | 0.003 (2) | 0.004 (2) |
| P3 | 0.0293 (5) | 0.0349 (5) | 0.0297 (6) | 0.0011 (4) | 0.0070 (4) | 0.0016 (4) |
| C25 | 0.035 (2) | 0.040 (2) | 0.040 (2) | -0.0016 (18) | 0.0122 (18) | 0.0009 (18) |
| C26 | 0.043 (3) | 0.052 (3) | 0.066 (3) | -0.005 (2) | 0.022 (2) | 0.008 (2) |
| C27 | 0.036 (2) | 0.061 (3) | 0.048 (3) | 0.000 (2) | 0.015 (2) | -0.002 (2) |
| C28 | 0.042 (2) | 0.064 (3) | 0.049 (3) | -0.013 (2) | 0.005 (2) | -0.008 (2) |
| C29 | 0.034 (2) | 0.034 (2) | 0.042 (2) | 0.0028 (17) | 0.0040 (18) | 0.0025 (17) |
| C30 | 0.044 (2) | 0.033 (2) | 0.061 (3) | 0.0018 (19) | 0.010 (2) | 0.0046 (19) |
| C31 | 0.044 (3) | 0.048 (3) | 0.056 (3) | 0.003 (2) | 0.004 (2) | 0.006 (2) |
| C32 | 0.046 (3) | 0.046 (3) | 0.058 (3) | 0.005 (2) | 0.013 (2) | -0.004 (2) |
| C33 | 0.035 (2) | 0.043 (2) | 0.035 (2) | -0.0020 (19) | 0.0113 (18) | -0.0051 (18) |
| C34 | 0.046 (2) | 0.044 (3) | 0.053 (3) | -0.001 (2) | 0.018 (2) | -0.0106 (19) |
| C35 | 0.036 (2) | 0.045 (2) | 0.046 (3) | -0.0022 (19) | 0.0138 (19) | -0.0045 (19) |
| C36 | 0.054 (3) | 0.069 (3) | 0.033 (2) | -0.006 (2) | 0.011 (2) | 0.001 (2) |

Geometric parameters (Å, °)

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|---------|-------------|----------|-----------|
| Zn1—C11 | 2.2690 (10) | C9—C12 | 1.536 (5) |
| Zn1—C12 | 2.2666 (10) | C10—H10A | 0.9800 |

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|-----------|-------------|----------|-----------|
| Zn1—C13 | 2.2219 (11) | C10—H10B | 0.9800 |
| Zn1—O1 | 2.024 (3) | C10—H10C | 0.9800 |
| O1—H10A | 0.817 (18) | C11—H11A | 0.9800 |
| O1—H10B | 0.821 (18) | C11—H11B | 0.9800 |
| Zn2—C14 | 2.2203 (11) | C11—H11C | 0.9800 |
| Zn2—C15 | 2.2666 (10) | C12—H12A | 0.9800 |
| Zn2—C16 | 2.2699 (10) | C12—H12B | 0.9800 |
| Zn2—O2 | 2.025 (3) | C12—H12C | 0.9800 |
| O2—H20A | 0.822 (18) | P2—H2P | 1.31 (3) |
| O2—H20B | 0.821 (18) | P2—C13 | 1.875 (4) |
| Zn3—C17 | 2.2199 (10) | P2—C17 | 1.860 (4) |
| Zn3—C18 | 2.2695 (10) | P2—C21 | 1.881 (4) |
| Zn3—C19 | 2.2686 (10) | C13—C14 | 1.542 (5) |
| Zn3—O3 | 2.028 (3) | C13—C15 | 1.529 (5) |
| O3—H30A | 0.847 (18) | C13—C16 | 1.539 (5) |
| O3—H30B | 0.833 (18) | C14—H14A | 0.9800 |
| C11B—C37B | 1.797 (7) | C14—H14B | 0.9800 |
| C12B—C38B | 1.793 (6) | C14—H14C | 0.9800 |
| C37B—H37A | 0.9900 | C15—H15A | 0.9800 |
| C37B—H37B | 0.9900 | C15—H15B | 0.9800 |
| C37B—C38B | 1.496 (7) | C15—H15C | 0.9800 |
| C38B—H38A | 0.9900 | C16—H16A | 0.9800 |
| C38B—H38B | 0.9900 | C16—H16B | 0.9800 |
| C11A—C37A | 1.798 (7) | C16—H16C | 0.9800 |
| C12A—C38A | 1.789 (7) | C17—C18 | 1.551 (5) |
| C37A—H37C | 0.9900 | C17—C19 | 1.542 (5) |
| C37A—H37D | 0.9900 | C17—C20 | 1.551 (5) |
| C37A—C38A | 1.495 (8) | C18—H18A | 0.9800 |
| C38A—H38C | 0.9900 | C18—H18B | 0.9800 |
| C38A—H38D | 0.9900 | C18—H18C | 0.9800 |
| C13B—C39B | 1.796 (4) | C19—H19A | 0.9800 |
| C14B—C40B | 1.799 (5) | C19—H19B | 0.9800 |
| C39B—H39A | 0.9900 | C19—H19C | 0.9800 |
| C39B—H39B | 0.9900 | C20—H20A | 0.9800 |
| C39B—C40B | 1.509 (6) | C20—H20B | 0.9800 |
| C40B—H40A | 0.9900 | C20—H20C | 0.9800 |
| C40B—H40B | 0.9900 | C21—C22 | 1.545 (5) |
| C14A—C40A | 1.797 (8) | C21—C23 | 1.538 (5) |
| C13A—C39A | 1.792 (8) | C21—C24 | 1.537 (5) |
| C39A—H39C | 0.9900 | C22—H22A | 0.9800 |
| C39A—H39D | 0.9900 | C22—H22B | 0.9800 |
| C39A—C40A | 1.493 (9) | C22—H22C | 0.9800 |
| C40A—H40C | 0.9900 | C23—H23A | 0.9800 |
| C40A—H40D | 0.9900 | C23—H23B | 0.9800 |
| C15A—C41A | 1.803 (7) | C23—H23C | 0.9800 |
| C16A—C42A | 1.790 (7) | C24—H24A | 0.9800 |
| C41A—H41A | 0.9900 | C24—H24B | 0.9800 |
| C41A—H41B | 0.9900 | C24—H24C | 0.9800 |

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|-------------|------------|---------------|-----------|
| C41A—C42A | 1.486 (8) | P3—H3P | 1.31 (3) |
| C42A—H42A | 0.9900 | P3—C25 | 1.877 (4) |
| C42A—H42B | 0.9900 | P3—C29 | 1.876 (4) |
| C15B—C41B | 1.803 (6) | P3—C33 | 1.879 (3) |
| C16B—C42B | 1.787 (6) | C25—C26 | 1.537 (5) |
| C41B—H41C | 0.9900 | C25—C27 | 1.531 (5) |
| C41B—H41D | 0.9900 | C25—C28 | 1.530 (5) |
| C41B—C42B | 1.484 (7) | C26—H26A | 0.9800 |
| C42B—H42C | 0.9900 | C26—H26B | 0.9800 |
| C42B—H42D | 0.9900 | C26—H26C | 0.9800 |
| P1—H1P | 1.31 (3) | C27—H27A | 0.9800 |
| P1—C1 | 1.868 (4) | C27—H27B | 0.9800 |
| P1—C5 | 1.882 (3) | C27—H27C | 0.9800 |
| P1—C9 | 1.879 (3) | C28—H28A | 0.9800 |
| C1—C2 | 1.532 (5) | C28—H28B | 0.9800 |
| C1—C3 | 1.541 (5) | C28—H28C | 0.9800 |
| C1—C4 | 1.540 (5) | C29—C30 | 1.539 (5) |
| C2—H2A | 0.9800 | C29—C31 | 1.537 (5) |
| C2—H2B | 0.9800 | C29—C32 | 1.541 (5) |
| C2—H2C | 0.9800 | C30—H30A | 0.9800 |
| C3—H3A | 0.9800 | C30—H30B | 0.9800 |
| C3—H3B | 0.9800 | C30—H30C | 0.9800 |
| C3—H3C | 0.9800 | C31—H31A | 0.9800 |
| C4—H4A | 0.9800 | C31—H31B | 0.9800 |
| C4—H4B | 0.9800 | C31—H31C | 0.9800 |
| C4—H4C | 0.9800 | C32—H32A | 0.9800 |
| C5—C6 | 1.528 (5) | C32—H32B | 0.9800 |
| C5—C7 | 1.539 (5) | C32—H32C | 0.9800 |
| C5—C8 | 1.533 (5) | C33—C34 | 1.530 (5) |
| C6—H6A | 0.9800 | C33—C35 | 1.534 (5) |
| C6—H6B | 0.9800 | C33—C36 | 1.552 (5) |
| C6—H6C | 0.9800 | C34—H34A | 0.9800 |
| C7—H7A | 0.9800 | C34—H34B | 0.9800 |
| C7—H7B | 0.9800 | C34—H34C | 0.9800 |
| C7—H7C | 0.9800 | C35—H35A | 0.9800 |
| C8—H8A | 0.9800 | C35—H35B | 0.9800 |
| C8—H8B | 0.9800 | C35—H35C | 0.9800 |
| C8—H8C | 0.9800 | C36—H36A | 0.9800 |
| C9—C10 | 1.527 (5) | C36—H36B | 0.9800 |
| C9—C11 | 1.547 (5) | C36—H36C | 0.9800 |
| C12—Zn1—C11 | 107.87 (4) | C9—C11—H11C | 109.5 |
| C13—Zn1—C11 | 116.22 (4) | H11A—C11—H11B | 109.5 |
| C13—Zn1—C12 | 114.58 (4) | H11A—C11—H11C | 109.5 |
| O1—Zn1—C11 | 102.71 (8) | H11B—C11—H11C | 109.5 |
| O1—Zn1—C12 | 104.68 (8) | C9—C12—H12A | 109.5 |
| O1—Zn1—C13 | 109.53 (8) | C9—C12—H12B | 109.5 |
| Zn1—O1—H10A | 103 (3) | C9—C12—H12C | 109.5 |

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| Zn1—O1—H1OB | 118 (3) | H12A—C12—H12B | 109.5 |
| H10A—O1—H1OB | 108 (4) | H12A—C12—H12C | 109.5 |
| Cl4—Zn2—Cl5 | 113.92 (4) | H12B—C12—H12C | 109.5 |
| Cl4—Zn2—Cl6 | 115.31 (4) | C13—P2—H2P | 103.5 (14) |
| Cl5—Zn2—Cl6 | 108.91 (4) | C13—P2—C21 | 113.85 (17) |
| O2—Zn2—Cl4 | 109.43 (8) | C17—P2—H2P | 105.3 (14) |
| O2—Zn2—Cl5 | 104.91 (8) | C17—P2—C13 | 114.82 (19) |
| O2—Zn2—Cl6 | 103.33 (8) | C17—P2—C21 | 114.73 (18) |
| Zn2—O2—H2OA | 105 (3) | C21—P2—H2P | 102.7 (14) |
| Zn2—O2—H2OB | 116 (3) | C14—C13—P2 | 110.9 (3) |
| H2OA—O2—H2OB | 116 (4) | C15—C13—P2 | 110.9 (3) |
| Cl7—Zn3—Cl8 | 115.54 (4) | C15—C13—C14 | 108.6 (3) |
| Cl7—Zn3—Cl9 | 114.59 (4) | C15—C13—C16 | 111.9 (3) |
| Cl9—Zn3—Cl8 | 108.37 (4) | C16—C13—P2 | 107.5 (3) |
| O3—Zn3—Cl7 | 109.29 (8) | C16—C13—C14 | 106.9 (3) |
| O3—Zn3—Cl8 | 102.93 (8) | C13—C14—H14A | 109.5 |
| O3—Zn3—Cl9 | 104.98 (8) | C13—C14—H14B | 109.5 |
| Zn3—O3—H3OA | 105 (3) | C13—C14—H14C | 109.5 |
| Zn3—O3—H3OB | 116 (3) | H14A—C14—H14B | 109.5 |
| H3OA—O3—H3OB | 115 (4) | H14A—C14—H14C | 109.5 |
| Cl1B—C37B—H37A | 109.2 | H14B—C14—H14C | 109.5 |
| Cl1B—C37B—H37B | 109.2 | C13—C15—H15A | 109.5 |
| H37A—C37B—H37B | 107.9 | C13—C15—H15B | 109.5 |
| C38B—C37B—Cl1B | 112.0 (11) | C13—C15—H15C | 109.5 |
| C38B—C37B—H37A | 109.2 | H15A—C15—H15B | 109.5 |
| C38B—C37B—H37B | 109.2 | H15A—C15—H15C | 109.5 |
| Cl2B—C38B—H38A | 109.8 | H15B—C15—H15C | 109.5 |
| Cl2B—C38B—H38B | 109.8 | C13—C16—H16A | 109.5 |
| C37B—C38B—Cl2B | 109.2 (10) | C13—C16—H16B | 109.5 |
| C37B—C38B—H38A | 109.8 | C13—C16—H16C | 109.5 |
| C37B—C38B—H38B | 109.8 | H16A—C16—H16B | 109.5 |
| H38A—C38B—H38B | 108.3 | H16A—C16—H16C | 109.5 |
| Cl1A—C37A—H37C | 110.0 | H16B—C16—H16C | 109.5 |
| Cl1A—C37A—H37D | 110.0 | C18—C17—P2 | 108.1 (3) |
| H37C—C37A—H37D | 108.4 | C18—C17—C20 | 106.4 (3) |
| C38A—C37A—Cl1A | 108.6 (12) | C19—C17—P2 | 112.0 (3) |
| C38A—C37A—H37C | 110.0 | C19—C17—C18 | 110.0 (3) |
| C38A—C37A—H37D | 110.0 | C19—C17—C20 | 110.2 (4) |
| Cl2A—C38A—H38C | 108.4 | C20—C17—P2 | 110.0 (3) |
| Cl2A—C38A—H38D | 108.4 | C17—C18—H18A | 109.5 |
| C37A—C38A—Cl2A | 115.4 (11) | C17—C18—H18B | 109.5 |
| C37A—C38A—H38C | 108.4 | C17—C18—H18C | 109.5 |
| C37A—C38A—H38D | 108.4 | H18A—C18—H18B | 109.5 |
| H38C—C38A—H38D | 107.5 | H18A—C18—H18C | 109.5 |
| Cl3B—C39B—H39A | 109.0 | H18B—C18—H18C | 109.5 |
| Cl3B—C39B—H39B | 109.0 | C17—C19—H19A | 109.5 |
| H39A—C39B—H39B | 107.8 | C17—C19—H19B | 109.5 |
| C40B—C39B—Cl3B | 112.8 (4) | C17—C19—H19C | 109.5 |

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| C40B—C39B—H39A | 109.0 | H19A—C19—H19B | 109.5 |
| C40B—C39B—H39B | 109.0 | H19A—C19—H19C | 109.5 |
| C14B—C40B—H40A | 109.9 | H19B—C19—H19C | 109.5 |
| C14B—C40B—H40B | 109.9 | C17—C20—H20A | 109.5 |
| C39B—C40B—C14B | 109.0 (5) | C17—C20—H20B | 109.5 |
| C39B—C40B—H40A | 109.9 | C17—C20—H20C | 109.5 |
| C39B—C40B—H40B | 109.9 | H20A—C20—H20B | 109.5 |
| H40A—C40B—H40B | 108.3 | H20A—C20—H20C | 109.5 |
| C13A—C39A—H39C | 111.5 | H20B—C20—H20C | 109.5 |
| C13A—C39A—H39D | 111.5 | C22—C21—P2 | 107.7 (2) |
| H39C—C39A—H39D | 109.3 | C23—C21—P2 | 110.7 (2) |
| C40A—C39A—C13A | 101 (2) | C23—C21—C22 | 106.6 (3) |
| C40A—C39A—H39C | 111.5 | C24—C21—P2 | 111.8 (3) |
| C40A—C39A—H39D | 111.5 | C24—C21—C22 | 110.3 (3) |
| C14A—C40A—H40C | 109.0 | C24—C21—C23 | 109.5 (3) |
| C14A—C40A—H40D | 109.0 | C21—C22—H22A | 109.5 |
| C39A—C40A—C14A | 113 (3) | C21—C22—H22B | 109.5 |
| C39A—C40A—H40C | 109.0 | C21—C22—H22C | 109.5 |
| C39A—C40A—H40D | 109.0 | H22A—C22—H22B | 109.5 |
| H40C—C40A—H40D | 107.8 | H22A—C22—H22C | 109.5 |
| C15A—C41A—H41A | 108.7 | H22B—C22—H22C | 109.5 |
| C15A—C41A—H41B | 108.7 | C21—C23—H23A | 109.5 |
| H41A—C41A—H41B | 107.6 | C21—C23—H23B | 109.5 |
| C42A—C41A—C15A | 114.2 (14) | C21—C23—H23C | 109.5 |
| C42A—C41A—H41A | 108.7 | H23A—C23—H23B | 109.5 |
| C42A—C41A—H41B | 108.7 | H23A—C23—H23C | 109.5 |
| C16A—C42A—H42A | 110.4 | H23B—C23—H23C | 109.5 |
| C16A—C42A—H42B | 110.4 | C21—C24—H24A | 109.5 |
| C41A—C42A—C16A | 106.7 (14) | C21—C24—H24B | 109.5 |
| C41A—C42A—H42A | 110.4 | C21—C24—H24C | 109.5 |
| C41A—C42A—H42B | 110.4 | H24A—C24—H24B | 109.5 |
| H42A—C42A—H42B | 108.6 | H24A—C24—H24C | 109.5 |
| C15B—C41B—H41C | 109.9 | H24B—C24—H24C | 109.5 |
| C15B—C41B—H41D | 109.9 | C25—P3—H3P | 102.8 (13) |
| H41C—C41B—H41D | 108.3 | C25—P3—C33 | 114.59 (16) |
| C42B—C41B—C15B | 109.0 (9) | C29—P3—H3P | 105.2 (13) |
| C42B—C41B—H41C | 109.9 | C29—P3—C25 | 113.83 (16) |
| C42B—C41B—H41D | 109.9 | C29—P3—C33 | 114.89 (16) |
| C16B—C42B—H42C | 108.8 | C33—P3—H3P | 103.6 (13) |
| C16B—C42B—H42D | 108.8 | C26—C25—P3 | 108.3 (2) |
| C41B—C42B—C16B | 113.7 (8) | C27—C25—P3 | 110.6 (3) |
| C41B—C42B—H42C | 108.8 | C27—C25—C26 | 105.7 (3) |
| C41B—C42B—H42D | 108.8 | C28—C25—P3 | 111.3 (2) |
| H42C—C42B—H42D | 107.7 | C28—C25—C26 | 110.4 (3) |
| C1—P1—H1P | 104.7 (13) | C28—C25—C27 | 110.3 (3) |
| C1—P1—C5 | 114.66 (16) | C25—C26—H26A | 109.5 |
| C1—P1—C9 | 114.03 (16) | C25—C26—H26B | 109.5 |
| C5—P1—H1P | 102.8 (14) | C25—C26—H26C | 109.5 |

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| C9—P1—H1P | 104.2 (14) | H26A—C26—H26B | 109.5 |
| C9—P1—C5 | 114.61 (16) | H26A—C26—H26C | 109.5 |
| C2—C1—P1 | 111.4 (3) | H26B—C26—H26C | 109.5 |
| C2—C1—C3 | 109.0 (3) | C25—C27—H27A | 109.5 |
| C2—C1—C4 | 111.0 (3) | C25—C27—H27B | 109.5 |
| C3—C1—P1 | 110.5 (2) | C25—C27—H27C | 109.5 |
| C4—C1—P1 | 108.6 (2) | H27A—C27—H27B | 109.5 |
| C4—C1—C3 | 106.2 (3) | H27A—C27—H27C | 109.5 |
| C1—C2—H2A | 109.5 | H27B—C27—H27C | 109.5 |
| C1—C2—H2B | 109.5 | C25—C28—H28A | 109.5 |
| C1—C2—H2C | 109.5 | C25—C28—H28B | 109.5 |
| H2A—C2—H2B | 109.5 | C25—C28—H28C | 109.5 |
| H2A—C2—H2C | 109.5 | H28A—C28—H28B | 109.5 |
| H2B—C2—H2C | 109.5 | H28A—C28—H28C | 109.5 |
| C1—C3—H3A | 109.5 | H28B—C28—H28C | 109.5 |
| C1—C3—H3B | 109.5 | C30—C29—P3 | 110.5 (2) |
| C1—C3—H3C | 109.5 | C30—C29—C32 | 106.3 (3) |
| H3A—C3—H3B | 109.5 | C31—C29—P3 | 111.1 (2) |
| H3A—C3—H3C | 109.5 | C31—C29—C30 | 109.3 (3) |
| H3B—C3—H3C | 109.5 | C31—C29—C32 | 111.0 (3) |
| C1—C4—H4A | 109.5 | C32—C29—P3 | 108.5 (2) |
| C1—C4—H4B | 109.5 | C29—C30—H30A | 109.5 |
| C1—C4—H4C | 109.5 | C29—C30—H30B | 109.5 |
| H4A—C4—H4B | 109.5 | C29—C30—H30C | 109.5 |
| H4A—C4—H4C | 109.5 | H30A—C30—H30B | 109.5 |
| H4B—C4—H4C | 109.5 | H30A—C30—H30C | 109.5 |
| C6—C5—P1 | 111.9 (3) | H30B—C30—H30C | 109.5 |
| C6—C5—C7 | 109.9 (3) | C29—C31—H31A | 109.5 |
| C6—C5—C8 | 110.2 (3) | C29—C31—H31B | 109.5 |
| C7—C5—P1 | 109.7 (2) | C29—C31—H31C | 109.5 |
| C8—C5—P1 | 108.7 (2) | H31A—C31—H31B | 109.5 |
| C8—C5—C7 | 106.3 (3) | H31A—C31—H31C | 109.5 |
| C5—C6—H6A | 109.5 | H31B—C31—H31C | 109.5 |
| C5—C6—H6B | 109.5 | C29—C32—H32A | 109.5 |
| C5—C6—H6C | 109.5 | C29—C32—H32B | 109.5 |
| H6A—C6—H6B | 109.5 | C29—C32—H32C | 109.5 |
| H6A—C6—H6C | 109.5 | H32A—C32—H32B | 109.5 |
| H6B—C6—H6C | 109.5 | H32A—C32—H32C | 109.5 |
| C5—C7—H7A | 109.5 | H32B—C32—H32C | 109.5 |
| C5—C7—H7B | 109.5 | C34—C33—P3 | 109.8 (2) |
| C5—C7—H7C | 109.5 | C34—C33—C35 | 106.7 (3) |
| H7A—C7—H7B | 109.5 | C34—C33—C36 | 109.7 (3) |
| H7A—C7—H7C | 109.5 | C35—C33—P3 | 109.3 (2) |
| H7B—C7—H7C | 109.5 | C35—C33—C36 | 109.9 (3) |
| C5—C8—H8A | 109.5 | C36—C33—P3 | 111.4 (3) |
| C5—C8—H8B | 109.5 | C33—C34—H34A | 109.5 |
| C5—C8—H8C | 109.5 | C33—C34—H34B | 109.5 |
| H8A—C8—H8B | 109.5 | C33—C34—H34C | 109.5 |

| | | | |
|---------------------|------------|----------------|------------|
| H8A—C8—H8C | 109.5 | H34A—C34—H34B | 109.5 |
| H8B—C8—H8C | 109.5 | H34A—C34—H34C | 109.5 |
| C10—C9—P1 | 112.1 (2) | H34B—C34—H34C | 109.5 |
| C10—C9—C11 | 111.0 (3) | C33—C35—H35A | 109.5 |
| C10—C9—C12 | 110.2 (3) | C33—C35—H35B | 109.5 |
| C11—C9—P1 | 107.1 (2) | C33—C35—H35C | 109.5 |
| C12—C9—P1 | 110.8 (3) | H35A—C35—H35B | 109.5 |
| C12—C9—C11 | 105.4 (3) | H35A—C35—H35C | 109.5 |
| C9—C10—H10A | 109.5 | H35B—C35—H35C | 109.5 |
| C9—C10—H10B | 109.5 | C33—C36—H36A | 109.5 |
| C9—C10—H10C | 109.5 | C33—C36—H36B | 109.5 |
| H10A—C10—H10B | 109.5 | C33—C36—H36C | 109.5 |
| H10A—C10—H10C | 109.5 | H36A—C36—H36B | 109.5 |
| H10B—C10—H10C | 109.5 | H36A—C36—H36C | 109.5 |
| C9—C11—H11A | 109.5 | H36B—C36—H36C | 109.5 |
| C9—C11—H11B | 109.5 | | |
| | | | |
| C11B—C37B—C38B—C12B | -68.6 (12) | C17—P2—C13—C14 | 36.4 (3) |
| C11A—C37A—C38A—C12A | -61.4 (14) | C17—P2—C13—C15 | -84.4 (3) |
| C13B—C39B—C40B—C14B | -67.1 (6) | C17—P2—C13—C16 | 152.9 (2) |
| C13A—C39A—C40A—C14A | 54 (4) | C17—P2—C21—C22 | -72.8 (3) |
| C15A—C41A—C42A—C16A | -64 (2) | C17—P2—C21—C23 | 171.0 (3) |
| C15B—C41B—C42B—C16B | -66.1 (12) | C17—P2—C21—C24 | 48.6 (3) |
| C1—P1—C5—C6 | -53.1 (3) | C21—P2—C13—C14 | 171.5 (3) |
| C1—P1—C5—C7 | -175.4 (2) | C21—P2—C13—C15 | 50.7 (3) |
| C1—P1—C5—C8 | 68.7 (3) | C21—P2—C13—C16 | -72.0 (3) |
| C1—P1—C9—C10 | 84.8 (3) | C21—P2—C17—C18 | 155.5 (3) |
| C1—P1—C9—C11 | -153.2 (2) | C21—P2—C17—C19 | -83.1 (3) |
| C1—P1—C9—C12 | -38.7 (3) | C21—P2—C17—C20 | 39.7 (3) |
| C5—P1—C1—C2 | 85.5 (3) | C25—P3—C29—C30 | -169.9 (2) |
| C5—P1—C1—C3 | -35.8 (3) | C25—P3—C29—C31 | -48.4 (3) |
| C5—P1—C1—C4 | -152.0 (2) | C25—P3—C29—C32 | 73.9 (3) |
| C5—P1—C9—C10 | -50.2 (3) | C25—P3—C33—C34 | -40.7 (3) |
| C5—P1—C9—C11 | 71.8 (3) | C25—P3—C33—C35 | -157.4 (2) |
| C5—P1—C9—C12 | -173.7 (2) | C25—P3—C33—C36 | 81.0 (3) |
| C9—P1—C1—C2 | -49.5 (3) | C29—P3—C25—C26 | -155.1 (2) |
| C9—P1—C1—C3 | -170.7 (2) | C29—P3—C25—C27 | -39.6 (3) |
| C9—P1—C1—C4 | 73.1 (3) | C29—P3—C25—C28 | 83.4 (3) |
| C9—P1—C5—C6 | 81.5 (3) | C29—P3—C33—C34 | -175.3 (2) |
| C9—P1—C5—C7 | -40.7 (3) | C29—P3—C33—C35 | 68.0 (3) |
| C9—P1—C5—C8 | -156.6 (2) | C29—P3—C33—C36 | -53.6 (3) |
| C13—P2—C17—C18 | -69.8 (3) | C33—P3—C25—C26 | 69.8 (3) |
| C13—P2—C17—C19 | 51.6 (4) | C33—P3—C25—C27 | -174.7 (2) |
| C13—P2—C17—C20 | 174.4 (3) | C33—P3—C25—C28 | -51.7 (3) |
| C13—P2—C21—C22 | 152.0 (3) | C33—P3—C29—C30 | -34.9 (3) |
| C13—P2—C21—C23 | 35.8 (3) | C33—P3—C29—C31 | 86.6 (3) |
| C13—P2—C21—C24 | -86.5 (3) | C33—P3—C29—C32 | -151.1 (2) |

Hydrogen-bond geometry (Å, °)

| <i>D</i> —H··· <i>A</i> | <i>D</i> —H | H··· <i>A</i> | <i>D</i> ··· <i>A</i> | <i>D</i> —H··· <i>A</i> |
|---------------------------------------|-------------|---------------|-----------------------|-------------------------|
| O1—H1 <i>OA</i> ···C12 ⁱ | 0.82 (2) | 2.37 (3) | 3.107 (3) | 150 (4) |
| O1—H1 <i>OB</i> ···C11 ⁱⁱ | 0.82 (2) | 2.27 (2) | 3.086 (3) | 173 (4) |
| O2—H2 <i>OA</i> ···C19 ⁱⁱⁱ | 0.82 (2) | 2.33 (2) | 3.120 (3) | 161 (4) |
| O2—H2 <i>OB</i> ···C18 ^{iv} | 0.82 (2) | 2.28 (2) | 3.095 (3) | 177 (4) |
| O3—H3 <i>OA</i> ···C15 ⁱⁱⁱ | 0.85 (2) | 2.30 (2) | 3.100 (3) | 158 (4) |
| O3—H3 <i>OB</i> ···C16 ^{iv} | 0.83 (2) | 2.28 (2) | 3.106 (3) | 173 (4) |

Symmetry codes: (i) $-x+1/2, y+1/2, -z+3/2$; (ii) $-x+1/2, y-1/2, -z+3/2$; (iii) $-x+3/2, y+1/2, -z+3/2$; (iv) $-x+3/2, y-1/2, -z+3/2$.