



Crystal structure of bromidopentakis(tetrahydrofuran- κ O)magnesium bis[1,2-bis(diphenylphosphanyl)benzene- κ^2P,P']cobaltate(−1) tetrahydrofuran disolvate

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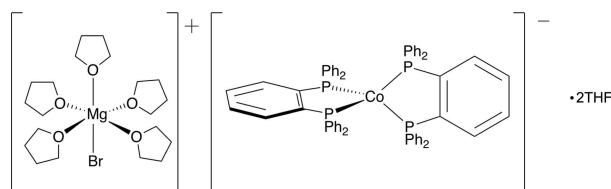
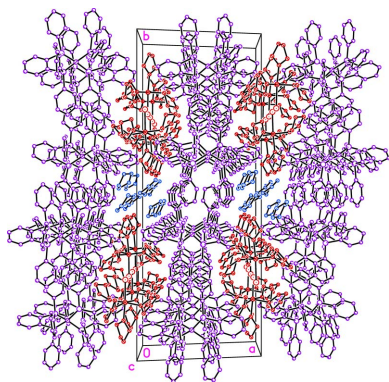
Supporting information: this article has supporting information at journals.iucr.org/e

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Structural characterization of the ionic title complex, $[\text{MgBr}(\text{THF})_5][\text{Co}(\text{dpbz})_2] \cdot 2\text{THF}$ [THF is tetrahydrofuran, $\text{C}_4\text{H}_8\text{O}$; dpbz is 1,2-bis(diphenylphosphanyl)benzene, $\text{C}_{30}\text{H}_{24}\text{P}_2$], revealed a well-separated cation and anion co-crystallized with two THF solvent molecules that interact with the cation *via* weak $\text{C}-\text{H} \cdots \text{O}$ contacts. The geometry about the cobalt center is pseudotetrahedral, as is expected for a d^{10} metal center, only deviating from an ideal tetrahedral geometry because of the restrictive bite angles of the bidentate phosphane ligands. Three THF ligands of the cation and one co-crystallized THF solvent molecule are each disordered over two orientations. In the extended structure, the cations and THF solvent molecules are arranged in (100) sheets that alternate with layers of anions, the latter of which show various π -interactions, which may explain the particular packing arrangement.

1. Chemical context

Phosphane ligands, especially aryl ones, have been used for many years to support transition metals in low oxidation states (Chatt & Watson, 1961; Chatt & Rowe, 1961). Bidentate phosphanes, or bisphosphanes, such as 1,2-bis(diphenylphosphanyl)benzene (dpbz), have the added benefit of the chelate effect (Cotton *et al.*, 1999). In an attempt to synthesize a cobalt(I) analog of the known iron(I) complex $\text{FeX}(\text{dpbz})_2$, $\text{X} = \text{Cl}, \text{Br}$, a species proposed to be an active catalyst in Negishi cross-coupling reactions (Adams *et al.*, 2012), CoBr_2 was reacted with four equivalents of *p*-tolylMgBr in tetrahydrofuran (THF) at 298 K. The unexpected result was a cobalt complex in the formal −1 oxidation state, formulated as $[\text{MgBr}(\text{THF})_5][\text{Co}(\text{dpbz})_2] \cdot 2\text{THF}$ **1** (Fig. 1). Herein we examine the crystal structure of **1** and compare it with the free bisphosphane and related cobalt species.



2. Structural commentary

The asymmetric unit of **1** contains one $[\text{MgBr}(\text{THF})_5]^+$ cation, one $[\text{Co}(\text{dpbz})_2]^-$ anion, and two co-crystallized THF solvent

Table 1
 Selected geometric parameters (Å, °).

| | | | |
|-----------|------------|-----------|------------|
| Co1–P1 | 2.1049 (6) | P2–C19 | 1.867 (2) |
| Co1–P2 | 2.0988 (6) | P2–C25 | 1.847 (2) |
| Co1–P3 | 2.0968 (6) | P3–C31 | 1.859 (2) |
| Co1–P4 | 2.1050 (6) | P3–C37 | 1.846 (2) |
| P1–C1 | 1.870 (2) | P3–C43 | 1.870 (2) |
| P1–C7 | 1.876 (2) | P4–C36 | 1.863 (2) |
| P1–C13 | 1.849 (2) | P4–C49 | 1.843 (2) |
| P2–C6 | 1.854 (2) | P4–C55 | 1.874 (2) |
| P1–Co1–P4 | 120.60 (3) | P3–Co1–P1 | 122.08 (3) |
| P2–Co1–P1 | 90.18 (2) | P3–Co1–P2 | 118.74 (3) |
| P2–Co1–P4 | 118.59 (2) | P3–Co1–P4 | 89.75 (2) |

molecules, all in general positions. The cation and anion are well separated. The average terminal P–Ph bond length in the anion of 1.859 (5) Å (Table 1) is about 0.02 Å longer than that observed in the free ligand [1.840 (2) Å, Levason *et al.*, 2006], which is consistent with backbonding from the d^{10} , formally Co^{-1} center into the σ^* orbitals of the P–C bonds. The average terminal P–Ph bond length in **1** of 1.861 (4) Å is identical to that found in $[\text{Co}(\text{dppe})_2]^-$ (dppe is 1,2-bis(diphenylphosphanyl)ethane), the only other structurally characterized four-coordinate bis(bisphosphane) cobalt(−) complex to date (Brennessel *et al.*, 2002).

The metal–phosphorus bond lengths are probably the best indicator that backbonding is occurring. The average Co–P bond lengths in **1** and $[\text{Co}(\text{dppe})_2]^-$ are 2.1014 (12) and 2.109 (1) Å, respectively. This distance increases by approximately 0.1 Å in structures containing $[\text{Co}(\text{dppe})_2]^+$ cations, for which the cobalt center is formally in the +1 oxidation state. The average Co–P bond lengths are 2.2032 (13) and 2.1930 (6) Å, respectively, for $[\text{Co}(\text{dppe})_2][\text{C}_{60}]$ -1,2-dichlorobenzene (Konarev *et al.*, 2011) and $[\text{Co}(\text{dppe})_2][\text{Ge}_9\{\text{Si}(\text{Me}_3)_3\}_3]\cdot\text{C}_7\text{H}_8$ (Kysliak *et al.*, 2016). The neutral Co^0 complex $\text{Co}(\text{dppp})_2$ (dppp = 1,2-bis(diphenylphosphanyl)propane; Kysliak *et al.*, 2016) has an average Co–P bond length of 2.173 (1) Å, which unsurprisingly lies between that of **1** and the two Co^{1+} cations.

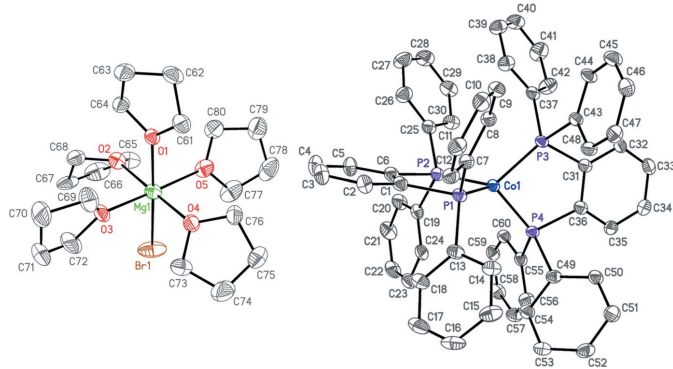

Figure 1
 Anisotropic displacement ellipsoid plot of **1** drawn at the 50% probability level with hydrogen atoms and solvent molecules omitted. Only the major component of the THF ligand disorder is shown. The reciprocal position of the two ions has been modified for clarity.

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

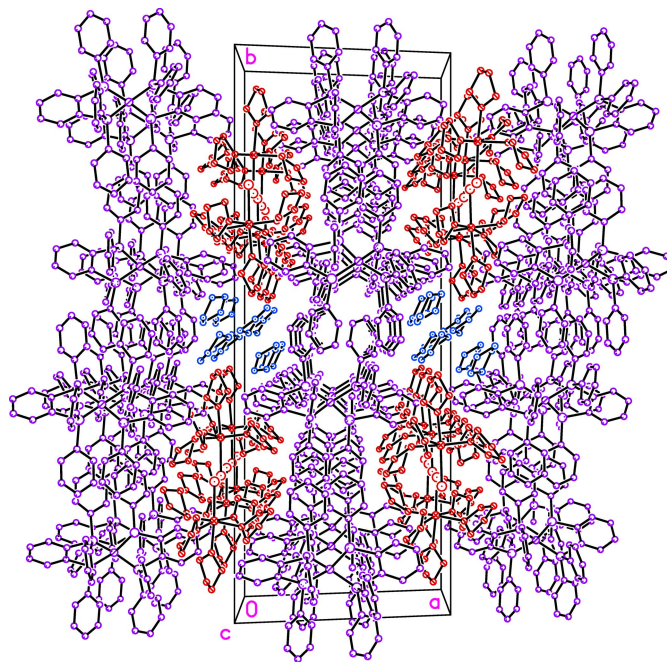
| $D-H\cdots A$ | $D-H$ | $H\cdots A$ | $D\cdots A$ | $D-H\cdots A$ |
|-----------------------|-------|-------------|-------------|---------------|
| C62–H62B \cdots O6 | 0.97 | 2.48 | 3.438 (4) | 167 |
| C63–H63B \cdots O7 | 0.97 | 2.59 | 3.555 (8) | 179 |
| C63–H63B \cdots O7' | 0.97 | 2.63 | 3.565 (6) | 162 |

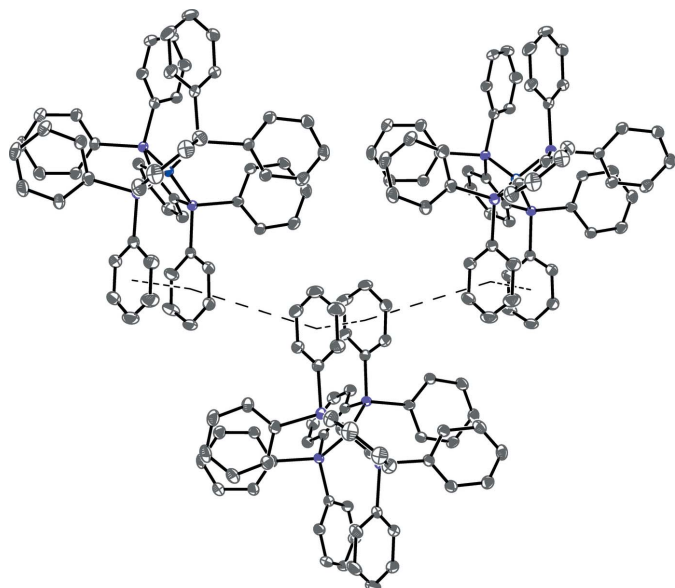
As is expected for a d^{10} cobalt center, the geometry of **1** is essentially tetrahedral, with a twist angle between the two P–Co–P planes of 89.465 (15)°, for which 90° would be ideal. The major deviation from perfect tetrahedral geometry, however, is due to the restrictive bite angles of the dpbz ligands [average 89.97 (3)°, Table 1].

Each terminal phenyl ring from one dpbz ligand is oriented to allow for possible parallel off-center π -system interactions (Martinez & Iverson, 2012) with those from the second dpbz ligand. The ring pair C25–C30/C37–C42 has the shortest centroid–centroid distance of 3.5325 (16) Å and the smallest angle between ring planes of 3.26 (13)°. Ring pairs C19–C24/C55–C60 and C13–C18/C49–C54 also have reasonable distances and angles of 3.8179 (15) and 4.0796 (16) Å and 11.66 (8) and 8.67 (16)°, respectively. Only the fourth pair, C13–C18/C43–C48, seems unlikely to have any significant intermolecular interaction with its analogous values of 4.4142 (11) Å and 36.99 (7)°.

3. Supramolecular features

Both co-crystallized THF solvent molecules interact with the cation *via* weak C–H \cdots O bonds (Table 2). The cations and


Figure 2
 In the bc planes are sheets of cations (red) and THF solvent molecules (blue) alternating with sheets of anions (purple). Hydrogen atoms have been omitted.

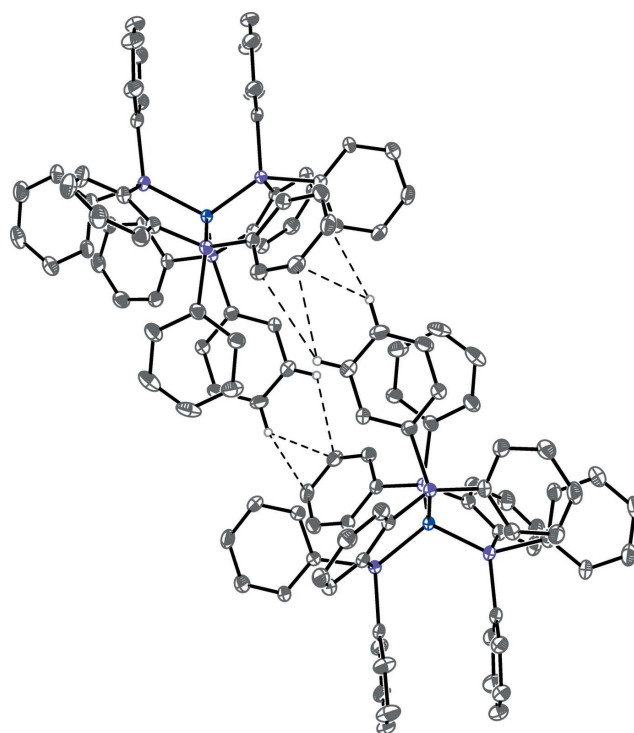

Figure 3

Anisotropic displacement ellipsoid plot of **1** drawn at the 50% probability level showing the extended intra- and intermolecular π -system interactions with hydrogen atoms omitted. The [001] direction (c axis) is to the right. (See Fig. 2 for view down [001].) Symmetry-equivalent molecules were generated by crystallographic twofold screw axes with symmetry operators $x, \frac{3}{2} - y, -\frac{1}{2} + z$ and $x, \frac{3}{2} - y, \frac{1}{2} + z$.

THF molecules of solvation are found in sheets normal to [100] that alternate with sheets of the anions (Fig. 2). Within each layer of anions there appear to be numerous potential π -system interactions (Martinez & Iverson, 2012; McGaughey *et al.*, 1998). Along [001] is an alternation between short intramolecular offset parallel stacking and longer intermolecular interactions with centroid–centroid distances of 3.533 (2) and 5.252 (2) Å, respectively (Fig. 3). On the opposite side of each molecule and also along the [001] direction is a second analogous set of potential π -system interactions, but with longer centroid–centroid distances of 4.080 (2) and 5.786 (2) Å; however, these rings are nearly coplanar (*i.e.* the open faces are not directed toward one another) and therefore they are unlikely to have any significant attractive intermolecular interactions. Upon further inspection, the one-dimensional chains along [001] are linked to other parallel chains by phenyl rings that are oriented correctly for edge-to-face C—H... π attractive interactions (Fig. 4), thus providing a possible explanation for the two-dimensional packing motif of anions in the bc planes.

4. Database survey

The only other structure containing a four-coordinate cobalt(−1) anion with two aryl bisphosphanes is the potassium 18-crown-6 salt of [Co(dppe)₂][−] (Brennessel *et al.*, 2002). Multiple species containing four-coordinate metals with two dpbz ligands are found in the Cambridge Structural Database (CSD, Version 5.40, November 2018; Groom *et al.*, 2016) with the following counts: Ni(dp bz)₂: five, Pt(dp bz)₂: two, [Cu(dp bz)₂]⁺: one, [Ag(dp bz)₂]⁺: five, [Au(dp bz)₂]⁺: thirteen.


Figure 4

Anisotropic displacement ellipsoid plot of **1** drawn at the 50% probability level of the edge-to-face π -system contacts that link the chains aligned along [001] in the [010] direction as well, thus offering an explanation for the observed two-dimensional sheets of anions. Hydrogen atoms except for those on carbon atoms C51 and C52 (and their symmetry equivalents) were omitted. The symmetry-equivalent molecule was generated by a crystallographic inversion center with symmetry operator $1 - x, 2 - y, 1 - z$.

Additionally there is one occurrence each of the square-planar cations [Rh(dp bz)₂]⁺ and [Ni(dp bz)₂]²⁺.

5. Synthesis and crystallization

CoBr₂ (99%, Sigma–Aldrich), dpbz (98%, Strem), *p*-tolyl-MgBr (1.0 M in THF, Sigma–Aldrich), THF (Sigma–Aldrich, anhydrous, 99.9%, inhibitor-free), and *n*-pentane (Sigma–Aldrich, >99%, anhydrous) were used in the synthesis of **1** without further purification. All reactions were performed in an MBraun inert-atmosphere (N₂) glovebox. CoBr₂ (27 mg, 0.12 mmol) and dpbz (99 mg, 0.22 mmol, 1.8 equiv.) were dissolved in 1 mL THF. *p*-TolylMgBr (494 μ L, 4 equiv.) was added to the cobalt solution at 0.33 mmol min^{−1} at room temperature. The resulting dark-red solution was allowed to stir at room temperature at 770 r.p.m. for 30 min. The solution was then filtered through Celite. Pentane (1 mL) was layered on top of the solution, and the solution was stored in a 243 K freezer until orange–brown crystalline blocks of **1** were observed.

6. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 3. Three THF ligands and one co-

Table 3
Experimental details.

| | |
|---|---|
| Crystal data | |
| Chemical formula | [MgBr(C ₄ H ₈ O) ₅][Co(C ₃₀ H ₂₄ P ₂) ₂]-2C ₄ H ₈ O |
| <i>M_r</i> | 1560.74 |
| Crystal system, space group | Monoclinic, <i>P</i> ₂ / <i>c</i> |
| Temperature (K) | 100 |
| <i>a</i> , <i>b</i> , <i>c</i> (Å) | 15.1096 (2), 38.1917 (3), 14.1266 (1) |
| β (°) | 106.102 (1) |
| <i>V</i> (Å ³) | 7832.11 (14) |
| <i>Z</i> | 4 |
| Radiation type | Cu <i>K</i> α |
| μ (mm ⁻¹) | 3.60 |
| Crystal size (mm) | 0.42 × 0.13 × 0.07 |
| Data collection | |
| Diffractometer | Rigaku XtaLAB Synergy, Dualflex, HyPix |
| Absorption correction | Multi-scan (<i>CrysAlis PRO</i> ; Rigaku OD, 2018) |
| <i>T</i> _{min} , <i>T</i> _{max} | 0.290, 1.000 |
| No. of measured, independent and observed [<i>I</i> > 2 σ (<i>I</i>)] reflections | 73364, 16393, 14853 |
| <i>R</i> _{int} | 0.048 |
| (sin θ / λ) _{max} (Å ⁻¹) | 0.634 |
| Refinement | |
| <i>R</i> [<i>F</i> ² > 2 σ (<i>F</i> ²)], <i>wR</i> (<i>F</i> ²), <i>S</i> | 0.048, 0.127, 1.06 |
| No. of reflections | 16393 |
| No. of parameters | 974 |
| No. of restraints | 91 |
| H-atom treatment | H-atom parameters constrained |
| $\Delta\rho_{\text{max}}$, $\Delta\rho_{\text{min}}$ (e Å ⁻³) | 0.64, -1.69 |

Computer programs: *CrysAlis PRO* (Rigaku OD, 2018), *SHELXT* (Sheldrick, 2015a), *SHELXL* (Sheldrick, 2015b) and *OLEX2* (Dolomanov *et al.*, 2009).

crystallized THF solvent molecule were modeled as disordered over two sets of site each: O2/C65–C68, 0.650 (8):0.350 (8), O3/C69–C72, 0.615 (8):0.385 (8), O5/C77–C80, 0.63 (2):0.37 (2), O7/C85–C88, 0.609 (4):0.391 (4). Analogous bond lengths and angles between the two positions of each disordered THF molecule were restrained to be similar. Anisotropic displacement parameters for proximal atoms were constrained to be equivalent.

H atoms were refined using riding models: aromatic, C–H = 0.93 Å, and methylene, C–H = 0.97 Å, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$.

The maximum residual peak of 0.64 e Å⁻³ and the deepest hole of -1.69 e Å⁻³ are found 0.84 and 0.83 Å from atoms H74A and Br1, respectively.

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References

- Adams, C. J., Bedford, R. B., Carter, E., Gower, N. J., Haddow, M. F., Harvey, J. N., Huwe, M., Cartes, M. A., Mansell, S. M., Mendoza, C., Murphy, D. M., Neeve, E. C. & Nunn, J. (2012). *J. Am. Chem. Soc.* **134**, 10333–10336.
- Brennessel, W. W., Young, V. G. Jr & Ellis, J. E. (2002). *Angew. Chem. Int. Ed.* **41**, 1211–1215.
- Chatt, J. & Rowe, G. A. (1961). *Nature*, **191**, 1191.
- Chatt, J. & Watson, H. R. (1961). *Nature*, **189**, 1003–1004.
- Cotton, F. A., Wilkinson, G., Murillo, C. A. & Bochmann, M. (1999). *Advanced Inorganic Chemistry* 6th ed., pp. 27–29. New York: John Wiley & Sons, Inc.
- Dolomanov, O. V., Bourhis, L. J., Gildea, R. J., Howard, J. A. K. & Puschmann, H. (2009). *J. Appl. Cryst.* **42**, 339–341.
- Groom, C. R., Bruno, I. J., Lightfoot, M. P. & Ward, S. C. (2016). *Acta Cryst.* **B72**, 171–179.
- Konarev, D. V., Kuźmin, A. V., Simonov, S. V., Khasanov, S. S., Yudanov, E. I. & Lyubovskaya, R. N. (2011). *Dalton Trans.* **40**, 4453–4458.
- Kysliak, O., Schrenk, C. & Schnepf, A. (2016). *Chem. Eur. J.* **22**, 18787–18793.
- Levason, W., Reid, G. & Webster, M. (2006). *Acta Cryst.* **C62**, o438–o440.
- Martinez, C. R. & Iverson, B. L. (2012). *Chem. Sci.* **3**, 2191–2201.
- McGaughey, G. B., Gagné, M. & Rappé, A. K. (1998). *J. Biol. Chem.* **273**, 15458–15463.
- Rigaku OD (2018). *CrysAlis PRO*. Rigaku Corporation, Yarnton, England.
- Sheldrick, G. M. (2015a). *Acta Cryst.* **A71**, 3–8.
- Sheldrick, G. M. (2015b). *Acta Cryst.* **C71**, 3–8.

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Computing details

Data collection: *CrysAlis PRO* (Rigaku OD, 2018); cell refinement: *CrysAlis PRO* (Rigaku OD, 2018); data reduction: *CrysAlis PRO* (Rigaku OD, 2018); program(s) used to solve structure: ShelXT (Sheldrick, 2015a); program(s) used to refine structure: *SHELXL* (Sheldrick, 2015b); molecular graphics: *OLEX2* (Dolomanov *et al.*, 2009); software used to prepare material for publication: *OLEX2* (Dolomanov *et al.*, 2009).

Bromidopentakis(tetrahydrofuran- κ O)magnesium bis[1,2-bis(diphenylphosphanyl)benzene- κ^2P,P']cobaltate(-1) tetrahydrofuran disolvate

Crystal data

[MgBr(C₄H₈O)₅][Co(C₃₀H₂₄P₂)₂] \cdot 2C₄H₈O

$M_r = 1560.74$

Monoclinic, $P2_1/c$

$a = 15.1096$ (2) Å

$b = 38.1917$ (3) Å

$c = 14.1266$ (1) Å

$\beta = 106.102$ (1)°

$V = 7832.11$ (14) Å³

$Z = 4$

$F(000) = 3288$

$D_x = 1.324$ Mg m⁻³

Cu $K\alpha$ radiation, $\lambda = 1.54184$ Å

Cell parameters from 33363 reflections

$\theta = 3.5\text{--}77.4^\circ$

$\mu = 3.60$ mm⁻¹

$T = 100$ K

Block, brown

$0.42 \times 0.13 \times 0.07$ mm

Data collection

Rigaku XtaLAB Synergy, Dualflex, HyPix diffractometer

Radiation source: micro-focus sealed X-ray tube, PhotonJet (Cu) X-ray Source

Mirror monochromator

ω scans

Absorption correction: multi-scan (CrysAlisPro; Rigaku OD, 2018)

$T_{\min} = 0.290$, $T_{\max} = 1.000$

73364 measured reflections

16393 independent reflections

14853 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.048$

$\theta_{\max} = 77.9^\circ$, $\theta_{\min} = 2.3^\circ$

$h = -17 \rightarrow 19$

$k = -48 \rightarrow 42$

$l = -17 \rightarrow 17$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.048$

$wR(F^2) = 0.127$

$S = 1.06$

16393 reflections

974 parameters

91 restraints

Primary atom site location: dual

Secondary atom site location: difference Fourier map

Hydrogen site location: inferred from neighbouring sites

H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0611P)^2 + 8.1124P]$$

where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.002$

$$\Delta\rho_{\max} = 0.64 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\min} = -1.69 \text{ e } \text{\AA}^{-3}$$

Special details

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

Refinement. Three THF ligands and one cocrystallized THF solvent molecule are modeled as disordered over two positions: O2/C65-C68, 0.650 (8):0.350 (8), O3/C69-C72, 0.615 (8):0.385 (8), O5/C77-C80, 0.63 (2):0.37 (2), O7/C85-C88, 0.609 (4):0.391 (4). Analogous bond lengths and angles between the two positions of each disordered THF molecule were restrained to be similar. Anisotropic displacement parameters for proximal atoms were constrained to be equivalent.

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

| | x | y | z | $U_{\text{iso}}^*/U_{\text{eq}}$ | Occ. (<1) |
|-----|--------------|-------------|--------------|----------------------------------|-----------|
| Co1 | 0.56200 (2) | 0.88268 (2) | 0.29552 (2) | 0.01606 (9) | |
| P1 | 0.67543 (4) | 0.90934 (2) | 0.38763 (4) | 0.01799 (11) | |
| P2 | 0.64893 (4) | 0.85876 (2) | 0.22175 (4) | 0.01868 (11) | |
| P3 | 0.47536 (4) | 0.85100 (2) | 0.35207 (4) | 0.01739 (11) | |
| P4 | 0.44633 (4) | 0.91051 (2) | 0.21208 (4) | 0.01730 (11) | |
| C1 | 0.78342 (16) | 0.89463 (6) | 0.35948 (16) | 0.0219 (4) | |
| C2 | 0.87300 (17) | 0.90505 (7) | 0.40934 (18) | 0.0271 (5) | |
| H2 | 0.882334 | 0.920804 | 0.461381 | 0.032* | |
| C3 | 0.94796 (18) | 0.89211 (7) | 0.38189 (19) | 0.0321 (5) | |
| H3 | 1.007331 | 0.898804 | 0.416351 | 0.038* | |
| C4 | 0.93442 (19) | 0.86910 (8) | 0.3028 (2) | 0.0352 (6) | |
| H4 | 0.984716 | 0.860446 | 0.284349 | 0.042* | |
| C5 | 0.84630 (17) | 0.85912 (7) | 0.25175 (18) | 0.0284 (5) | |
| H5 | 0.837316 | 0.844301 | 0.197676 | 0.034* | |
| C6 | 0.77029 (16) | 0.87121 (6) | 0.28094 (16) | 0.0228 (4) | |
| C7 | 0.71708 (15) | 0.90607 (6) | 0.52539 (15) | 0.0200 (4) | |
| C8 | 0.68730 (16) | 0.87745 (6) | 0.56995 (16) | 0.0216 (4) | |
| H8 | 0.649168 | 0.860760 | 0.531045 | 0.026* | |
| C9 | 0.71407 (17) | 0.87364 (6) | 0.67193 (17) | 0.0262 (5) | |
| H9 | 0.692940 | 0.854599 | 0.700506 | 0.031* | |
| C10 | 0.77197 (17) | 0.89796 (7) | 0.73140 (17) | 0.0264 (5) | |
| H10 | 0.789580 | 0.895353 | 0.799533 | 0.032* | |
| C11 | 0.80335 (17) | 0.92628 (6) | 0.68783 (17) | 0.0248 (5) | |
| H11 | 0.843014 | 0.942527 | 0.726858 | 0.030* | |
| C12 | 0.77547 (16) | 0.93034 (6) | 0.58588 (16) | 0.0227 (4) | |
| H12 | 0.796104 | 0.949558 | 0.557585 | 0.027* | |
| C13 | 0.68770 (16) | 0.95732 (6) | 0.37907 (15) | 0.0223 (4) | |
| C14 | 0.64101 (17) | 0.97900 (6) | 0.42872 (16) | 0.0246 (5) | |
| H14 | 0.604953 | 0.969028 | 0.465297 | 0.029* | |
| C15 | 0.64753 (19) | 1.01517 (6) | 0.42440 (18) | 0.0307 (5) | |
| H15 | 0.617267 | 1.029186 | 0.459358 | 0.037* | |

| | | | | |
|-----|--------------|-------------|---------------|------------|
| C16 | 0.6992 (2) | 1.03052 (6) | 0.36797 (19) | 0.0343 (6) |
| H16 | 0.704354 | 1.054750 | 0.365632 | 0.041* |
| C17 | 0.74250 (19) | 1.00939 (7) | 0.31555 (19) | 0.0346 (6) |
| H17 | 0.775663 | 1.019538 | 0.276206 | 0.042* |
| C18 | 0.73743 (18) | 0.97309 (6) | 0.32057 (17) | 0.0277 (5) |
| H18 | 0.767287 | 0.959235 | 0.284812 | 0.033* |
| C19 | 0.64638 (16) | 0.87156 (6) | 0.09346 (16) | 0.0227 (4) |
| C20 | 0.65586 (18) | 0.84885 (7) | 0.01949 (17) | 0.0280 (5) |
| H20 | 0.667014 | 0.825189 | 0.033357 | 0.034* |
| C21 | 0.6487 (2) | 0.86137 (8) | −0.07511 (19) | 0.0346 (6) |
| H21 | 0.653822 | 0.845911 | −0.124195 | 0.042* |
| C22 | 0.63415 (19) | 0.89669 (8) | −0.09651 (18) | 0.0330 (6) |
| H22 | 0.629573 | 0.904947 | −0.159615 | 0.040* |
| C23 | 0.62645 (17) | 0.91957 (7) | −0.02309 (18) | 0.0291 (5) |
| H23 | 0.617585 | 0.943347 | −0.036667 | 0.035* |
| C24 | 0.63193 (16) | 0.90709 (6) | 0.07081 (17) | 0.0240 (5) |
| H24 | 0.625892 | 0.922629 | 0.119317 | 0.029* |
| C25 | 0.65782 (16) | 0.81068 (6) | 0.21354 (16) | 0.0224 (4) |
| C26 | 0.73116 (18) | 0.79079 (7) | 0.26938 (19) | 0.0304 (5) |
| H26 | 0.782408 | 0.801993 | 0.309945 | 0.036* |
| C27 | 0.7287 (2) | 0.75444 (7) | 0.2652 (2) | 0.0388 (6) |
| H27 | 0.778131 | 0.741617 | 0.303426 | 0.047* |
| C28 | 0.6540 (2) | 0.73719 (7) | 0.2053 (2) | 0.0368 (6) |
| H28 | 0.653172 | 0.712870 | 0.201900 | 0.044* |
| C29 | 0.5805 (2) | 0.75637 (7) | 0.15036 (19) | 0.0319 (5) |
| H29 | 0.529601 | 0.744936 | 0.109823 | 0.038* |
| C30 | 0.58174 (18) | 0.79270 (6) | 0.15510 (17) | 0.0274 (5) |
| H30 | 0.530978 | 0.805271 | 0.118672 | 0.033* |
| C31 | 0.35262 (16) | 0.86047 (6) | 0.28835 (15) | 0.0201 (4) |
| C32 | 0.27655 (16) | 0.84320 (6) | 0.30413 (17) | 0.0231 (4) |
| H32 | 0.285582 | 0.824841 | 0.349060 | 0.028* |
| C33 | 0.18766 (17) | 0.85306 (6) | 0.25364 (18) | 0.0272 (5) |
| H33 | 0.137412 | 0.840906 | 0.262961 | 0.033* |
| C34 | 0.17416 (17) | 0.88139 (6) | 0.18868 (18) | 0.0266 (5) |
| H34 | 0.114784 | 0.888345 | 0.155026 | 0.032* |
| C35 | 0.24949 (16) | 0.89911 (6) | 0.17452 (16) | 0.0229 (4) |
| H35 | 0.240158 | 0.918037 | 0.131435 | 0.028* |
| C36 | 0.33930 (16) | 0.88898 (6) | 0.22401 (15) | 0.0200 (4) |
| C37 | 0.48157 (16) | 0.80276 (6) | 0.34774 (15) | 0.0202 (4) |
| C38 | 0.56174 (17) | 0.78692 (6) | 0.40489 (17) | 0.0256 (5) |
| H38 | 0.609041 | 0.800857 | 0.442388 | 0.031* |
| C39 | 0.57257 (19) | 0.75087 (6) | 0.40711 (17) | 0.0290 (5) |
| H39 | 0.625634 | 0.740837 | 0.447858 | 0.035* |
| C40 | 0.5045 (2) | 0.72977 (6) | 0.34878 (19) | 0.0313 (5) |
| H40 | 0.511021 | 0.705546 | 0.350722 | 0.038* |
| C41 | 0.4267 (2) | 0.74522 (7) | 0.2875 (2) | 0.0373 (6) |
| H41 | 0.381890 | 0.731312 | 0.245942 | 0.045* |
| C42 | 0.41491 (19) | 0.78124 (6) | 0.28750 (19) | 0.0308 (5) |

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|------|--------------|--------------|---------------|--------------|-----------|
| H42 | 0.361715 | 0.791149 | 0.246673 | 0.037* | |
| C43 | 0.46386 (16) | 0.85598 (5) | 0.47980 (16) | 0.0199 (4) | |
| C44 | 0.45917 (17) | 0.82847 (6) | 0.54279 (16) | 0.0237 (5) | |
| H44 | 0.455585 | 0.805586 | 0.519505 | 0.028* | |
| C45 | 0.45976 (19) | 0.83461 (6) | 0.64000 (18) | 0.0294 (5) | |
| H45 | 0.459109 | 0.815859 | 0.681794 | 0.035* | |
| C46 | 0.46132 (18) | 0.86856 (7) | 0.67472 (17) | 0.0270 (5) | |
| H46 | 0.462255 | 0.872724 | 0.739850 | 0.032* | |
| C47 | 0.46148 (17) | 0.89631 (6) | 0.61135 (17) | 0.0249 (5) | |
| H47 | 0.460033 | 0.919183 | 0.633482 | 0.030* | |
| C48 | 0.46377 (16) | 0.89022 (6) | 0.51557 (16) | 0.0222 (4) | |
| H48 | 0.465280 | 0.909073 | 0.474367 | 0.027* | |
| C49 | 0.42490 (16) | 0.95620 (6) | 0.24138 (15) | 0.0207 (4) | |
| C50 | 0.36400 (16) | 0.96529 (6) | 0.29604 (16) | 0.0228 (4) | |
| H50 | 0.331658 | 0.947800 | 0.317968 | 0.027* | |
| C51 | 0.35115 (18) | 0.99999 (6) | 0.31800 (18) | 0.0278 (5) | |
| H51 | 0.310294 | 1.005477 | 0.354377 | 0.033* | |
| C52 | 0.39851 (18) | 1.02656 (6) | 0.28634 (18) | 0.0269 (5) | |
| H52 | 0.388231 | 1.049855 | 0.299325 | 0.032* | |
| C53 | 0.46147 (17) | 1.01792 (6) | 0.23501 (17) | 0.0254 (5) | |
| H53 | 0.494433 | 1.035528 | 0.214478 | 0.030* | |
| C54 | 0.47575 (16) | 0.98322 (6) | 0.21399 (16) | 0.0219 (4) | |
| H54 | 0.519613 | 0.977807 | 0.181308 | 0.026* | |
| C55 | 0.41173 (15) | 0.91347 (6) | 0.07424 (15) | 0.0207 (4) | |
| C56 | 0.38055 (17) | 0.94338 (6) | 0.01859 (16) | 0.0253 (5) | |
| H56 | 0.371206 | 0.963866 | 0.050098 | 0.030* | |
| C57 | 0.36315 (18) | 0.94309 (7) | −0.08338 (17) | 0.0287 (5) | |
| H57 | 0.343390 | 0.963431 | −0.119193 | 0.034* | |
| C58 | 0.37506 (18) | 0.91275 (7) | −0.13187 (16) | 0.0282 (5) | |
| H58 | 0.363781 | 0.912604 | −0.200015 | 0.034* | |
| C59 | 0.40407 (17) | 0.88249 (6) | −0.07749 (17) | 0.0254 (5) | |
| H59 | 0.411482 | 0.861881 | −0.109503 | 0.030* | |
| C60 | 0.42199 (16) | 0.88287 (6) | 0.02395 (16) | 0.0218 (4) | |
| H60 | 0.441202 | 0.862402 | 0.059370 | 0.026* | |
| Br1 | 0.07583 (2) | 0.75114 (2) | 0.27391 (3) | 0.05868 (12) | |
| Mg1 | 0.07915 (6) | 0.68385 (2) | 0.30221 (6) | 0.02865 (18) | |
| O1 | 0.09259 (12) | 0.62946 (4) | 0.33147 (12) | 0.0261 (3) | |
| O2 | 0.2044 (6) | 0.6882 (4) | 0.4168 (6) | 0.0328 (14) | 0.650 (8) |
| C65 | 0.2808 (13) | 0.7075 (4) | 0.3963 (9) | 0.0435 (19) | 0.650 (8) |
| H65A | 0.260351 | 0.719528 | 0.333538 | 0.052* | 0.650 (8) |
| H65B | 0.330335 | 0.691638 | 0.394153 | 0.052* | 0.650 (8) |
| C66 | 0.3132 (7) | 0.7336 (2) | 0.4795 (7) | 0.061 (2) | 0.650 (8) |
| H66A | 0.312449 | 0.757270 | 0.454073 | 0.073* | 0.650 (8) |
| H66B | 0.375151 | 0.728174 | 0.519051 | 0.073* | 0.650 (8) |
| C67 | 0.2439 (5) | 0.72976 (13) | 0.5401 (4) | 0.0528 (15) | 0.650 (8) |
| H67A | 0.272513 | 0.734185 | 0.609470 | 0.063* | 0.650 (8) |
| H67B | 0.191779 | 0.745358 | 0.516600 | 0.063* | 0.650 (8) |
| C68 | 0.2156 (6) | 0.69171 (19) | 0.5216 (6) | 0.0409 (17) | 0.650 (8) |

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|------|---------------|-------------|--------------|-------------|-----------|
| H68A | 0.263053 | 0.676108 | 0.559338 | 0.049* | 0.650 (8) |
| H68B | 0.158368 | 0.687026 | 0.537742 | 0.049* | 0.650 (8) |
| O2' | 0.2129 (11) | 0.6891 (7) | 0.4020 (12) | 0.0328 (14) | 0.350 (8) |
| C65' | 0.283 (3) | 0.7096 (8) | 0.3735 (18) | 0.0435 (19) | 0.350 (8) |
| H65C | 0.259385 | 0.718214 | 0.306607 | 0.052* | 0.350 (8) |
| H65D | 0.337444 | 0.695667 | 0.378115 | 0.052* | 0.350 (8) |
| C66' | 0.3036 (16) | 0.7397 (5) | 0.4469 (13) | 0.061 (2) | 0.350 (8) |
| H66C | 0.366119 | 0.748169 | 0.457361 | 0.073* | 0.350 (8) |
| H66D | 0.260966 | 0.759007 | 0.425123 | 0.073* | 0.350 (8) |
| C67' | 0.2905 (10) | 0.7227 (3) | 0.5392 (8) | 0.0528 (15) | 0.350 (8) |
| H67C | 0.349808 | 0.718043 | 0.585804 | 0.063* | 0.350 (8) |
| H67D | 0.255683 | 0.738016 | 0.570256 | 0.063* | 0.350 (8) |
| C68' | 0.2388 (12) | 0.6888 (4) | 0.5082 (12) | 0.0409 (17) | 0.350 (8) |
| H68C | 0.277851 | 0.668846 | 0.533571 | 0.049* | 0.350 (8) |
| H68D | 0.184670 | 0.687730 | 0.532226 | 0.049* | 0.350 (8) |
| O3 | -0.0021 (10) | 0.6865 (6) | 0.4036 (10) | 0.034 (2) | 0.615 (8) |
| C69 | -0.053 (2) | 0.6572 (5) | 0.429 (2) | 0.0478 (18) | 0.615 (8) |
| H69A | -0.015193 | 0.636296 | 0.440744 | 0.057* | 0.615 (8) |
| H69B | -0.108062 | 0.652537 | 0.375647 | 0.057* | 0.615 (8) |
| C70 | -0.0782 (5) | 0.6678 (2) | 0.5203 (5) | 0.0741 (19) | 0.615 (8) |
| H70A | -0.134850 | 0.656603 | 0.523682 | 0.089* | 0.615 (8) |
| H70B | -0.029452 | 0.661916 | 0.579042 | 0.089* | 0.615 (8) |
| C71 | -0.0899 (8) | 0.7065 (3) | 0.5088 (7) | 0.092 (3) | 0.615 (8) |
| H71A | -0.150997 | 0.712406 | 0.467801 | 0.111* | 0.615 (8) |
| H71B | -0.079965 | 0.717898 | 0.572281 | 0.111* | 0.615 (8) |
| C72 | -0.018 (3) | 0.7169 (6) | 0.460 (3) | 0.069 (3) | 0.615 (8) |
| H72A | -0.038528 | 0.736769 | 0.417164 | 0.083* | 0.615 (8) |
| H72B | 0.038596 | 0.723300 | 0.509514 | 0.083* | 0.615 (8) |
| O3' | 0.0124 (19) | 0.6851 (10) | 0.4153 (18) | 0.034 (2) | 0.385 (8) |
| C69' | -0.054 (3) | 0.6578 (9) | 0.420 (4) | 0.0478 (18) | 0.385 (8) |
| H69C | -0.023401 | 0.637953 | 0.457594 | 0.057* | 0.385 (8) |
| H69D | -0.086481 | 0.649934 | 0.353927 | 0.057* | 0.385 (8) |
| C70' | -0.1195 (8) | 0.6747 (4) | 0.4689 (8) | 0.0741 (19) | 0.385 (8) |
| H70C | -0.170933 | 0.685534 | 0.421139 | 0.089* | 0.385 (8) |
| H70D | -0.142676 | 0.657841 | 0.507567 | 0.089* | 0.385 (8) |
| C71' | -0.0591 (12) | 0.7015 (5) | 0.5333 (13) | 0.092 (3) | 0.385 (8) |
| H71C | -0.016780 | 0.690800 | 0.590404 | 0.111* | 0.385 (8) |
| H71D | -0.095267 | 0.719204 | 0.554886 | 0.111* | 0.385 (8) |
| C72' | -0.009 (5) | 0.7164 (9) | 0.464 (5) | 0.069 (3) | 0.385 (8) |
| H72C | -0.048369 | 0.732368 | 0.417296 | 0.083* | 0.385 (8) |
| H72D | 0.046244 | 0.728652 | 0.499914 | 0.083* | 0.385 (8) |
| O4 | -0.04687 (12) | 0.67674 (5) | 0.19181 (13) | 0.0308 (4) | |
| O5 | 0.14741 (14) | 0.67323 (6) | 0.19244 (15) | 0.0393 (4) | |
| C61 | 0.0442 (2) | 0.60163 (7) | 0.2688 (2) | 0.0357 (6) | |
| H61A | 0.048397 | 0.604591 | 0.201990 | 0.043* | |
| H61B | -0.020300 | 0.601421 | 0.267637 | 0.043* | |
| C62 | 0.0908 (2) | 0.56810 (7) | 0.3130 (2) | 0.0388 (6) | |
| H62A | 0.143673 | 0.562937 | 0.288974 | 0.047* | |

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|------|---------------|--------------|--------------|-------------|----------|
| H62B | 0.048477 | 0.548477 | 0.298570 | 0.047* | |
| C63 | 0.1199 (2) | 0.57638 (7) | 0.4218 (2) | 0.0378 (6) | |
| H63A | 0.170251 | 0.561467 | 0.457036 | 0.045* | |
| H63B | 0.068850 | 0.573869 | 0.450490 | 0.045* | |
| C64 | 0.15005 (19) | 0.61415 (6) | 0.42258 (19) | 0.0307 (5) | |
| H64A | 0.140981 | 0.626320 | 0.479427 | 0.037* | |
| H64B | 0.214657 | 0.615542 | 0.424626 | 0.037* | |
| C73 | -0.1252 (2) | 0.69921 (7) | 0.1900 (2) | 0.0387 (6) | |
| H73A | -0.104208 | 0.721455 | 0.221390 | 0.046* | |
| H73B | -0.164713 | 0.688165 | 0.224965 | 0.046* | |
| C74 | -0.1766 (3) | 0.70470 (10) | 0.0843 (3) | 0.0631 (10) | |
| H74A | -0.174948 | 0.729176 | 0.066577 | 0.076* | |
| H74B | -0.240390 | 0.697599 | 0.072408 | 0.076* | |
| C75 | -0.1287 (2) | 0.68231 (8) | 0.0248 (2) | 0.0430 (7) | |
| H75A | -0.089869 | 0.696463 | -0.004307 | 0.052* | |
| H75B | -0.173152 | 0.669851 | -0.027032 | 0.052* | |
| C76 | -0.0716 (2) | 0.65707 (7) | 0.10046 (19) | 0.0340 (6) | |
| H76A | -0.107210 | 0.636466 | 0.106415 | 0.041* | |
| H76B | -0.016999 | 0.649800 | 0.082313 | 0.041* | |
| C77 | 0.1439 (18) | 0.6960 (4) | 0.1093 (10) | 0.0552 (19) | 0.63 (2) |
| H77A | 0.191600 | 0.713717 | 0.126867 | 0.066* | 0.63 (2) |
| H77B | 0.084462 | 0.707410 | 0.087106 | 0.066* | 0.63 (2) |
| C78 | 0.1598 (10) | 0.6716 (3) | 0.0293 (8) | 0.065 (2) | 0.63 (2) |
| H78A | 0.102009 | 0.665872 | -0.018558 | 0.077* | 0.63 (2) |
| H78B | 0.200498 | 0.682462 | -0.004627 | 0.077* | 0.63 (2) |
| C77' | 0.136 (3) | 0.6945 (7) | 0.1048 (19) | 0.0552 (19) | 0.37 (2) |
| H77C | 0.156540 | 0.718280 | 0.122321 | 0.066* | 0.37 (2) |
| H77D | 0.071724 | 0.695078 | 0.066605 | 0.066* | 0.37 (2) |
| C78' | 0.1951 (15) | 0.6769 (3) | 0.0463 (15) | 0.065 (2) | 0.37 (2) |
| H78C | 0.254901 | 0.688101 | 0.058996 | 0.077* | 0.37 (2) |
| H78D | 0.164921 | 0.677615 | -0.023935 | 0.077* | 0.37 (2) |
| C79 | 0.2041 (2) | 0.63911 (11) | 0.0853 (2) | 0.0565 (9) | |
| H79A | 0.163914 | 0.618969 | 0.066437 | 0.068* | 0.63 (2) |
| H79B | 0.262407 | 0.634057 | 0.072041 | 0.068* | 0.63 (2) |
| H79C | 0.148479 | 0.625563 | 0.058270 | 0.068* | 0.37 (2) |
| H79D | 0.256559 | 0.627066 | 0.073572 | 0.068* | 0.37 (2) |
| C80 | 0.2183 (2) | 0.64766 (9) | 0.1924 (2) | 0.0442 (7) | |
| H80A | 0.211180 | 0.626936 | 0.229260 | 0.053* | |
| H80B | 0.279190 | 0.657418 | 0.220931 | 0.053* | |
| O6 | -0.05333 (17) | 0.49866 (6) | 0.22536 (16) | 0.0474 (5) | |
| C81 | -0.0188 (3) | 0.49417 (10) | 0.1421 (2) | 0.0534 (8) | |
| H81A | 0.038407 | 0.481068 | 0.160298 | 0.064* | |
| H81B | -0.007326 | 0.516744 | 0.116385 | 0.064* | |
| C82 | -0.0908 (2) | 0.47449 (9) | 0.0663 (3) | 0.0492 (8) | |
| H82A | -0.062744 | 0.458644 | 0.029350 | 0.059* | |
| H82B | -0.131323 | 0.490446 | 0.020755 | 0.059* | |
| C83 | -0.1428 (3) | 0.45449 (9) | 0.1261 (3) | 0.0595 (10) | |
| H83A | -0.208385 | 0.459082 | 0.102317 | 0.071* | |

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| H83B | -0.132509 | 0.429510 | 0.122733 | 0.071* | |
| C84 | -0.1046 (2) | 0.46776 (10) | 0.2301 (3) | 0.0582 (9) | |
| H84A | -0.154348 | 0.472953 | 0.259008 | 0.070* | |
| H84B | -0.065064 | 0.450215 | 0.270418 | 0.070* | |
| O7 | -0.0684 (6) | 0.5669 (2) | 0.5248 (6) | 0.0581 (11) | 0.391 (4) |
| C85 | -0.1238 (7) | 0.5650 (3) | 0.5912 (8) | 0.0498 (14) | 0.391 (4) |
| H85A | -0.171262 | 0.582890 | 0.575925 | 0.060* | 0.391 (4) |
| H85B | -0.086465 | 0.568282 | 0.658530 | 0.060* | 0.391 (4) |
| C86 | -0.1665 (8) | 0.5290 (3) | 0.5781 (8) | 0.059 (2) | 0.391 (4) |
| H86A | -0.127405 | 0.512239 | 0.622100 | 0.071* | 0.391 (4) |
| H86B | -0.226620 | 0.529216 | 0.590166 | 0.071* | 0.391 (4) |
| C87 | -0.1741 (10) | 0.5205 (3) | 0.4720 (9) | 0.085 (2) | 0.391 (4) |
| H87A | -0.170803 | 0.495447 | 0.461927 | 0.102* | 0.391 (4) |
| H87B | -0.230730 | 0.529562 | 0.428331 | 0.102* | 0.391 (4) |
| C88 | -0.0918 (8) | 0.5389 (3) | 0.4575 (8) | 0.0597 (19) | 0.391 (4) |
| H88A | -0.040437 | 0.522732 | 0.468120 | 0.072* | 0.391 (4) |
| H88B | -0.105216 | 0.547718 | 0.390648 | 0.072* | 0.391 (4) |
| O7' | -0.0982 (4) | 0.57383 (12) | 0.4729 (4) | 0.0581 (11) | 0.609 (4) |
| C85' | -0.1206 (4) | 0.58116 (18) | 0.5618 (4) | 0.0498 (14) | 0.609 (4) |
| H85C | -0.173875 | 0.596458 | 0.549324 | 0.060* | 0.609 (4) |
| H85D | -0.069352 | 0.592606 | 0.608368 | 0.060* | 0.609 (4) |
| C86' | -0.1416 (5) | 0.5462 (2) | 0.6032 (5) | 0.059 (2) | 0.609 (4) |
| H86C | -0.100306 | 0.542011 | 0.668204 | 0.071* | 0.609 (4) |
| H86D | -0.204677 | 0.545470 | 0.607165 | 0.071* | 0.609 (4) |
| C87' | -0.1266 (8) | 0.51961 (19) | 0.5302 (6) | 0.085 (2) | 0.609 (4) |
| H87C | -0.066081 | 0.508980 | 0.552718 | 0.102* | 0.609 (4) |
| H87D | -0.173175 | 0.501398 | 0.518011 | 0.102* | 0.609 (4) |
| C88' | -0.1352 (6) | 0.54173 (19) | 0.4404 (5) | 0.0597 (19) | 0.609 (4) |
| H88C | -0.101999 | 0.531100 | 0.398034 | 0.072* | 0.609 (4) |
| H88D | -0.199375 | 0.544161 | 0.403442 | 0.072* | 0.609 (4) |

Atomic displacement parameters (\AA^2)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|-----|--------------|--------------|--------------|---------------|--------------|---------------|
| Co1 | 0.02125 (18) | 0.01477 (17) | 0.01464 (16) | 0.00069 (13) | 0.00911 (13) | 0.00017 (12) |
| P1 | 0.0234 (3) | 0.0164 (2) | 0.0162 (2) | -0.00055 (19) | 0.0087 (2) | -0.00049 (18) |
| P2 | 0.0241 (3) | 0.0183 (3) | 0.0165 (2) | 0.0010 (2) | 0.0104 (2) | -0.00129 (19) |
| P3 | 0.0245 (3) | 0.0140 (2) | 0.0164 (2) | 0.00024 (19) | 0.0102 (2) | 0.00049 (18) |
| P4 | 0.0234 (3) | 0.0152 (2) | 0.0155 (2) | 0.00125 (19) | 0.0091 (2) | 0.00117 (18) |
| C1 | 0.0258 (11) | 0.0219 (11) | 0.0202 (10) | 0.0008 (8) | 0.0100 (8) | 0.0019 (8) |
| C2 | 0.0251 (12) | 0.0293 (12) | 0.0284 (11) | -0.0031 (9) | 0.0101 (9) | -0.0021 (9) |
| C3 | 0.0230 (12) | 0.0400 (14) | 0.0349 (13) | -0.0034 (10) | 0.0110 (10) | -0.0006 (11) |
| C4 | 0.0272 (13) | 0.0466 (16) | 0.0372 (14) | 0.0029 (11) | 0.0179 (11) | -0.0023 (12) |
| C5 | 0.0257 (12) | 0.0358 (13) | 0.0281 (12) | 0.0011 (10) | 0.0146 (10) | -0.0041 (10) |
| C6 | 0.0279 (12) | 0.0232 (11) | 0.0204 (10) | 0.0005 (9) | 0.0119 (9) | 0.0012 (8) |
| C7 | 0.0246 (11) | 0.0194 (10) | 0.0181 (10) | 0.0025 (8) | 0.0096 (8) | 0.0005 (8) |
| C8 | 0.0249 (11) | 0.0205 (10) | 0.0214 (10) | 0.0024 (8) | 0.0100 (9) | 0.0014 (8) |
| C9 | 0.0295 (12) | 0.0256 (11) | 0.0266 (11) | 0.0038 (9) | 0.0130 (9) | 0.0066 (9) |

| | | | | | | |
|-----|-------------|-------------|-------------|--------------|-------------|--------------|
| C10 | 0.0294 (12) | 0.0338 (13) | 0.0178 (10) | 0.0068 (10) | 0.0094 (9) | 0.0004 (9) |
| C11 | 0.0278 (12) | 0.0237 (11) | 0.0227 (11) | 0.0044 (9) | 0.0069 (9) | -0.0028 (9) |
| C12 | 0.0270 (11) | 0.0201 (10) | 0.0217 (10) | 0.0028 (9) | 0.0081 (9) | 0.0010 (8) |
| C13 | 0.0290 (12) | 0.0212 (11) | 0.0166 (9) | -0.0013 (9) | 0.0063 (8) | 0.0023 (8) |
| C14 | 0.0335 (12) | 0.0193 (11) | 0.0208 (10) | -0.0002 (9) | 0.0074 (9) | 0.0008 (8) |
| C15 | 0.0421 (14) | 0.0213 (11) | 0.0246 (11) | 0.0037 (10) | 0.0023 (10) | -0.0011 (9) |
| C16 | 0.0441 (15) | 0.0173 (11) | 0.0352 (13) | -0.0042 (10) | 0.0004 (11) | 0.0071 (10) |
| C17 | 0.0374 (14) | 0.0322 (13) | 0.0319 (13) | -0.0078 (11) | 0.0057 (11) | 0.0132 (11) |
| C18 | 0.0326 (13) | 0.0266 (12) | 0.0251 (11) | -0.0028 (10) | 0.0099 (10) | 0.0059 (9) |
| C19 | 0.0223 (11) | 0.0269 (11) | 0.0213 (10) | -0.0016 (9) | 0.0101 (8) | -0.0001 (9) |
| C20 | 0.0381 (14) | 0.0301 (12) | 0.0215 (11) | -0.0004 (10) | 0.0175 (10) | -0.0015 (9) |
| C21 | 0.0447 (15) | 0.0408 (15) | 0.0233 (12) | -0.0049 (12) | 0.0178 (11) | -0.0050 (10) |
| C22 | 0.0359 (14) | 0.0452 (15) | 0.0210 (11) | -0.0046 (11) | 0.0131 (10) | 0.0061 (10) |
| C23 | 0.0299 (13) | 0.0325 (13) | 0.0269 (12) | -0.0017 (10) | 0.0113 (10) | 0.0070 (10) |
| C24 | 0.0252 (11) | 0.0275 (12) | 0.0227 (11) | -0.0021 (9) | 0.0126 (9) | 0.0003 (9) |
| C25 | 0.0305 (12) | 0.0212 (11) | 0.0198 (10) | 0.0015 (9) | 0.0144 (9) | -0.0011 (8) |
| C26 | 0.0336 (13) | 0.0265 (12) | 0.0323 (12) | 0.0050 (10) | 0.0110 (10) | -0.0008 (10) |
| C27 | 0.0443 (16) | 0.0272 (13) | 0.0470 (15) | 0.0134 (11) | 0.0159 (13) | 0.0037 (11) |
| C28 | 0.0541 (17) | 0.0200 (12) | 0.0433 (15) | 0.0052 (11) | 0.0251 (13) | -0.0024 (10) |
| C29 | 0.0461 (15) | 0.0240 (12) | 0.0292 (12) | -0.0046 (10) | 0.0166 (11) | -0.0058 (10) |
| C30 | 0.0377 (14) | 0.0226 (11) | 0.0237 (11) | 0.0007 (10) | 0.0117 (10) | -0.0012 (9) |
| C31 | 0.0289 (11) | 0.0172 (10) | 0.0173 (9) | 0.0004 (8) | 0.0115 (8) | -0.0022 (8) |
| C32 | 0.0253 (11) | 0.0219 (11) | 0.0258 (11) | -0.0009 (9) | 0.0132 (9) | 0.0004 (9) |
| C33 | 0.0271 (12) | 0.0269 (12) | 0.0320 (12) | -0.0045 (9) | 0.0155 (10) | -0.0031 (9) |
| C34 | 0.0238 (12) | 0.0276 (12) | 0.0289 (12) | 0.0032 (9) | 0.0082 (9) | -0.0016 (9) |
| C35 | 0.0252 (11) | 0.0221 (11) | 0.0229 (10) | 0.0018 (9) | 0.0090 (9) | -0.0005 (8) |
| C36 | 0.0277 (11) | 0.0184 (10) | 0.0167 (9) | -0.0004 (8) | 0.0107 (8) | -0.0023 (8) |
| C37 | 0.0290 (11) | 0.0189 (10) | 0.0167 (9) | 0.0006 (8) | 0.0129 (8) | -0.0012 (8) |
| C38 | 0.0328 (13) | 0.0219 (11) | 0.0225 (10) | 0.0034 (9) | 0.0084 (9) | -0.0031 (9) |
| C39 | 0.0413 (14) | 0.0242 (12) | 0.0225 (11) | 0.0094 (10) | 0.0107 (10) | 0.0002 (9) |
| C40 | 0.0489 (16) | 0.0163 (11) | 0.0331 (13) | 0.0034 (10) | 0.0185 (11) | -0.0026 (9) |
| C41 | 0.0405 (15) | 0.0237 (13) | 0.0464 (15) | -0.0011 (11) | 0.0099 (12) | -0.0118 (11) |
| C42 | 0.0354 (14) | 0.0219 (12) | 0.0319 (12) | 0.0011 (10) | 0.0038 (10) | -0.0070 (10) |
| C43 | 0.0248 (11) | 0.0163 (10) | 0.0212 (10) | -0.0001 (8) | 0.0109 (8) | -0.0001 (8) |
| C44 | 0.0359 (13) | 0.0195 (10) | 0.0209 (10) | -0.0001 (9) | 0.0162 (9) | 0.0014 (8) |
| C45 | 0.0427 (14) | 0.0253 (12) | 0.0245 (11) | 0.0028 (10) | 0.0164 (10) | 0.0050 (9) |
| C46 | 0.0342 (13) | 0.0316 (13) | 0.0189 (10) | 0.0031 (10) | 0.0136 (9) | -0.0015 (9) |
| C47 | 0.0305 (12) | 0.0212 (11) | 0.0256 (11) | 0.0013 (9) | 0.0123 (9) | -0.0061 (9) |
| C48 | 0.0302 (12) | 0.0172 (10) | 0.0216 (10) | 0.0009 (8) | 0.0113 (9) | 0.0012 (8) |
| C49 | 0.0270 (11) | 0.0177 (10) | 0.0176 (10) | 0.0006 (8) | 0.0064 (8) | 0.0003 (8) |
| C50 | 0.0310 (12) | 0.0195 (10) | 0.0221 (10) | -0.0001 (9) | 0.0144 (9) | -0.0001 (8) |
| C51 | 0.0356 (13) | 0.0229 (11) | 0.0296 (12) | 0.0013 (10) | 0.0166 (10) | -0.0036 (9) |
| C52 | 0.0355 (13) | 0.0161 (10) | 0.0295 (12) | 0.0021 (9) | 0.0096 (10) | -0.0024 (9) |
| C53 | 0.0313 (12) | 0.0178 (11) | 0.0269 (11) | -0.0012 (9) | 0.0077 (9) | 0.0044 (9) |
| C54 | 0.0256 (11) | 0.0203 (11) | 0.0213 (10) | 0.0004 (8) | 0.0091 (8) | 0.0015 (8) |
| C55 | 0.0238 (11) | 0.0231 (11) | 0.0180 (10) | -0.0003 (8) | 0.0103 (8) | 0.0004 (8) |
| C56 | 0.0332 (12) | 0.0233 (11) | 0.0208 (10) | 0.0035 (9) | 0.0100 (9) | 0.0015 (8) |
| C57 | 0.0360 (13) | 0.0271 (12) | 0.0232 (11) | 0.0002 (10) | 0.0083 (10) | 0.0049 (9) |

| | | | | | | |
|------|--------------|--------------|-------------|---------------|--------------|--------------|
| C58 | 0.0341 (13) | 0.0364 (13) | 0.0155 (10) | -0.0030 (10) | 0.0092 (9) | 0.0014 (9) |
| C59 | 0.0279 (12) | 0.0273 (12) | 0.0238 (11) | -0.0022 (9) | 0.0118 (9) | -0.0031 (9) |
| C60 | 0.0244 (11) | 0.0215 (11) | 0.0215 (10) | 0.0002 (8) | 0.0094 (9) | 0.0015 (8) |
| Br1 | 0.04265 (19) | 0.02299 (15) | 0.1025 (3) | -0.00182 (12) | 0.00695 (18) | 0.01037 (16) |
| Mg1 | 0.0286 (4) | 0.0227 (4) | 0.0349 (4) | -0.0027 (3) | 0.0092 (3) | -0.0004 (3) |
| O1 | 0.0312 (9) | 0.0215 (8) | 0.0269 (8) | -0.0027 (7) | 0.0102 (7) | -0.0027 (6) |
| O2 | 0.0312 (18) | 0.0277 (11) | 0.039 (3) | -0.0074 (14) | 0.0084 (18) | -0.005 (2) |
| C65 | 0.0319 (17) | 0.028 (2) | 0.070 (6) | -0.0095 (15) | 0.013 (4) | 0.000 (4) |
| C66 | 0.055 (3) | 0.030 (4) | 0.086 (7) | -0.014 (3) | -0.001 (4) | 0.002 (4) |
| C67 | 0.063 (4) | 0.023 (2) | 0.049 (2) | 0.011 (2) | -0.022 (3) | -0.0091 (18) |
| C68 | 0.047 (5) | 0.0291 (19) | 0.037 (3) | 0.004 (2) | -0.003 (2) | -0.0057 (17) |
| O2' | 0.0312 (18) | 0.0277 (11) | 0.039 (3) | -0.0074 (14) | 0.0084 (18) | -0.005 (2) |
| C65' | 0.0319 (17) | 0.028 (2) | 0.070 (6) | -0.0095 (15) | 0.013 (4) | 0.000 (4) |
| C66' | 0.055 (3) | 0.030 (4) | 0.086 (7) | -0.014 (3) | -0.001 (4) | 0.002 (4) |
| C67' | 0.063 (4) | 0.023 (2) | 0.049 (2) | 0.011 (2) | -0.022 (3) | -0.0091 (18) |
| C68' | 0.047 (5) | 0.0291 (19) | 0.037 (3) | 0.004 (2) | -0.003 (2) | -0.0057 (17) |
| O3 | 0.034 (5) | 0.035 (2) | 0.030 (3) | 0.004 (4) | 0.006 (4) | -0.013 (3) |
| C69 | 0.0457 (19) | 0.060 (2) | 0.047 (5) | -0.0039 (16) | 0.027 (3) | -0.003 (3) |
| C70 | 0.065 (4) | 0.120 (5) | 0.051 (4) | 0.035 (4) | 0.039 (3) | 0.015 (4) |
| C71 | 0.113 (9) | 0.124 (6) | 0.044 (5) | 0.071 (5) | 0.028 (5) | -0.007 (4) |
| C72 | 0.088 (7) | 0.060 (2) | 0.051 (3) | 0.026 (3) | 0.007 (5) | -0.032 (2) |
| O3' | 0.034 (5) | 0.035 (2) | 0.030 (3) | 0.004 (4) | 0.006 (4) | -0.013 (3) |
| C69' | 0.0457 (19) | 0.060 (2) | 0.047 (5) | -0.0039 (16) | 0.027 (3) | -0.003 (3) |
| C70' | 0.065 (4) | 0.120 (5) | 0.051 (4) | 0.035 (4) | 0.039 (3) | 0.015 (4) |
| C71' | 0.113 (9) | 0.124 (6) | 0.044 (5) | 0.071 (5) | 0.028 (5) | -0.007 (4) |
| C72' | 0.088 (7) | 0.060 (2) | 0.051 (3) | 0.026 (3) | 0.007 (5) | -0.032 (2) |
| O4 | 0.0291 (9) | 0.0343 (10) | 0.0292 (9) | 0.0012 (7) | 0.0082 (7) | -0.0039 (7) |
| O5 | 0.0358 (10) | 0.0463 (11) | 0.0411 (10) | 0.0019 (8) | 0.0197 (8) | 0.0146 (9) |
| C61 | 0.0407 (15) | 0.0279 (13) | 0.0362 (13) | -0.0042 (11) | 0.0068 (11) | -0.0085 (11) |
| C62 | 0.0438 (16) | 0.0248 (13) | 0.0461 (15) | 0.0012 (11) | 0.0096 (12) | -0.0080 (11) |
| C63 | 0.0471 (16) | 0.0230 (12) | 0.0421 (15) | -0.0025 (11) | 0.0107 (12) | 0.0026 (11) |
| C64 | 0.0365 (14) | 0.0242 (12) | 0.0300 (12) | -0.0012 (10) | 0.0069 (10) | 0.0006 (9) |
| C73 | 0.0326 (14) | 0.0311 (14) | 0.0503 (16) | 0.0035 (11) | 0.0079 (12) | -0.0040 (12) |
| C74 | 0.080 (3) | 0.0407 (18) | 0.055 (2) | 0.0168 (18) | -0.0041 (18) | -0.0001 (15) |
| C75 | 0.0474 (17) | 0.0438 (16) | 0.0341 (14) | -0.0034 (13) | 0.0050 (12) | 0.0079 (12) |
| C76 | 0.0408 (15) | 0.0336 (14) | 0.0273 (12) | -0.0023 (11) | 0.0088 (11) | -0.0018 (10) |
| C77 | 0.051 (4) | 0.064 (3) | 0.056 (2) | -0.006 (2) | 0.023 (2) | 0.027 (2) |
| C78 | 0.051 (6) | 0.109 (5) | 0.039 (4) | 0.014 (4) | 0.022 (4) | 0.022 (3) |
| C77' | 0.051 (4) | 0.064 (3) | 0.056 (2) | -0.006 (2) | 0.023 (2) | 0.027 (2) |
| C78' | 0.051 (6) | 0.109 (5) | 0.039 (4) | 0.014 (4) | 0.022 (4) | 0.022 (3) |
| C79 | 0.0411 (18) | 0.086 (3) | 0.0442 (18) | 0.0067 (17) | 0.0147 (14) | -0.0014 (17) |
| C80 | 0.0328 (15) | 0.060 (2) | 0.0445 (16) | 0.0087 (13) | 0.0185 (12) | 0.0094 (14) |
| O6 | 0.0610 (14) | 0.0383 (11) | 0.0480 (12) | -0.0062 (10) | 0.0235 (10) | -0.0034 (9) |
| C81 | 0.056 (2) | 0.063 (2) | 0.0456 (17) | -0.0177 (17) | 0.0202 (15) | -0.0021 (16) |
| C82 | 0.0487 (18) | 0.0391 (16) | 0.0546 (19) | 0.0049 (14) | 0.0058 (15) | -0.0066 (14) |
| C83 | 0.0475 (19) | 0.0470 (19) | 0.096 (3) | -0.0077 (15) | 0.0401 (19) | -0.0279 (19) |
| C84 | 0.0416 (18) | 0.061 (2) | 0.067 (2) | -0.0089 (15) | 0.0075 (16) | 0.0274 (18) |
| O7 | 0.071 (3) | 0.059 (2) | 0.058 (3) | -0.005 (2) | 0.040 (3) | 0.004 (2) |

| | | | | | | |
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| C85 | 0.043 (2) | 0.065 (5) | 0.042 (3) | 0.007 (3) | 0.013 (2) | -0.003 (3) |
| C86 | 0.044 (4) | 0.086 (7) | 0.051 (4) | 0.006 (3) | 0.018 (3) | 0.027 (4) |
| C87 | 0.139 (8) | 0.056 (3) | 0.075 (5) | 0.017 (4) | 0.054 (5) | 0.025 (4) |
| C88 | 0.083 (6) | 0.059 (3) | 0.040 (3) | 0.027 (4) | 0.021 (4) | 0.009 (2) |
| O7' | 0.071 (3) | 0.059 (2) | 0.058 (3) | -0.005 (2) | 0.040 (3) | 0.004 (2) |
| C85' | 0.043 (2) | 0.065 (5) | 0.042 (3) | 0.007 (3) | 0.013 (2) | -0.003 (3) |
| C86' | 0.044 (4) | 0.086 (7) | 0.051 (4) | 0.006 (3) | 0.018 (3) | 0.027 (4) |
| C87' | 0.139 (8) | 0.056 (3) | 0.075 (5) | 0.017 (4) | 0.054 (5) | 0.025 (4) |
| C88' | 0.083 (6) | 0.059 (3) | 0.040 (3) | 0.027 (4) | 0.021 (4) | 0.009 (2) |

Geometric parameters (Å, °)

| | | | |
|---------|------------|-----------|------------|
| Co1—P1 | 2.1049 (6) | C65—H65A | 0.9700 |
| Co1—P2 | 2.0988 (6) | C65—H65B | 0.9700 |
| Co1—P3 | 2.0968 (6) | C65—C66 | 1.516 (8) |
| Co1—P4 | 2.1050 (6) | C66—H66A | 0.9700 |
| P1—C1 | 1.870 (2) | C66—H66B | 0.9700 |
| P1—C7 | 1.876 (2) | C66—C67 | 1.532 (10) |
| P1—C13 | 1.849 (2) | C67—H67A | 0.9700 |
| P2—C6 | 1.854 (2) | C67—H67B | 0.9700 |
| P2—C19 | 1.867 (2) | C67—C68 | 1.517 (7) |
| P2—C25 | 1.847 (2) | C68—H68A | 0.9700 |
| P3—C31 | 1.859 (2) | C68—H68B | 0.9700 |
| P3—C37 | 1.846 (2) | O2'—C65' | 1.456 (13) |
| P3—C43 | 1.870 (2) | O2'—C68' | 1.441 (12) |
| P4—C36 | 1.863 (2) | C65'—H65C | 0.9700 |
| P4—C49 | 1.843 (2) | C65'—H65D | 0.9700 |
| P4—C55 | 1.874 (2) | C65'—C66' | 1.523 (14) |
| C1—C2 | 1.399 (3) | C66'—H66C | 0.9700 |
| C1—C6 | 1.396 (3) | C66'—H66D | 0.9700 |
| C2—H2 | 0.9300 | C66'—C67' | 1.518 (14) |
| C2—C3 | 1.386 (4) | C67'—H67C | 0.9700 |
| C3—H3 | 0.9300 | C67'—H67D | 0.9700 |
| C3—C4 | 1.391 (4) | C67'—C68' | 1.512 (12) |
| C4—H4 | 0.9300 | C68'—H68C | 0.9700 |
| C4—C5 | 1.380 (4) | C68'—H68D | 0.9700 |
| C5—H5 | 0.9300 | O3—C69 | 1.458 (10) |
| C5—C6 | 1.402 (3) | O3—C72 | 1.466 (8) |
| C7—C8 | 1.397 (3) | C69—H69A | 0.9700 |
| C7—C12 | 1.396 (3) | C69—H69B | 0.9700 |
| C8—H8 | 0.9300 | C69—C70 | 1.500 (14) |
| C8—C9 | 1.392 (3) | C70—H70A | 0.9700 |
| C9—H9 | 0.9300 | C70—H70B | 0.9700 |
| C9—C10 | 1.388 (4) | C70—C71 | 1.494 (11) |
| C10—H10 | 0.9300 | C71—H71A | 0.9700 |
| C10—C11 | 1.391 (3) | C71—H71B | 0.9700 |
| C11—H11 | 0.9300 | C71—C72 | 1.491 (13) |
| C11—C12 | 1.393 (3) | C72—H72A | 0.9700 |

| | | | |
|---------|-----------|-----------|------------|
| C12—H12 | 0.9300 | C72—H72B | 0.9700 |
| C13—C14 | 1.396 (3) | O3'—C69' | 1.458 (13) |
| C13—C18 | 1.398 (3) | O3'—C72' | 1.465 (13) |
| C14—H14 | 0.9300 | C69'—H69C | 0.9700 |
| C14—C15 | 1.388 (3) | C69'—H69D | 0.9700 |
| C15—H15 | 0.9300 | C69'—C70' | 1.502 (17) |
| C15—C16 | 1.390 (4) | C70'—H70C | 0.9700 |
| C16—H16 | 0.9300 | C70'—H70D | 0.9700 |
| C16—C17 | 1.378 (4) | C70'—C71' | 1.500 (16) |
| C17—H17 | 0.9300 | C71'—H71C | 0.9700 |
| C17—C18 | 1.391 (4) | C71'—H71D | 0.9700 |
| C18—H18 | 0.9300 | C71'—C72' | 1.500 (17) |
| C19—C20 | 1.396 (3) | C72'—H72C | 0.9700 |
| C19—C24 | 1.397 (3) | C72'—H72D | 0.9700 |
| C20—H20 | 0.9300 | O4—C73 | 1.457 (3) |
| C20—C21 | 1.395 (3) | O4—C76 | 1.450 (3) |
| C21—H21 | 0.9300 | O5—C77 | 1.449 (5) |
| C21—C22 | 1.386 (4) | O5—C77' | 1.450 (8) |
| C22—H22 | 0.9300 | O5—C80 | 1.450 (4) |
| C22—C23 | 1.386 (4) | C61—H61A | 0.9700 |
| C23—H23 | 0.9300 | C61—H61B | 0.9700 |
| C23—C24 | 1.390 (3) | C61—C62 | 1.510 (4) |
| C24—H24 | 0.9300 | C62—H62A | 0.9700 |
| C25—C26 | 1.394 (3) | C62—H62B | 0.9700 |
| C25—C30 | 1.396 (3) | C62—C63 | 1.511 (4) |
| C26—H26 | 0.9300 | C63—H63A | 0.9700 |
| C26—C27 | 1.389 (4) | C63—H63B | 0.9700 |
| C27—H27 | 0.9300 | C63—C64 | 1.512 (3) |
| C27—C28 | 1.377 (4) | C64—H64A | 0.9700 |
| C28—H28 | 0.9300 | C64—H64B | 0.9700 |
| C28—C29 | 1.376 (4) | C73—H73A | 0.9700 |
| C29—H29 | 0.9300 | C73—H73B | 0.9700 |
| C29—C30 | 1.389 (3) | C73—C74 | 1.495 (5) |
| C30—H30 | 0.9300 | C74—H74A | 0.9700 |
| C31—C32 | 1.396 (3) | C74—H74B | 0.9700 |
| C31—C36 | 1.397 (3) | C74—C75 | 1.515 (5) |
| C32—H32 | 0.9300 | C75—H75A | 0.9700 |
| C32—C33 | 1.387 (3) | C75—H75B | 0.9700 |
| C33—H33 | 0.9300 | C75—C76 | 1.518 (4) |
| C33—C34 | 1.397 (3) | C76—H76A | 0.9700 |
| C34—H34 | 0.9300 | C76—H76B | 0.9700 |
| C34—C35 | 1.386 (3) | C77—H77A | 0.9700 |
| C35—H35 | 0.9300 | C77—H77B | 0.9700 |
| C35—C36 | 1.398 (3) | C77—C78 | 1.533 (8) |
| C37—C38 | 1.394 (3) | C78—H78A | 0.9700 |
| C37—C42 | 1.393 (3) | C78—H78B | 0.9700 |
| C38—H38 | 0.9300 | C78—C79 | 1.525 (6) |
| C38—C39 | 1.386 (3) | C77'—H77C | 0.9700 |

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| C39—H39 | 0.9300 | C77'—H77D | 0.9700 |
| C39—C40 | 1.384 (4) | C77'—C78' | 1.531 (9) |
| C40—H40 | 0.9300 | C78'—H78C | 0.9700 |
| C40—C41 | 1.384 (4) | C78'—H78D | 0.9700 |
| C41—H41 | 0.9300 | C78'—C79 | 1.539 (8) |
| C41—C42 | 1.387 (3) | C79—H79A | 0.9700 |
| C42—H42 | 0.9300 | C79—H79B | 0.9700 |
| C43—C44 | 1.391 (3) | C79—H79C | 0.9700 |
| C43—C48 | 1.402 (3) | C79—H79D | 0.9700 |
| C44—H44 | 0.9300 | C79—C80 | 1.504 (5) |
| C44—C45 | 1.391 (3) | C80—H80A | 0.9700 |
| C45—H45 | 0.9300 | C80—H80B | 0.9700 |
| C45—C46 | 1.384 (3) | O6—C81 | 1.423 (4) |
| C46—H46 | 0.9300 | O6—C84 | 1.423 (4) |
| C46—C47 | 1.388 (3) | C81—H81A | 0.9700 |
| C47—H47 | 0.9300 | C81—H81B | 0.9700 |
| C47—C48 | 1.383 (3) | C81—C82 | 1.500 (5) |
| C48—H48 | 0.9300 | C82—H82A | 0.9700 |
| C49—C50 | 1.399 (3) | C82—H82B | 0.9700 |
| C49—C54 | 1.404 (3) | C82—C83 | 1.511 (5) |
| C50—H50 | 0.9300 | C83—H83A | 0.9700 |
| C50—C51 | 1.387 (3) | C83—H83B | 0.9700 |
| C51—H51 | 0.9300 | C83—C84 | 1.510 (6) |
| C51—C52 | 1.386 (3) | C84—H84A | 0.9700 |
| C52—H52 | 0.9300 | C84—H84B | 0.9700 |
| C52—C53 | 1.386 (3) | O7—C85 | 1.422 (10) |
| C53—H53 | 0.9300 | O7—C88 | 1.409 (11) |
| C53—C54 | 1.388 (3) | C85—H85A | 0.9700 |
| C54—H54 | 0.9300 | C85—H85B | 0.9700 |
| C55—C56 | 1.393 (3) | C85—C86 | 1.508 (12) |
| C55—C60 | 1.399 (3) | C86—H86A | 0.9700 |
| C56—H56 | 0.9300 | C86—H86B | 0.9700 |
| C56—C57 | 1.391 (3) | C86—C87 | 1.506 (13) |
| C57—H57 | 0.9300 | C87—H87A | 0.9700 |
| C57—C58 | 1.383 (4) | C87—H87B | 0.9700 |
| C58—H58 | 0.9300 | C87—C88 | 1.492 (13) |
| C58—C59 | 1.390 (3) | C88—H88A | 0.9700 |
| C59—H59 | 0.9300 | C88—H88B | 0.9700 |
| C59—C60 | 1.383 (3) | O7'—C85' | 1.417 (7) |
| C60—H60 | 0.9300 | O7'—C88' | 1.372 (8) |
| Br1—Mg1 | 2.5990 (9) | C85'—H85C | 0.9700 |
| Mg1—O1 | 2.1166 (18) | C85'—H85D | 0.9700 |
| Mg1—O2 | 2.128 (6) | C85'—C86' | 1.527 (8) |
| Mg1—O2' | 2.129 (12) | C86'—H86C | 0.9700 |
| Mg1—O3 | 2.132 (7) | C86'—H86D | 0.9700 |
| Mg1—O3' | 2.114 (11) | C86'—C87' | 1.507 (11) |
| Mg1—O4 | 2.1177 (19) | C87'—H87C | 0.9700 |
| Mg1—O5 | 2.126 (2) | C87'—H87D | 0.9700 |

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| O1—C61 | 1.445 (3) | C87'—C88' | 1.500 (8) |
| O1—C64 | 1.460 (3) | C88'—H88C | 0.9700 |
| O2—C65 | 1.464 (7) | C88'—H88D | 0.9700 |
| O2—C68 | 1.450 (7) | | |
| P1—Co1—P4 | 120.60 (3) | O2—C68—H68A | 111.3 |
| P2—Co1—P1 | 90.18 (2) | O2—C68—H68B | 111.3 |
| P2—Co1—P4 | 118.59 (2) | C67—C68—H68A | 111.3 |
| P3—Co1—P1 | 122.08 (3) | C67—C68—H68B | 111.3 |
| P3—Co1—P2 | 118.74 (3) | H68A—C68—H68B | 109.2 |
| P3—Co1—P4 | 89.75 (2) | C65'—O2'—Mg1 | 120 (2) |
| C1—P1—Co1 | 109.40 (8) | C68'—O2'—Mg1 | 128.5 (14) |
| C1—P1—C7 | 98.00 (10) | C68'—O2'—C65' | 106.5 (12) |
| C7—P1—Co1 | 126.18 (7) | O2'—C65'—H65C | 110.9 |
| C13—P1—Co1 | 121.21 (8) | O2'—C65'—H65D | 110.9 |
| C13—P1—C1 | 100.06 (10) | O2'—C65'—C66' | 104.3 (14) |
| C13—P1—C7 | 97.06 (10) | H65C—C65'—H65D | 108.9 |
| C6—P2—Co1 | 109.95 (7) | C66'—C65'—H65C | 110.9 |
| C6—P2—C19 | 97.01 (10) | C66'—C65'—H65D | 110.9 |
| C19—P2—Co1 | 121.38 (8) | C65'—C66'—H66C | 111.4 |
| C25—P2—Co1 | 122.03 (7) | C65'—C66'—H66D | 111.4 |
| C25—P2—C6 | 101.73 (11) | H66C—C66'—H66D | 109.2 |
| C25—P2—C19 | 100.47 (10) | C67'—C66'—C65' | 102.0 (11) |
| C31—P3—Co1 | 110.35 (7) | C67'—C66'—H66C | 111.4 |
| C31—P3—C43 | 95.75 (10) | C67'—C66'—H66D | 111.4 |
| C37—P3—Co1 | 121.35 (7) | C66'—C67'—H67C | 110.2 |
| C37—P3—C31 | 103.36 (10) | C66'—C67'—H67D | 110.2 |
| C37—P3—C43 | 98.79 (9) | H67C—C67'—H67D | 108.5 |
| C43—P3—Co1 | 122.95 (7) | C68'—C67'—C66' | 107.3 (9) |
| C36—P4—Co1 | 109.46 (7) | C68'—C67'—H67C | 110.2 |
| C36—P4—C55 | 96.99 (10) | C68'—C67'—H67D | 110.2 |
| C49—P4—Co1 | 121.52 (7) | O2'—C68'—C67' | 105.3 (11) |
| C49—P4—C36 | 100.76 (10) | O2'—C68'—H68C | 110.7 |
| C49—P4—C55 | 99.23 (10) | O2'—C68'—H68D | 110.7 |
| C55—P4—Co1 | 124.14 (7) | C67'—C68'—H68C | 110.7 |
| C2—C1—P1 | 125.93 (18) | C67'—C68'—H68D | 110.7 |
| C6—C1—P1 | 114.93 (17) | H68C—C68'—H68D | 108.8 |
| C6—C1—C2 | 119.1 (2) | C69—O3—Mg1 | 124.7 (11) |
| C1—C2—H2 | 119.7 | C69—O3—C72 | 107.3 (7) |
| C3—C2—C1 | 120.6 (2) | C72—O3—Mg1 | 128.1 (13) |
| C3—C2—H2 | 119.7 | O3—C69—H69A | 110.5 |
| C2—C3—H3 | 120.0 | O3—C69—H69B | 110.5 |
| C2—C3—C4 | 120.0 (2) | O3—C69—C70 | 106.3 (8) |
| C4—C3—H3 | 120.0 | H69A—C69—H69B | 108.7 |
| C3—C4—H4 | 120.0 | C70—C69—H69A | 110.5 |
| C5—C4—C3 | 120.0 (2) | C70—C69—H69B | 110.5 |
| C5—C4—H4 | 120.0 | C69—C70—H70A | 111.2 |
| C4—C5—H5 | 119.8 | C69—C70—H70B | 111.2 |

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| C4—C5—C6 | 120.4 (2) | H70A—C70—H70B | 109.1 |
| C6—C5—H5 | 119.8 | C71—C70—C69 | 102.9 (10) |
| C1—C6—P2 | 115.32 (17) | C71—C70—H70A | 111.2 |
| C1—C6—C5 | 119.8 (2) | C71—C70—H70B | 111.2 |
| C5—C6—P2 | 124.85 (18) | C70—C71—H71A | 111.1 |
| C8—C7—P1 | 117.54 (17) | C70—C71—H71B | 111.1 |
| C12—C7—P1 | 124.33 (16) | H71A—C71—H71B | 109.0 |
| C12—C7—C8 | 118.1 (2) | C72—C71—C70 | 103.5 (12) |
| C7—C8—H8 | 119.6 | C72—C71—H71A | 111.1 |
| C9—C8—C7 | 120.7 (2) | C72—C71—H71B | 111.1 |
| C9—C8—H8 | 119.6 | O3—C72—C71 | 106.9 (8) |
| C8—C9—H9 | 119.7 | O3—C72—H72A | 110.3 |
| C10—C9—C8 | 120.7 (2) | O3—C72—H72B | 110.3 |
| C10—C9—H9 | 119.7 | C71—C72—H72A | 110.3 |
| C9—C10—H10 | 120.4 | C71—C72—H72B | 110.3 |
| C9—C10—C11 | 119.2 (2) | H72A—C72—H72B | 108.6 |
| C11—C10—H10 | 120.4 | C69'—O3'—Mg1 | 119.6 (19) |
| C10—C11—H11 | 119.9 | C69'—O3'—C72' | 108.5 (13) |
| C10—C11—C12 | 120.1 (2) | C72'—O3'—Mg1 | 126 (2) |
| C12—C11—H11 | 119.9 | O3'—C69'—H69C | 110.7 |
| C7—C12—H12 | 119.4 | O3'—C69'—H69D | 110.7 |
| C11—C12—C7 | 121.2 (2) | O3'—C69'—C70' | 105.3 (11) |
| C11—C12—H12 | 119.4 | H69C—C69'—H69D | 108.8 |
| C14—C13—P1 | 118.76 (17) | C70'—C69'—H69C | 110.7 |
| C14—C13—C18 | 118.1 (2) | C70'—C69'—H69D | 110.7 |
| C18—C13—P1 | 123.06 (18) | C69'—C70'—H70C | 111.4 |
| C13—C14—H14 | 119.5 | C69'—C70'—H70D | 111.4 |
| C15—C14—C13 | 121.1 (2) | H70C—C70'—H70D | 109.3 |
| C15—C14—H14 | 119.5 | C71'—C70'—C69' | 101.9 (15) |
| C14—C15—H15 | 119.9 | C71'—C70'—H70C | 111.4 |
| C14—C15—C16 | 120.2 (2) | C71'—C70'—H70D | 111.4 |
| C16—C15—H15 | 119.9 | C70'—C71'—H71C | 111.5 |
| C15—C16—H16 | 120.4 | C70'—C71'—H71D | 111.5 |
| C17—C16—C15 | 119.1 (2) | C70'—C71'—C72' | 101 (2) |
| C17—C16—H16 | 120.4 | H71C—C71'—H71D | 109.3 |
| C16—C17—H17 | 119.5 | C72'—C71'—H71C | 111.5 |
| C16—C17—C18 | 121.0 (2) | C72'—C71'—H71D | 111.5 |
| C18—C17—H17 | 119.5 | O3'—C72'—C71' | 102.3 (14) |
| C13—C18—H18 | 119.8 | O3'—C72'—H72C | 111.3 |
| C17—C18—C13 | 120.4 (2) | O3'—C72'—H72D | 111.3 |
| C17—C18—H18 | 119.8 | C71'—C72'—H72C | 111.3 |
| C20—C19—P2 | 125.82 (18) | C71'—C72'—H72D | 111.3 |
| C20—C19—C24 | 118.3 (2) | H72C—C72'—H72D | 109.2 |
| C24—C19—P2 | 115.87 (16) | C73—O4—Mg1 | 119.83 (16) |
| C19—C20—H20 | 119.8 | C76—O4—Mg1 | 132.79 (16) |
| C21—C20—C19 | 120.5 (2) | C76—O4—C73 | 106.4 (2) |
| C21—C20—H20 | 119.8 | C77—O5—Mg1 | 124.1 (6) |
| C20—C21—H21 | 119.7 | C77—O5—C80 | 105.3 (7) |

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| C22—C21—C20 | 120.5 (2) | C77'—O5—Mg1 | 123.2 (9) |
| C22—C21—H21 | 119.7 | C77'—O5—C80 | 106.9 (10) |
| C21—C22—H22 | 120.3 | C80—O5—Mg1 | 129.57 (16) |
| C23—C22—C21 | 119.5 (2) | O1—C61—H61A | 110.6 |
| C23—C22—H22 | 120.3 | O1—C61—H61B | 110.6 |
| C22—C23—H23 | 119.9 | O1—C61—C62 | 105.8 (2) |
| C22—C23—C24 | 120.2 (2) | H61A—C61—H61B | 108.7 |
| C24—C23—H23 | 119.9 | C62—C61—H61A | 110.6 |
| C19—C24—H24 | 119.5 | C62—C61—H61B | 110.6 |
| C23—C24—C19 | 121.0 (2) | C61—C62—H62A | 111.3 |
| C23—C24—H24 | 119.5 | C61—C62—H62B | 111.3 |
| C26—C25—P2 | 124.51 (19) | C61—C62—C63 | 102.3 (2) |
| C26—C25—C30 | 117.5 (2) | H62A—C62—H62B | 109.2 |
| C30—C25—P2 | 117.65 (18) | C63—C62—H62A | 111.3 |
| C25—C26—H26 | 119.6 | C63—C62—H62B | 111.3 |
| C27—C26—C25 | 120.9 (3) | C62—C63—H63A | 111.4 |
| C27—C26—H26 | 119.6 | C62—C63—H63B | 111.4 |
| C26—C27—H27 | 119.6 | C62—C63—C64 | 102.0 (2) |
| C28—C27—C26 | 120.8 (3) | H63A—C63—H63B | 109.2 |
| C28—C27—H27 | 119.6 | C64—C63—H63A | 111.4 |
| C27—C28—H28 | 120.4 | C64—C63—H63B | 111.4 |
| C29—C28—C27 | 119.2 (2) | O1—C64—C63 | 105.8 (2) |
| C29—C28—H28 | 120.4 | O1—C64—H64A | 110.6 |
| C28—C29—H29 | 119.8 | O1—C64—H64B | 110.6 |
| C28—C29—C30 | 120.4 (3) | C63—C64—H64A | 110.6 |
| C30—C29—H29 | 119.8 | C63—C64—H64B | 110.6 |
| C25—C30—H30 | 119.4 | H64A—C64—H64B | 108.7 |
| C29—C30—C25 | 121.2 (2) | O4—C73—H73A | 110.3 |
| C29—C30—H30 | 119.4 | O4—C73—H73B | 110.3 |
| C32—C31—P3 | 125.87 (17) | O4—C73—C74 | 107.1 (3) |
| C32—C31—C36 | 119.7 (2) | H73A—C73—H73B | 108.5 |
| C36—C31—P3 | 114.28 (17) | C74—C73—H73A | 110.3 |
| C31—C32—H32 | 119.6 | C74—C73—H73B | 110.3 |
| C33—C32—C31 | 120.8 (2) | C73—C74—H74A | 110.5 |
| C33—C32—H32 | 119.6 | C73—C74—H74B | 110.5 |
| C32—C33—H33 | 120.2 | C73—C74—C75 | 106.2 (3) |
| C32—C33—C34 | 119.5 (2) | H74A—C74—H74B | 108.7 |
| C34—C33—H33 | 120.2 | C75—C74—H74A | 110.5 |
| C33—C34—H34 | 120.1 | C75—C74—H74B | 110.5 |
| C35—C34—C33 | 119.8 (2) | C74—C75—H75A | 111.1 |
| C35—C34—H34 | 120.1 | C74—C75—H75B | 111.1 |
| C34—C35—H35 | 119.5 | C74—C75—C76 | 103.5 (2) |
| C34—C35—C36 | 121.0 (2) | H75A—C75—H75B | 109.0 |
| C36—C35—H35 | 119.5 | C76—C75—H75A | 111.1 |
| C31—C36—P4 | 115.53 (17) | C76—C75—H75B | 111.1 |
| C31—C36—C35 | 119.1 (2) | O4—C76—C75 | 104.7 (2) |
| C35—C36—P4 | 125.39 (17) | O4—C76—H76A | 110.8 |
| C38—C37—P3 | 117.26 (17) | O4—C76—H76B | 110.8 |

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| C42—C37—P3 | 125.07 (18) | C75—C76—H76A | 110.8 |
| C42—C37—C38 | 117.6 (2) | C75—C76—H76B | 110.8 |
| C37—C38—H38 | 119.2 | H76A—C76—H76B | 108.9 |
| C39—C38—C37 | 121.5 (2) | O5—C77—H77A | 110.8 |
| C39—C38—H38 | 119.2 | O5—C77—H77B | 110.8 |
| C38—C39—H39 | 120.0 | O5—C77—C78 | 104.9 (8) |
| C40—C39—C38 | 120.1 (2) | H77A—C77—H77B | 108.8 |
| C40—C39—H39 | 120.0 | C78—C77—H77A | 110.8 |
| C39—C40—H40 | 120.5 | C78—C77—H77B | 110.8 |
| C39—C40—C41 | 119.1 (2) | C77—C78—H78A | 110.9 |
| C41—C40—H40 | 120.5 | C77—C78—H78B | 110.9 |
| C40—C41—H41 | 119.7 | H78A—C78—H78B | 108.9 |
| C40—C41—C42 | 120.6 (3) | C79—C78—C77 | 104.3 (7) |
| C42—C41—H41 | 119.7 | C79—C78—H78A | 110.9 |
| C37—C42—H42 | 119.5 | C79—C78—H78B | 110.9 |
| C41—C42—C37 | 121.0 (2) | O5—C77'—H77C | 110.7 |
| C41—C42—H42 | 119.5 | O5—C77'—H77D | 110.7 |
| C44—C43—P3 | 125.10 (16) | O5—C77'—C78' | 105.4 (14) |
| C44—C43—C48 | 117.9 (2) | H77C—C77'—H77D | 108.8 |
| C48—C43—P3 | 116.90 (16) | C78'—C77'—H77C | 110.7 |
| C43—C44—H44 | 119.5 | C78'—C77'—H77D | 110.7 |
| C45—C44—C43 | 121.1 (2) | C77'—C78'—H78C | 111.2 |
| C45—C44—H44 | 119.5 | C77'—C78'—H78D | 111.2 |
| C44—C45—H45 | 119.9 | C77'—C78'—C79 | 103.0 (14) |
| C46—C45—C44 | 120.2 (2) | H78C—C78'—H78D | 109.1 |
| C46—C45—H45 | 119.9 | C79—C78'—H78C | 111.2 |
| C45—C46—H46 | 120.3 | C79—C78'—H78D | 111.2 |
| C45—C46—C47 | 119.3 (2) | C78—C79—H79A | 110.7 |
| C47—C46—H46 | 120.3 | C78—C79—H79B | 110.7 |
| C46—C47—H47 | 119.7 | C78'—C79—H79C | 112.3 |
| C48—C47—C46 | 120.5 (2) | C78'—C79—H79D | 112.3 |
| C48—C47—H47 | 119.7 | H79A—C79—H79B | 108.8 |
| C43—C48—H48 | 119.6 | H79C—C79—H79D | 109.9 |
| C47—C48—C43 | 120.8 (2) | C80—C79—C78 | 105.4 (5) |
| C47—C48—H48 | 119.6 | C80—C79—C78' | 97.5 (9) |
| C50—C49—P4 | 122.85 (17) | C80—C79—H79A | 110.7 |
| C50—C49—C54 | 117.6 (2) | C80—C79—H79B | 110.7 |
| C54—C49—P4 | 119.39 (17) | C80—C79—H79C | 112.3 |
| C49—C50—H50 | 119.5 | C80—C79—H79D | 112.3 |
| C51—C50—C49 | 121.0 (2) | O5—C80—C79 | 104.3 (2) |
| C51—C50—H50 | 119.5 | O5—C80—H80A | 110.9 |
| C50—C51—H51 | 119.6 | O5—C80—H80B | 110.9 |
| C52—C51—C50 | 120.7 (2) | C79—C80—H80A | 110.9 |
| C52—C51—H51 | 119.6 | C79—C80—H80B | 110.9 |
| C51—C52—H52 | 120.5 | H80A—C80—H80B | 108.9 |
| C51—C52—C53 | 119.0 (2) | C81—O6—C84 | 105.6 (3) |
| C53—C52—H52 | 120.5 | O6—C81—H81A | 110.3 |
| C52—C53—H53 | 119.7 | O6—C81—H81B | 110.3 |

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|-------------|-------------|---------------|-----------|
| C52—C53—C54 | 120.6 (2) | O6—C81—C82 | 106.9 (3) |
| C54—C53—H53 | 119.7 | H81A—C81—H81B | 108.6 |
| C49—C54—H54 | 119.5 | C82—C81—H81A | 110.3 |
| C53—C54—C49 | 120.9 (2) | C82—C81—H81B | 110.3 |
| C53—C54—H54 | 119.5 | C81—C82—H82A | 111.0 |
| C56—C55—P4 | 126.10 (17) | C81—C82—H82B | 111.0 |
| C56—C55—C60 | 117.8 (2) | C81—C82—C83 | 103.9 (3) |
| C60—C55—P4 | 116.04 (16) | H82A—C82—H82B | 109.0 |
| C55—C56—H56 | 119.5 | C83—C82—H82A | 111.0 |
| C57—C56—C55 | 121.0 (2) | C83—C82—H82B | 111.0 |
| C57—C56—H56 | 119.5 | C82—C83—H83A | 110.8 |
| C56—C57—H57 | 119.8 | C82—C83—H83B | 110.8 |
| C58—C57—C56 | 120.4 (2) | H83A—C83—H83B | 108.8 |
| C58—C57—H57 | 119.8 | C84—C83—C82 | 104.9 (3) |
| C57—C58—H58 | 120.4 | C84—C83—H83A | 110.8 |
| C57—C58—C59 | 119.2 (2) | C84—C83—H83B | 110.8 |
| C59—C58—H58 | 120.4 | O6—C84—C83 | 107.3 (3) |
| C58—C59—H59 | 119.8 | O6—C84—H84A | 110.2 |
| C60—C59—C58 | 120.4 (2) | O6—C84—H84B | 110.2 |
| C60—C59—H59 | 119.8 | C83—C84—H84A | 110.2 |
| C55—C60—H60 | 119.4 | C83—C84—H84B | 110.2 |
| C59—C60—C55 | 121.2 (2) | H84A—C84—H84B | 108.5 |
| C59—C60—H60 | 119.4 | C88—O7—C85 | 108.9 (8) |
| O1—Mg1—Br1 | 175.61 (6) | O7—C85—H85A | 110.5 |
| O1—Mg1—O2 | 84.9 (4) | O7—C85—H85B | 110.5 |
| O1—Mg1—O2' | 86.6 (8) | O7—C85—C86 | 106.0 (8) |
| O1—Mg1—O3 | 87.6 (6) | H85A—C85—H85B | 108.7 |
| O1—Mg1—O4 | 92.05 (8) | C86—C85—H85A | 110.5 |
| O1—Mg1—O5 | 85.15 (8) | C86—C85—H85B | 110.5 |
| O2—Mg1—Br1 | 90.8 (4) | C85—C86—H86A | 111.1 |
| O2—Mg1—O3 | 92.4 (5) | C85—C86—H86B | 111.1 |
| O2'—Mg1—Br1 | 89.0 (8) | H86A—C86—H86B | 109.1 |
| O3—Mg1—Br1 | 93.8 (6) | C87—C86—C85 | 103.4 (8) |
| O3'—Mg1—Br1 | 95.8 (11) | C87—C86—H86A | 111.1 |
| O3'—Mg1—O1 | 85.1 (11) | C87—C86—H86B | 111.1 |
| O3'—Mg1—O2' | 93.5 (10) | C86—C87—H87A | 111.5 |
| O3'—Mg1—O4 | 92.3 (9) | C86—C87—H87B | 111.5 |
| O3'—Mg1—O5 | 170.2 (11) | H87A—C87—H87B | 109.3 |
| O4—Mg1—Br1 | 92.21 (6) | C88—C87—C86 | 101.4 (9) |
| O4—Mg1—O2 | 176.7 (4) | C88—C87—H87A | 111.5 |
| O4—Mg1—O2' | 173.9 (6) | C88—C87—H87B | 111.5 |
| O4—Mg1—O3 | 86.2 (5) | O7—C88—C87 | 109.1 (8) |
| O4—Mg1—O5 | 87.55 (8) | O7—C88—H88A | 109.9 |
| O5—Mg1—Br1 | 93.96 (6) | O7—C88—H88B | 109.9 |
| O5—Mg1—O2 | 93.5 (3) | C87—C88—H88A | 109.9 |
| O5—Mg1—O2' | 86.4 (6) | C87—C88—H88B | 109.9 |
| O5—Mg1—O3 | 170.2 (6) | H88A—C88—H88B | 108.3 |
| C61—O1—Mg1 | 126.68 (16) | C88'—O7'—C85' | 107.3 (5) |

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| C61—O1—C64 | 108.74 (19) | O7'—C85'—H85C | 110.3 |
| C64—O1—Mg1 | 124.51 (14) | O7'—C85'—H85D | 110.3 |
| C65—O2—Mg1 | 118.2 (9) | O7'—C85'—C86' | 107.0 (5) |
| C68—O2—Mg1 | 127.7 (7) | H85C—C85'—H85D | 108.6 |
| C68—O2—C65 | 106.3 (6) | C86'—C85'—H85C | 110.3 |
| O2—C65—H65A | 110.5 | C86'—C85'—H85D | 110.3 |
| O2—C65—H65B | 110.5 | C85'—C86'—H86C | 111.0 |
| O2—C65—C66 | 106.3 (7) | C85'—C86'—H86D | 111.0 |
| H65A—C65—H65B | 108.7 | H86C—C86'—H86D | 109.0 |
| C66—C65—H65A | 110.5 | C87'—C86'—C85' | 104.0 (5) |
| C66—C65—H65B | 110.5 | C87'—C86'—H86C | 111.0 |
| C65—C66—H66A | 110.9 | C87'—C86'—H86D | 111.0 |
| C65—C66—H66B | 110.9 | C86'—C87'—H87C | 111.4 |
| C65—C66—C67 | 104.2 (6) | C86'—C87'—H87D | 111.4 |
| H66A—C66—H66B | 108.9 | H87C—C87'—H87D | 109.3 |
| C67—C66—H66A | 110.9 | C88'—C87'—C86' | 101.8 (5) |
| C67—C66—H66B | 110.9 | C88'—C87'—H87C | 111.4 |
| C66—C67—H67A | 111.5 | C88'—C87'—H87D | 111.4 |
| C66—C67—H67B | 111.5 | O7'—C88'—C87' | 106.8 (6) |
| H67A—C67—H67B | 109.3 | O7'—C88'—H88C | 110.4 |
| C68—C67—C66 | 101.6 (5) | O7'—C88'—H88D | 110.4 |
| C68—C67—H67A | 111.5 | C87'—C88'—H88C | 110.4 |
| C68—C67—H67B | 111.5 | C87'—C88'—H88D | 110.4 |
| O2—C68—C67 | 102.2 (7) | H88C—C88'—H88D | 108.6 |
| | | | |
| Co1—P1—C1—C2 | -176.18 (19) | C36—P4—C49—C54 | -161.79 (18) |
| Co1—P1—C1—C6 | 3.97 (19) | C36—P4—C55—C56 | 102.6 (2) |
| Co1—P1—C7—C8 | 17.7 (2) | C36—P4—C55—C60 | -79.98 (19) |
| Co1—P1—C7—C12 | -162.17 (16) | C36—C31—C32—C33 | 2.7 (3) |
| Co1—P1—C13—C14 | 83.10 (19) | C37—P3—C31—C32 | 45.8 (2) |
| Co1—P1—C13—C18 | -93.1 (2) | C37—P3—C31—C36 | -138.16 (16) |
| Co1—P2—C6—C1 | -2.31 (19) | C37—P3—C43—C44 | -1.9 (2) |
| Co1—P2—C6—C5 | 178.27 (19) | C37—P3—C43—C48 | 175.40 (18) |
| Co1—P2—C19—C20 | -142.05 (19) | C37—C38—C39—C40 | 2.6 (4) |
| Co1—P2—C19—C24 | 36.7 (2) | C38—C37—C42—C41 | 2.4 (4) |
| Co1—P2—C25—C26 | -103.6 (2) | C38—C39—C40—C41 | 1.0 (4) |
| Co1—P2—C25—C30 | 69.39 (18) | C39—C40—C41—C42 | -2.8 (4) |
| Co1—P3—C31—C32 | 176.95 (17) | C40—C41—C42—C37 | 1.1 (4) |
| Co1—P3—C31—C36 | -6.98 (17) | C42—C37—C38—C39 | -4.3 (3) |
| Co1—P3—C37—C38 | 66.63 (18) | C43—P3—C31—C32 | -54.7 (2) |
| Co1—P3—C37—C42 | -109.3 (2) | C43—P3—C31—C36 | 121.37 (16) |
| Co1—P3—C43—C44 | -138.55 (19) | C43—P3—C37—C38 | -70.98 (18) |
| Co1—P3—C43—C48 | 38.7 (2) | C43—P3—C37—C42 | 113.1 (2) |
| Co1—P4—C36—C31 | 3.70 (17) | C43—C44—C45—C46 | 2.6 (4) |
| Co1—P4—C36—C35 | -176.30 (17) | C44—C43—C48—C47 | 1.6 (4) |
| Co1—P4—C49—C50 | -98.33 (19) | C44—C45—C46—C47 | 0.6 (4) |
| Co1—P4—C49—C54 | 77.26 (19) | C45—C46—C47—C48 | -2.6 (4) |
| Co1—P4—C55—C56 | -138.13 (18) | C46—C47—C48—C43 | 1.5 (4) |

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| Co1—P4—C55—C60 | 39.3 (2) | C48—C43—C44—C45 | -3.6 (4) |
| P1—C1—C2—C3 | 179.60 (19) | C49—P4—C36—C31 | -125.46 (16) |
| P1—C1—C6—P2 | -1.1 (2) | C49—P4—C36—C35 | 54.5 (2) |
| P1—C1—C6—C5 | 178.40 (18) | C49—P4—C55—C56 | 0.5 (2) |
| P1—C7—C8—C9 | -178.85 (17) | C49—P4—C55—C60 | 177.87 (18) |
| P1—C7—C12—C11 | 179.77 (17) | C49—C50—C51—C52 | -0.1 (4) |
| P1—C13—C14—C15 | -179.72 (18) | C50—C49—C54—C53 | -4.1 (3) |
| P1—C13—C18—C17 | 178.61 (19) | C50—C51—C52—C53 | -2.1 (4) |
| P2—C19—C20—C21 | 177.2 (2) | C51—C52—C53—C54 | 1.2 (4) |
| P2—C19—C24—C23 | -178.42 (19) | C52—C53—C54—C49 | 2.0 (4) |
| P2—C25—C26—C27 | 174.1 (2) | C54—C49—C50—C51 | 3.1 (3) |
| P2—C25—C30—C29 | -175.55 (18) | C55—P4—C36—C31 | 133.72 (17) |
| P3—C31—C32—C33 | 178.55 (17) | C55—P4—C36—C35 | -46.3 (2) |
| P3—C31—C36—P4 | 2.0 (2) | C55—P4—C49—C50 | 121.6 (2) |
| P3—C31—C36—C35 | -177.99 (16) | C55—P4—C49—C54 | -62.79 (19) |
| P3—C37—C38—C39 | 179.49 (18) | C55—C56—C57—C58 | 1.1 (4) |
| P3—C37—C42—C41 | 178.3 (2) | C56—C55—C60—C59 | 1.6 (3) |
| P3—C43—C44—C45 | 173.6 (2) | C56—C57—C58—C59 | 0.4 (4) |
| P3—C43—C48—C47 | -175.85 (19) | C57—C58—C59—C60 | -0.9 (4) |
| P4—C49—C50—C51 | 178.80 (19) | C58—C59—C60—C55 | -0.2 (4) |
| P4—C49—C54—C53 | -179.91 (18) | C60—C55—C56—C57 | -2.1 (4) |
| P4—C55—C56—C57 | 175.25 (19) | Mg1—O1—C61—C62 | 169.44 (17) |
| P4—C55—C60—C59 | -175.99 (18) | Mg1—O1—C64—C63 | 165.75 (17) |
| C1—P1—C7—C8 | -103.54 (18) | Mg1—O2—C65—C66 | 129.4 (10) |
| C1—P1—C7—C12 | 76.6 (2) | Mg1—O2—C68—C67 | -107.0 (9) |
| C1—P1—C13—C14 | -156.77 (19) | Mg1—O2'—C65'—C66' | 117 (2) |
| C1—P1—C13—C18 | 27.0 (2) | Mg1—O2'—C68'—C67' | -124.5 (18) |
| C1—C2—C3—C4 | 1.3 (4) | Mg1—O3—C69—C70 | 165.0 (13) |
| C2—C1—C6—P2 | 179.09 (17) | Mg1—O3—C72—C71 | 172.0 (16) |
| C2—C1—C6—C5 | -1.5 (3) | Mg1—O3'—C69'—C70' | -152 (2) |
| C2—C3—C4—C5 | -0.1 (4) | Mg1—O3'—C72'—C71' | 178 (2) |
| C3—C4—C5—C6 | -1.9 (4) | Mg1—O4—C73—C74 | 145.5 (2) |
| C4—C5—C6—P2 | -177.9 (2) | Mg1—O4—C76—C75 | -133.0 (2) |
| C4—C5—C6—C1 | 2.7 (4) | Mg1—O5—C77—C78 | -153.6 (9) |
| C6—P2—C19—C20 | 99.4 (2) | Mg1—O5—C77'—C78' | -178.6 (16) |
| C6—P2—C19—C24 | -81.83 (19) | Mg1—O5—C80—C79 | 151.2 (2) |
| C6—P2—C25—C26 | 19.1 (2) | O1—C61—C62—C63 | 32.8 (3) |
| C6—P2—C25—C30 | -167.86 (17) | O2—C65—C66—C67 | -5.4 (15) |
| C6—C1—C2—C3 | -0.6 (4) | C65—O2—C68—C67 | 41.1 (12) |
| C7—P1—C1—C2 | -43.2 (2) | C65—C66—C67—C68 | 29.3 (10) |
| C7—P1—C1—C6 | 136.93 (17) | C66—C67—C68—O2 | -43.0 (7) |
| C7—P1—C13—C14 | -57.3 (2) | C68—O2—C65—C66 | -22.3 (16) |
| C7—P1—C13—C18 | 126.4 (2) | O2'—C65'—C66'—C67' | 33 (3) |
| C7—C8—C9—C10 | -0.9 (4) | C65'—O2'—C68'—C67' | 29 (3) |
| C8—C7—C12—C11 | -0.1 (3) | C65'—C66'—C67'—C68' | -16 (2) |
| C8—C9—C10—C11 | -0.2 (4) | C66'—C67'—C68'—O2' | -7 (2) |
| C9—C10—C11—C12 | 1.1 (3) | C68'—O2'—C65'—C66' | -39 (3) |
| C10—C11—C12—C7 | -1.0 (3) | O3—C69—C70—C71 | 32 (2) |

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| C12—C7—C8—C9 | 1.0 (3) | C69—O3—C72—C71 | -8 (3) |
| C13—P1—C1—C2 | 55.5 (2) | C69—C70—C71—C72 | -36 (3) |
| C13—P1—C1—C6 | -124.38 (18) | C70—C71—C72—O3 | 28 (3) |
| C13—P1—C7—C8 | 155.21 (18) | C72—O3—C69—C70 | -15 (3) |
| C13—P1—C7—C12 | -24.6 (2) | O3'—C69'—C70'—C71' | -29 (4) |
| C13—C14—C15—C16 | 1.7 (4) | C69'—O3'—C72'—C71' | 25 (5) |
| C14—C13—C18—C17 | 2.3 (4) | C69'—C70'—C71'—C72' | 44 (4) |
| C14—C15—C16—C17 | 0.9 (4) | C70'—C71'—C72'—O3' | -42 (5) |
| C15—C16—C17—C18 | -1.8 (4) | C72'—O3'—C69'—C70' | 3 (5) |
| C16—C17—C18—C13 | 0.2 (4) | O4—C73—C74—C75 | 4.0 (4) |
| C18—C13—C14—C15 | -3.3 (3) | O5—C77—C78—C79 | -19.2 (19) |
| C19—P2—C6—C1 | 124.76 (18) | O5—C77'—C78'—C79 | 22 (3) |
| C19—P2—C6—C5 | -54.7 (2) | C61—O1—C64—C63 | -11.4 (3) |
| C19—P2—C25—C26 | 118.6 (2) | C61—C62—C63—C64 | -38.8 (3) |
| C19—P2—C25—C30 | -68.35 (19) | C62—C63—C64—O1 | 31.4 (3) |
| C19—C20—C21—C22 | 1.4 (4) | C64—O1—C61—C62 | -13.5 (3) |
| C20—C19—C24—C23 | 0.4 (4) | C73—O4—C76—C75 | 35.2 (3) |
| C20—C21—C22—C23 | -0.1 (4) | C73—C74—C75—C76 | 16.7 (4) |
| C21—C22—C23—C24 | -1.0 (4) | C74—C75—C76—O4 | -31.7 (3) |
| C22—C23—C24—C19 | 0.8 (4) | C76—O4—C73—C74 | -24.5 (3) |
| C24—C19—C20—C21 | -1.6 (4) | C77—O5—C80—C79 | -40.3 (11) |
| C25—P2—C6—C1 | -132.97 (17) | C77—C78—C79—C80 | -4.8 (14) |
| C25—P2—C6—C5 | 47.6 (2) | C78—C79—C80—O5 | 27.1 (7) |
| C25—P2—C19—C20 | -3.9 (2) | C77'—O5—C80—C79 | -35 (2) |
| C25—P2—C19—C24 | 174.82 (18) | C77'—C78'—C79—C80 | -41 (2) |
| C25—C26—C27—C28 | 0.5 (4) | C78'—C79—C80—O5 | 46.9 (8) |
| C26—C25—C30—C29 | -2.0 (3) | C80—O5—C77—C78 | 37.2 (18) |
| C26—C27—C28—C29 | -1.1 (4) | C80—O5—C77'—C78' | 7 (3) |
| C27—C28—C29—C30 | 0.2 (4) | O6—C81—C82—C83 | -25.5 (4) |
| C28—C29—C30—C25 | 1.4 (4) | C81—O6—C84—C83 | -30.0 (4) |
| C30—C25—C26—C27 | 1.1 (4) | C81—C82—C83—C84 | 7.0 (4) |
| C31—P3—C37—C38 | -169.09 (17) | C82—C83—C84—O6 | 13.6 (4) |
| C31—P3—C37—C42 | 15.0 (2) | C84—O6—C81—C82 | 34.9 (4) |
| C31—P3—C43—C44 | 102.6 (2) | O7—C85—C86—C87 | -29.7 (12) |
| C31—P3—C43—C48 | -80.09 (19) | C85—O7—C88—C87 | 7.5 (13) |
| C31—C32—C33—C34 | -2.2 (3) | C85—C86—C87—C88 | 32.6 (12) |
| C32—C31—C36—P4 | 178.34 (16) | C86—C87—C88—O7 | -25.5 (13) |
| C32—C31—C36—C35 | -1.7 (3) | C88—O7—C85—C86 | 14.1 (12) |
| C32—C33—C34—C35 | 0.7 (4) | O7'—C85'—C86'—C87' | 1.2 (8) |
| C33—C34—C35—C36 | 0.3 (3) | C85'—O7'—C88'—C87' | -36.0 (8) |
| C34—C35—C36—P4 | -179.81 (17) | C85'—C86'—C87'—C88' | -20.8 (9) |
| C34—C35—C36—C31 | 0.2 (3) | C86'—C87'—C88'—O7' | 35.3 (9) |
| C36—P4—C49—C50 | 22.6 (2) | C88'—O7'—C85'—C86' | 21.4 (7) |

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

| $D-H\cdots A$ | $D-H$ | $H\cdots A$ | $D\cdots A$ | $D-H\cdots A$ |
|----------------------|-------|-------------|-------------|---------------|
| C62—H62B \cdots O6 | 0.97 | 2.48 | 3.438 (4) | 167 |

| | | | | |
|----------------|------|------|-----------|-----|
| C63—H63B···O7 | 0.97 | 2.59 | 3.555 (8) | 179 |
| C63—H63B···O7' | 0.97 | 2.63 | 3.565 (6) | 162 |
