

Acta Cryst. (1999). C55, IUC9900103 [doi:10.1107/S0108270199098911]

trans-Bis(tert-butylisocyanido-C)bis(triphenylphosphine-P)rhodium(I) nido-7,8-dicarbaundecaborane

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Abstract

In the crystal structure of $[\text{Rh}(\text{t-BuNC})_2(\text{PPh}_3)_2]^+[\text{7,8-nido-C}_2\text{H}_{12}\text{B}_9]^-$ the Rh atom lies on an inversion centre and has a square-planar geometry with Rh—P 2.3217 (6), Rh—C 1.954 (2) Å, P—Rh—C 86.74 (7)°. The carborane anion lies about a twofold axis and has disordered C and B atoms.

Comment

Disorder of the atoms in both cation and anion precludes any meaningful discussion of the geometry of this complex.

Experimental

The title compound was synthesized in 33% yield from the reaction of a CH_2Cl_2 solution of [3,3-bis(triphenylphosphine)-3-hydrido-1,2-dicarba-3-rhodadodecaborane] with *tert*-BuNC at reflux temperature for 8 h followed by stirring at 293 K for a further 18 h. Crystals suitable for X-ray analysis were grown by evaporation of a CH_2Cl_2 solution of the compound.

Refinement

The compound crystallized in the monoclinic system; space group $C2/c$ or Cc from the systematic absences. Both options were tried and $C2/c$ chosen.

The $[\text{C}_2\text{H}_{12}\text{B}_9]^-$ anion lies about a two fold axis and is disordered with the two C and nine B atoms distributed unequally over 12 sites; as a result of the imposed twofold symmetry, only six of these atom sites are unique. Allowing for these six sites as B atoms (labelled as B1 - B6) and allowing the occupancies to refine, indicated that four of the sites (B3 - B6) were consistent with their being unit occupancy B atoms and that the carbon-boron disorder was limited to sites labelled as B1 and B2. To keep the electron count approximately correct, and noting that there should be 28.5 non-H electrons in half of a C_2B_9 cage, we allowed for the B1, B2 sites as B atoms but with occupancies linked so that their occupancy sum was 1.70 (occupancies refined to B1 0.846 (11), B2 0.854 (11); *i.e.* each of the B1 and B2 sites is effectively shared by (0.50 C and 0.25B). In this way we effectively allowed for the required 28.5 electrons in the half of the C_2B_9 cage ($4 \times 5.0 + 1.70 \times 5.0 = 28.5$).

In the $[\text{C}_2\text{H}_{12}\text{B}_9]^-$ cage, each of the C and B atoms should have one terminal H atom and one of the B sites should have an additional H atom bonded to it; because of the disorder noted above, we were not able to decide the location of the twelfth H atom. Difference maps of the asymmetric unit showed six maxima (one per B or C/B site of the cage) consistent with the H atoms of terminal B—H or C—H groups and we allowed for these as riding atoms (B(or C)—H 1.10 Å) with unit occupancy. Changing the H atom occupancies on the C/B sites B1 and B2 had no significant effect on the dimensions or R -factors.

The Rh atom of the $[(\text{Ph}_3\text{P})_2(\text{CNBu}^t)_2\text{Rh}]^+$ cation lies on an inversion centre and the methyl groups of the unique *tert*-butyl group are disordered unequally over two sites (0.669 (6)/0.331 (6)); the minor site C atoms were refined with isotropic vibration parameters. Cation H atoms were treated as riding atoms (C—H 0.93 and 0.96 Å)

Structure solution and refinement in space group *Cc* was also considered. With $Z=4$, this choice of space group requires no crystallographic symmetry for either cation or anion. As all attempts consistently yielded a scrambled anion cage as found for the *C2/c* solution and refinements always yielded non-sensible anisotropic vibration parameters; this approach was abandoned.

Computing details

Data collection: *CAD-4-PC* Software (Enraf-Nonius, 1992); cell refinement: *SET4 & CELDIM* in *CAD-4-PC* Software (Enraf-Nonius, 1992); data reduction: *DATRD2* in *NRCVAX96* (Gabe *et al.*, 1989); program(s) used to solve structure: *NRCVAX96* via Patterson heavy-atom method; program(s) used to refine structure: *NRCVAX96* and *SHELXL97* (Sheldrick, 1997); software used to prepare material for publication: *NRCVAX96*, *SHELXL97* and *WORDPERFECT* macro PREP8 (Ferguson, 1998).

trans-Bis(triphenylphosphine)bis(*tert*-butylisocyanide)rhodium(I) *nido*-7,8-dicarbaundecaborane.

Crystal data

$[\text{Rh}(\text{C}_5\text{H}_9\text{N}_1)_2(\text{C}_{18}\text{H}_{15}\text{P}_1)_2](\text{C}_2\text{H}_{12}\text{B}_9)$	$V = 4931.8 (9) \text{ \AA}^3$
$M_r = 927.12$	$Z = 4$
Monoclinic, <i>C2/c</i>	Mo $K\alpha$
$a = 23.7306 (14) \text{ \AA}$	$\mu = 0.45 \text{ mm}^{-1}$
$b = 11.9542 (12) \text{ \AA}$	$T = 294 (1) \text{ K}$
$c = 17.468 (3) \text{ \AA}$	$0.49 \times 0.48 \times 0.21 \text{ mm}$
$\beta = 95.597 (19)^\circ$	

Data collection

Enraf-Nonius CAD-4 diffractometer	4206 reflections with $I > 2\sigma(I)$
Absorption correction: Gaussian (ABSO in NRCVAX; Gabe <i>et al.</i> , 1989)	$R_{\text{int}} = 0.010$
$T_{\min} = 0.822$, $T_{\max} = 0.918$	3 standard reflections
5506 measured reflections	every 120 min
5339 independent reflections	intensity decay: no decay, variation 1.1%

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.034$	37 restraints
$wR(F^2) = 0.090$	H-atom parameters constrained
$S = 1.19$	$\Delta\rho_{\max} = 0.65 \text{ e \AA}^{-3}$
5339 reflections	$\Delta\rho_{\min} = -0.40 \text{ e \AA}^{-3}$
303 parameters	

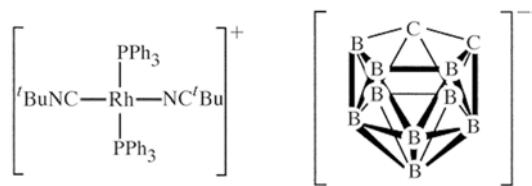
Acknowledgements

We thank Dr J. F. Gallagher and Dublin City University for the funds to purchase the X-ray tube used in the data collections.

References

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Scheme 1



supplementary materials

trans-Bis(triphenylphosphine)bis(tert-butylisocyanide)rhodium(I) nido-7,8-dicarbaundecaborane.***Crystal data***

[Rh(C ₅ H ₉ N ₁) ₂ (C ₁₈ H ₁₅ P ₁) ₂](C ₂ H ₁₂ B ₉)	? #Insert any comments here.
$M_r = 927.12$	$D_x = 1.249 \text{ Mg m}^{-3}$
Monoclinic, $C2/c$	Mo $K\alpha$ radiation
$a = 23.7306 (14) \text{ \AA}$	$\lambda = 0.71073 \text{ \AA}$
$b = 11.9542 (12) \text{ \AA}$	Cell parameters from 25 reflections
$c = 17.468 (3) \text{ \AA}$	$\theta = 16.7\text{--}23.6^\circ$
$\beta = 95.597 (19)^\circ$	$\mu = 0.45 \text{ mm}^{-1}$
$V = 4931.8 (9) \text{ \AA}^3$	$T = 294 (1) \text{ K}$
$Z = 4$	Plate, colourless
$F_{000} = 1928$	$0.49 \times 0.48 \times 0.21 \text{ mm}$

Data collection

Enraf-Nonius CAD-4 diffractometer	$R_{\text{int}} = 0.010$
Radiation source: X-ray tube	$\theta_{\text{max}} = 26.9^\circ$
Monochromator: graphite	$\theta_{\text{min}} = 2.2^\circ$
$T = 294(1) \text{ K}$	$h = -30 \rightarrow 30$
$\theta/2\theta$ scans	$k = 0 \rightarrow 15$
Absorption correction: Gaussian (ABSO in NRCVAX; Gabe et al., 1989)	$l = 0 \rightarrow 22$
$T_{\text{min}} = 0.822$, $T_{\text{max}} = 0.918$	3 standard reflections
5506 measured reflections	every 120 min
5339 independent reflections	intensity decay: no decay, variation 1.1%
4206 reflections with $I > 2\sigma(I)$	

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.034$	H-atom parameters constrained
$wR(F^2) = 0.090$	$w = 1/[\sigma^2(F_o^2) + (0.0448P)^2 + 1.2748P]$ where $P = (F_o^2 + 2F_c^2)/3$
$S = 1.19$	$(\Delta/\sigma)_{\text{max}} = <0.001$
5339 reflections	$\Delta\rho_{\text{max}} = 0.65 \text{ e \AA}^{-3}$
303 parameters	$\Delta\rho_{\text{min}} = -0.40 \text{ e \AA}^{-3}$
37 restraints	Extinction correction: none
Primary atom site location: structure-invariant direct methods	

supplementary materials

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)
Rh1	0.2500	0.2500	0.0000	0.03055 (9)	
P1	0.18349 (2)	0.14578 (5)	0.05820 (3)	0.03089 (14)	
C1	0.29472 (10)	0.1128 (2)	-0.00029 (14)	0.0364 (5)	
N1	0.31841 (9)	0.02938 (19)	0.00239 (13)	0.0457 (5)	
C2	0.34491 (12)	-0.0798 (2)	0.01578 (16)	0.0528 (7)	
C3	0.3033 (2)	-0.1518 (5)	0.0549 (4)	0.101 (3)	0.670 (6)
C4	0.3957 (2)	-0.0628 (5)	0.0749 (3)	0.085 (2)	0.670 (6)
C5	0.3603 (4)	-0.1271 (6)	-0.0560 (3)	0.149 (4)	0.670 (6)
C3A	0.3567 (6)	-0.1092 (13)	0.0981 (4)	0.128 (7)*	0.330 (6)
C4A	0.3994 (3)	-0.0802 (11)	-0.0239 (6)	0.090 (4)*	0.330 (6)
C5A	0.3045 (4)	-0.1616 (9)	-0.0265 (6)	0.084 (4)*	0.330 (6)
C11	0.21589 (10)	0.0677 (2)	0.14148 (13)	0.0364 (5)	
C12	0.26577 (11)	0.1092 (3)	0.17886 (15)	0.0484 (6)	
C13	0.29145 (14)	0.0560 (3)	0.24343 (17)	0.0654 (9)	
C14	0.26790 (15)	-0.0392 (3)	0.27079 (18)	0.0706 (10)	
C15	0.21840 (16)	-0.0808 (3)	0.23442 (19)	0.0702 (10)	
C16	0.19229 (13)	-0.0280 (3)	0.16924 (17)	0.0550 (7)	
C21	0.12641 (10)	0.22363 (19)	0.09724 (14)	0.0342 (5)	
C22	0.08396 (11)	0.2691 (2)	0.04636 (16)	0.0464 (7)	
C23	0.04132 (12)	0.3326 (3)	0.07307 (18)	0.0579 (8)	
C24	0.04082 (13)	0.3523 (3)	0.15060 (19)	0.0611 (8)	
C25	0.08306 (15)	0.3085 (3)	0.20129 (18)	0.0655 (9)	
C26	0.12559 (13)	0.2436 (2)	0.17534 (16)	0.0507 (7)	
C31	0.14397 (10)	0.0437 (2)	-0.00381 (13)	0.0362 (5)	
C32	0.16246 (12)	0.0170 (2)	-0.07455 (15)	0.0473 (6)	
C33	0.13192 (15)	-0.0576 (3)	-0.12333 (18)	0.0633 (9)	
C34	0.08302 (15)	-0.1056 (3)	-0.1027 (2)	0.0644 (9)	
C35	0.06472 (13)	-0.0808 (3)	-0.0327 (2)	0.0607 (8)	
C36	0.09423 (11)	-0.0053 (2)	0.01628 (17)	0.0495 (6)	
B1	0.4978 (3)	0.3319 (4)	0.1726 (3)	0.079 (2)	0.846 (11)
B2	0.4685 (2)	0.3694 (4)	0.2515 (4)	0.085 (2)	0.855 (11)
B3	0.4417 (2)	0.2630 (4)	0.2961 (3)	0.0756 (13)	
B4	0.46244 (15)	0.1435 (3)	0.2468 (2)	0.0598 (9)	
B5	0.49753 (18)	0.1882 (4)	0.1673 (2)	0.0650 (10)	
B6	0.43758 (19)	0.2620 (4)	0.1945 (3)	0.0724 (12)	
H3A	0.2942	-0.1157	0.1012	0.151*	0.670 (6)
H3B	0.3200	-0.2234	0.0673	0.151*	0.670 (6)
H3C	0.2694	-0.1617	0.0208	0.151*	0.670 (6)
H4A	0.3832	-0.0322	0.1212	0.128*	0.670 (6)
H4B	0.4219	-0.0121	0.0548	0.128*	0.670 (6)
H4C	0.4140	-0.1334	0.0862	0.128*	0.670 (6)
H5A	0.3870	-0.0787	-0.0774	0.223*	0.670 (6)
H5B	0.3271	-0.1343	-0.0916	0.223*	0.670 (6)
H5C	0.3771	-0.1994	-0.0463	0.223*	0.670 (6)
H3A1	0.3830	-0.0565	0.1227	0.192*	0.330 (6)

H3A2	0.3725	-0.1830	0.1026	0.192*	0.330 (6)
H3A3	0.3220	-0.1070	0.1223	0.192*	0.330 (6)
H4A1	0.3904	-0.0702	-0.0783	0.135*	0.330 (6)
H4A2	0.4186	-0.1503	-0.0145	0.135*	0.330 (6)
H4A3	0.4235	-0.0203	-0.0040	0.135*	0.330 (6)
H5A1	0.2969	-0.1386	-0.0791	0.126*	0.330 (6)
H5A2	0.2697	-0.1633	-0.0026	0.126*	0.330 (6)
H5A3	0.3212	-0.2348	-0.0245	0.126*	0.330 (6)
H12	0.2821	0.1732	0.1604	0.058*	
H13	0.3247	0.0847	0.2684	0.079*	
H14	0.2854	-0.0753	0.3139	0.085*	
H15	0.2022	-0.1446	0.2534	0.084*	
H16	0.1590	-0.0572	0.1444	0.066*	
H22	0.0842	0.2568	-0.0062	0.056*	
H23	0.0129	0.3621	0.0385	0.070*	
H24	0.0121	0.3949	0.1686	0.073*	
H25	0.0830	0.3226	0.2537	0.079*	
H26	0.1536	0.2135	0.2102	0.061*	
H32	0.1954	0.0491	-0.0893	0.057*	
H33	0.1447	-0.0752	-0.1705	0.076*	
H34	0.0625	-0.1546	-0.1361	0.077*	
H35	0.0323	-0.1149	-0.0179	0.073*	
H36	0.0807	0.0128	0.0629	0.059*	
H1	0.4957	0.3821	0.1196	0.094*	
H2	0.4464	0.4499	0.2543	0.102*	
H3	0.4029	0.2653	0.3258	0.091*	
H4	0.4378	0.0654	0.2447	0.072*	
H5	0.4961	0.1405	0.1133	0.078*	
H6	0.3970	0.2645	0.1581	0.087*	

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Rh1	0.02694 (13)	0.02842 (13)	0.03716 (14)	0.00416 (11)	0.00767 (9)	0.00550 (11)
P1	0.0280 (3)	0.0321 (3)	0.0330 (3)	0.0007 (2)	0.0053 (2)	0.0034 (2)
C1	0.0305 (12)	0.0356 (13)	0.0443 (13)	0.0039 (10)	0.0090 (10)	0.0056 (10)
N1	0.0390 (12)	0.0400 (13)	0.0583 (14)	0.0098 (10)	0.0053 (10)	0.0034 (10)
C2	0.0532 (17)	0.0373 (15)	0.0674 (19)	0.0183 (13)	0.0033 (14)	0.0027 (13)
C3	0.070 (4)	0.051 (3)	0.184 (7)	0.008 (3)	0.027 (4)	0.032 (4)
C4	0.070 (4)	0.068 (3)	0.110 (5)	0.016 (3)	-0.034 (3)	0.011 (3)
C5	0.270 (9)	0.113 (5)	0.064 (4)	0.122 (6)	0.023 (5)	-0.007 (4)
C11	0.0350 (13)	0.0398 (13)	0.0349 (12)	0.0043 (10)	0.0061 (9)	0.0055 (10)
C12	0.0448 (15)	0.0586 (17)	0.0416 (14)	-0.0016 (13)	0.0029 (11)	0.0022 (12)
C13	0.0515 (18)	0.104 (3)	0.0390 (15)	0.0086 (18)	-0.0044 (13)	0.0016 (16)
C14	0.073 (2)	0.095 (3)	0.0434 (16)	0.029 (2)	0.0011 (15)	0.0208 (17)
C15	0.090 (3)	0.063 (2)	0.059 (2)	0.0075 (19)	0.0123 (18)	0.0288 (16)
C16	0.0568 (18)	0.0507 (17)	0.0566 (17)	-0.0015 (14)	0.0011 (13)	0.0183 (13)
C21	0.0294 (11)	0.0348 (13)	0.0392 (12)	-0.0018 (9)	0.0085 (9)	-0.0005 (9)

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C22	0.0380 (14)	0.0580 (19)	0.0435 (14)	0.0107 (12)	0.0048 (11)	-0.0014 (12)
C23	0.0378 (15)	0.071 (2)	0.0642 (18)	0.0156 (14)	0.0024 (13)	-0.0028 (15)
C24	0.0456 (17)	0.065 (2)	0.075 (2)	0.0121 (15)	0.0179 (15)	-0.0147 (16)
C25	0.069 (2)	0.082 (2)	0.0478 (17)	0.0153 (19)	0.0162 (15)	-0.0172 (16)
C26	0.0505 (16)	0.0603 (18)	0.0417 (14)	0.0107 (14)	0.0060 (11)	0.0008 (13)
C31	0.0359 (13)	0.0340 (12)	0.0380 (12)	0.0030 (10)	-0.0004 (9)	0.0008 (10)
C32	0.0523 (16)	0.0488 (16)	0.0411 (14)	0.0014 (13)	0.0061 (11)	-0.0046 (12)
C33	0.078 (2)	0.059 (2)	0.0519 (17)	0.0047 (17)	0.0003 (15)	-0.0172 (14)
C34	0.072 (2)	0.0434 (17)	0.073 (2)	-0.0020 (16)	-0.0176 (17)	-0.0139 (15)
C35	0.0456 (17)	0.0500 (18)	0.084 (2)	-0.0098 (14)	-0.0045 (15)	-0.0048 (16)
C36	0.0424 (15)	0.0483 (16)	0.0577 (17)	-0.0072 (13)	0.0052 (12)	-0.0029 (13)
B1	0.103 (4)	0.065 (3)	0.073 (3)	0.011 (3)	0.034 (3)	0.027 (3)
B2	0.106 (4)	0.052 (3)	0.102 (5)	0.020 (3)	0.035 (4)	0.017 (3)
B3	0.076 (3)	0.083 (3)	0.072 (3)	0.018 (2)	0.032 (2)	0.013 (2)
B4	0.052 (2)	0.057 (2)	0.069 (2)	-0.0087 (17)	-0.0017 (16)	0.0113 (18)
B5	0.064 (2)	0.077 (3)	0.053 (2)	0.005 (2)	0.0061 (17)	-0.0049 (19)
B6	0.064 (2)	0.082 (3)	0.071 (2)	0.020 (2)	0.0062 (19)	0.022 (2)

Geometric parameters (\AA , $^\circ$)

Rh1—C1	1.954 (2)	C31—C32	1.388 (3)
Rh1—C1 ⁱ	1.954 (2)	C31—C36	1.393 (4)
Rh1—P1 ⁱ	2.3217 (6)	C32—C33	1.387 (4)
Rh1—P1	2.3217 (6)	C33—C34	1.374 (5)
P1—C31	1.829 (2)	C34—C35	1.370 (5)
P1—C21	1.829 (2)	C35—C36	1.386 (4)
P1—C11	1.833 (2)	B1—B2 ⁱⁱ	1.546 (8)
C1—N1	1.143 (3)	B1—B2	1.664 (8)
N1—C2	1.458 (3)	B1—B3 ⁱⁱ	1.700 (8)
C2—C5	1.454 (4)	B1—B5	1.721 (7)
C2—C3A	1.480 (6)	B1—B6	1.730 (7)
C2—C5A	1.511 (5)	B2—B2 ⁱⁱ	1.500 (11)
C2—C3	1.521 (5)	B2—B1 ⁱⁱ	1.546 (8)
C2—C4	1.523 (4)	B2—B3	1.651 (7)
C2—C4A	1.526 (6)	B2—B6	1.743 (8)
C11—C16	1.383 (4)	B3—B1 ⁱⁱ	1.700 (8)
C11—C12	1.387 (4)	B3—B5 ⁱⁱ	1.763 (6)
C12—C13	1.384 (4)	B3—B4	1.763 (6)
C13—C14	1.373 (5)	B3—B6	1.769 (6)
C14—C15	1.373 (5)	B4—B6	1.757 (5)
C15—C16	1.393 (4)	B4—B5	1.770 (6)
C21—C26	1.387 (4)	B4—B4 ⁱⁱ	1.775 (7)
C21—C22	1.388 (4)	B4—B5 ⁱⁱ	1.778 (5)
C22—C23	1.382 (4)	B5—B3 ⁱⁱ	1.763 (6)
C23—C24	1.376 (4)	B5—B6	1.777 (6)
C24—C25	1.375 (5)	B5—B4 ⁱⁱ	1.778 (5)
C25—C26	1.383 (4)		

C1—Rh1—C1 ⁱ	180.0	B2 ⁱⁱ —B1—B3 ⁱⁱ	61.0 (3)
C1—Rh1—P1 ⁱ	93.25 (7)	B2—B1—B3 ⁱⁱ	105.8 (4)
C1 ⁱ —Rh1—P1 ⁱ	86.75 (7)	B2 ⁱⁱ —B1—B5	109.5 (4)
C1—Rh1—P1	86.74 (7)	B2—B1—B5	108.3 (3)
C1 ⁱ —Rh1—P1	93.25 (7)	B3 ⁱⁱ —B1—B5	62.0 (3)
P1 ⁱ —Rh1—P1	180.00 (2)	B2 ⁱⁱ —B1—B6	108.7 (4)
C31—P1—C21	101.81 (11)	B2—B1—B6	61.8 (3)
C31—P1—C11	106.35 (11)	B3 ⁱⁱ —B1—B6	112.6 (3)
C21—P1—C11	103.40 (11)	B5—B1—B6	62.0 (3)
C31—P1—Rh1	115.47 (8)	B2 ⁱⁱ —B2—B1 ⁱⁱ	66.2 (5)
C21—P1—Rh1	116.74 (8)	B2 ⁱⁱ —B2—B3	116.6 (4)
C11—P1—Rh1	111.74 (8)	B1 ⁱⁱ —B2—B3	64.1 (3)
N1—C1—Rh1	175.6 (2)	B2 ⁱⁱ —B2—B1	58.2 (4)
C1—N1—C2	171.9 (3)	B1 ⁱⁱ —B2—B1	114.2 (4)
C5—C2—N1	110.5 (3)	B3—B2—B1	113.3 (4)
C5—C2—C3A	134.9 (7)	B2 ⁱⁱ —B2—B6	110.2 (4)
N1—C2—C3A	113.9 (7)	B1 ⁱⁱ —B2—B6	115.5 (4)
C5—C2—C5A	61.9 (4)	B3—B2—B6	62.8 (3)
N1—C2—C5A	104.9 (5)	B1—B2—B6	61.0 (3)
C3A—C2—C5A	111.8 (4)	B2—B3—B1 ⁱⁱ	54.9 (3)
C5—C2—C3	113.3 (3)	B2—B3—B5 ⁱⁱ	102.8 (4)
N1—C2—C3	106.9 (3)	B1 ⁱⁱ —B3—B5 ⁱⁱ	59.6 (3)
C3A—C2—C3	59.9 (5)	B2—B3—B4	104.7 (3)
C5A—C2—C3	56.2 (4)	B1 ⁱⁱ —B3—B4	106.4 (3)
C5—C2—C4	112.8 (3)	B5 ⁱⁱ —B3—B4	60.6 (2)
N1—C2—C4	106.9 (3)	B2—B3—B6	61.1 (3)
C3A—C2—C4	46.4 (5)	B1 ⁱⁱ —B3—B6	106.7 (3)
C5A—C2—C4	147.2 (5)	B5 ⁱⁱ —B3—B6	108.9 (3)
C3—C2—C4	106.0 (3)	B4—B3—B6	59.7 (2)
C5—C2—C4A	46.5 (4)	B6—B4—B3	60.4 (2)
N1—C2—C4A	107.5 (5)	B6—B4—B5	60.5 (2)
C3A—C2—C4A	110.7 (4)	B3—B4—B5	108.3 (3)
C5A—C2—C4A	107.6 (4)	B6—B4—B4 ⁱⁱ	108.5 (2)
C3—C2—C4A	144.9 (6)	B3—B4—B4 ⁱⁱ	107.2 (3)
C4—C2—C4A	70.1 (4)	B5—B4—B4 ⁱⁱ	60.2 (3)
C16—C11—C12	119.2 (2)	B6—B4—B5 ⁱⁱ	108.7 (3)
C16—C11—P1	123.2 (2)	B3—B4—B5 ⁱⁱ	59.7 (3)
C12—C11—P1	117.66 (19)	B5—B4—B5 ⁱⁱ	108.5 (3)
C13—C12—C11	120.5 (3)	B4 ⁱⁱ —B4—B5 ⁱⁱ	59.8 (2)
C14—C13—C12	120.2 (3)	B1—B5—B3 ⁱⁱ	58.4 (3)
C15—C14—C13	119.9 (3)	B1—B5—B4	105.0 (3)
C14—C15—C16	120.4 (3)	B3 ⁱⁱ —B5—B4	107.4 (3)
C11—C16—C15	119.9 (3)	B1—B5—B6	59.2 (3)

supplementary materials

C26—C21—C22	118.9 (2)	B3 ⁱⁱ —B5—B6	107.4 (3)
C26—C21—P1	122.43 (19)	B4—B5—B6	59.4 (2)
C22—C21—P1	118.56 (19)	B1—B5—B4 ⁱⁱ	104.9 (3)
C23—C22—C21	120.6 (3)	B3 ⁱⁱ —B5—B4 ⁱⁱ	59.7 (2)
C24—C23—C22	120.2 (3)	B4—B5—B4 ⁱⁱ	60.0 (2)
C25—C24—C23	119.5 (3)	B6—B5—B4 ⁱⁱ	107.5 (3)
C24—C25—C26	120.8 (3)	B1—B6—B2	57.3 (3)
C25—C26—C21	120.0 (3)	B1—B6—B4	105.2 (3)
C32—C31—C36	118.5 (2)	B2—B6—B4	101.2 (3)
C32—C31—P1	119.5 (2)	B1—B6—B3	104.6 (4)
C36—C31—P1	122.0 (2)	B2—B6—B3	56.1 (3)
C33—C32—C31	120.2 (3)	B4—B6—B3	60.0 (2)
C34—C33—C32	120.8 (3)	B1—B6—B5	58.8 (3)
C35—C34—C33	119.6 (3)	B2—B6—B5	102.4 (3)
C34—C35—C36	120.5 (3)	B4—B6—B5	60.1 (2)
C35—C36—C31	120.5 (3)	B3—B6—B5	107.7 (3)
B2 ⁱⁱ —B1—B2	55.6 (4)		

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