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Isopropyltriphenylphosphonium bromide monohydrate

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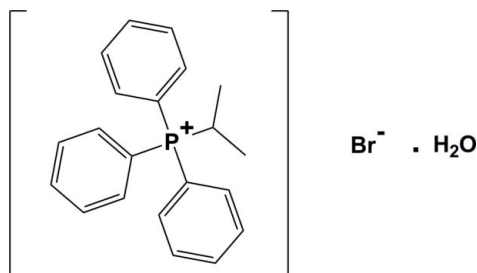
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 Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.007$ Å; disorder in solvent or counterion; R factor = 0.049; wR factor = 0.140; data-to-parameter ratio = 18.6.

In the title water-solvated salt, $\text{C}_{21}\text{H}_{22}\text{P}^+\cdot\text{Br}^-\cdot\text{H}_2\text{O}$, the ionic components are linked by short $\text{C}-\text{H}\cdots\text{Br}$ contacts along the a -axis direction. The two half occupied water molecules are connected to each other by strong $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds and they are also linked to the bromide anion by short $\text{O}-\text{H}\cdots\text{Br}$ contacts.

Related literature

For information on phase-transfer catalysts, see: Asai *et al.* (1994). For the crystal structure of tetraphenylphosphonium bromide, see: Alcock *et al.* (1985). For standard bond lengths, see: Allen *et al.* (1987).



Experimental

Crystal data

 $\text{C}_{21}\text{H}_{22}\text{P}^+\cdot\text{Br}^-\cdot\text{H}_2\text{O}$
 $M_r = 403.28$

 Orthorhombic, $P2_12_12_1$
 $a = 9.078$ (5) Å

 $b = 13.043$ (5) Å

 $c = 17.755$ (5) Å

 $V = 2102.3$ (15) Å³
 $Z = 4$

 Mo $K\alpha$ radiation
 $\mu = 2.04$ mm⁻¹
 $T = 293$ K
 $0.20 \times 0.15 \times 0.13$ mm

Data collection

 Bruker APEXII CCD area-detector diffractometer
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.668$, $T_{\max} = 0.688$

 12177 measured reflections
 4250 independent reflections
 3331 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.034$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.049$
 $wR(F^2) = 0.140$
 $S = 1.03$
 4250 reflections
 228 parameters
 6 restraints

 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.82$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.28$ e Å⁻³
 Absolute structure: Flack (1983),
 1799 Friedel pairs
 Flack parameter: 0.014 (13)

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{O1}-\text{H1O1}\cdots\text{Br1}^{\text{i}}$	0.85	2.86	3.701 (8)	169
$\text{O1}-\text{H2O1}\cdots\text{O2}$	0.85	2.20	2.960 (11)	149
$\text{O2}-\text{H1O2}\cdots\text{Br1}^{\text{ii}}$	0.85	2.68	3.526 (7)	175
$\text{O2}-\text{H2O2}\cdots\text{Br1}^{\text{iii}}$	0.85	2.81	3.598 (8)	154
$\text{C19}-\text{H19}\cdots\text{Br1}^{\text{iv}}$	0.98	2.76	3.723 (4)	169

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

Data collection: APEX2 (Bruker, 2005); cell refinement: SAINT (Bruker, 2005); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: SHELXTL (Sheldrick, 2008); software used to prepare material for publication: SHELXTL.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: SU2327).

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supporting information

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Isopropyltriphenylphosphonium bromide monohydrate

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S1. Comment

The title compound belongs to a family of phase transfer catalysts, which usually contain large alkyl or aryl ions, such as R_4N^+ , R_4P^+ , R_4B^+ etc. Because of their ease of dissolution in water, such salts have a convenient to satisfactory hydrophobic hydration (Asai, *et al.*, 1994).

The asymmetry unit of the title structure consists of one isopropyltriphenylphosphonium cation, one bromide anion and two half occupied water molecules (Fig. 1). In contrast with tetraphenylphosphonium salts, the major character of the cation of the title compound is that one phenyl group has been substituted by an isopropyl group. The fluctuation of the $C_{\text{phenyl}}\text{—P1}$ bond lengths in the title compound [from 1.789 (4)–1.806 (4) Å] is similar to that in the crystal structure of tetraphenylphosphonium bromide [1.801 (3) Å; Alcock, *et al.*, 1985]. The bonds between P1 and the C atoms of phenyl rings are $Csp^2\text{—Psp}^3$ bonds, but the connection between P1 and isopropyl group is typically an $Csp^3\text{—Psp}^3$ bond (Allen, *et al.*, 1987). The bond length of C19—P1 [1.818 (3) Å] associated with the isopropyl group is longer than that involving the phenyl groups, which vary from 1.789 (4) - 1.806 (4) Å.

In the crystal the water molecules are linked to the Br anion by short O—H \cdots Br contacts, and the two half occupied water molecules are connected to one another by strong O—H \cdots O hydrogen bonds (Table 1 and Fig. 2). The large cations and bromide anions are linked by short C19—H19 \cdots Br1 contacts (Table 1 and Fig. 2). These weak interactions also that play an important role in the stabilization of the crystal structure.

S2. Experimental

Triphenyl phosphine (10.5 g) and 2-bromopropane (4.2 mL) were placed in a teflon lined tube. The sealed tube was placed in an autoclave and heated to 433 K for 48 h, then cooled at a rate of 10 K/min. Colourless block-like crystals of the title compound were obtained.

S3. Refinement

The water H atoms were located in difference Fourier maps and were subsequently treated as riding atoms: O—H = 0.85 Å with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{O})$. The C-bound H-atoms were included in calculated positions and treated as riding atoms: C—H = 0.93, 0.96, and 0.98 Å for CH(aromatic), CH₃, and CH(methine) H-atoms, respectively, with $U_{\text{eq}}(\text{parent C-atom})$, where $k = 1.5$ for CH₃ H-atoms and $k = 1.2$ for all other H-atoms.

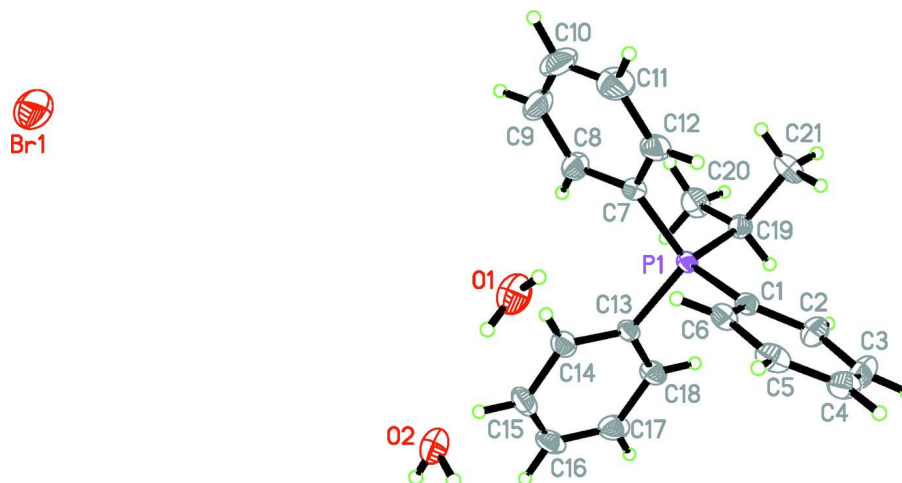


Figure 1

A view of the molecular structure of the title compound, with the numbering scheme and thermal ellipsoids drawn at the 30% probability level.

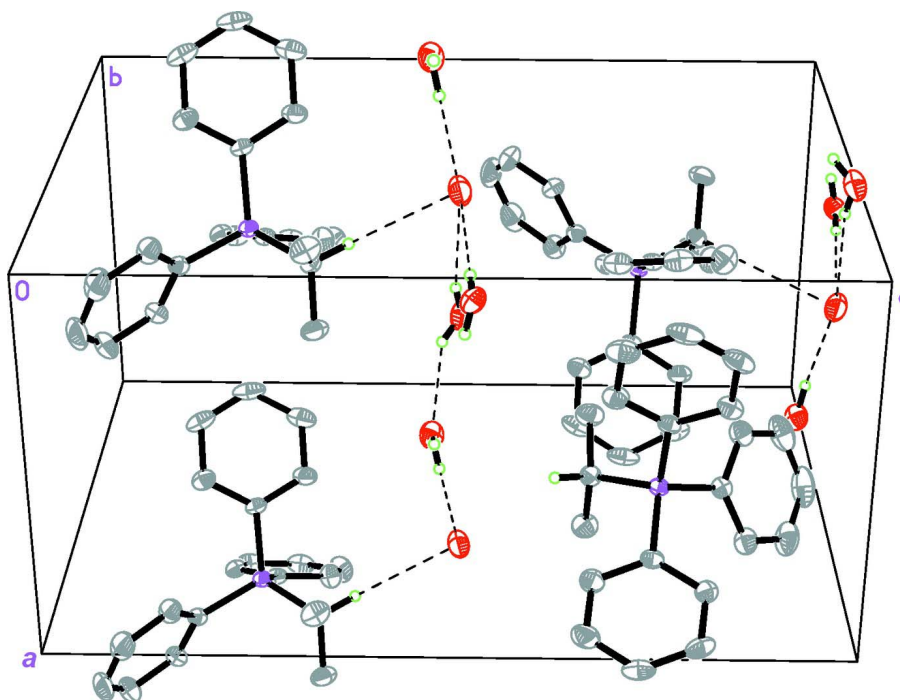


Figure 2

A view of the crystal packing of the title compound, showing the O-H...Br and O-H...O hydrogen bonds, and the C-H...Br contacts as dashed lines [H atoms not involved in these contacts have been omitted for clarity].

Isopropyltriphenylphosphonium bromide monohydrate

Crystal data

$C_{21}H_{22}P^+ \cdot Br^- \cdot H_2O$

$M_r = 403.28$

Orthorhombic, $P2_12_12_1$

Hall symbol: P 2ac 2ab

$a = 9.078 (5) \text{ \AA}$

$b = 13.043 (5) \text{ \AA}$

$c = 17.755 (5) \text{ \AA}$
 $V = 2102.3 (15) \text{ \AA}^3$
 $Z = 4$
 $F(000) = 832$
 $D_x = 1.274 \text{ Mg m}^{-3}$
 Mo $K\alpha$ radiation, $\lambda = 0.71069 \text{ \AA}$

Cell parameters from 3565 reflections
 $\theta = 2.5\text{--}23.5^\circ$
 $\mu = 2.04 \text{ mm}^{-1}$
 $T = 293 \text{ K}$
 ROD, colourless
 $0.20 \times 0.15 \times 0.13 \text{ mm}$

Data collection

Brucker APEXII CCD area-detector
 diffractometer
 Radiation source: fine-focus sealed tube
 Graphite monochromator
 phi and ω scans
 Absorption correction: multi-scan
 (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.668, T_{\max} = 0.688$

12177 measured reflections
 4250 independent reflections
 3331 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.034$
 $\theta_{\max} = 26.4^\circ, \theta_{\min} = 1.9^\circ$
 $h = -7 \rightarrow 11$
 $k = -16 \rightarrow 15$
 $l = -21 \rightarrow 21$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.049$
 $wR(F^2) = 0.140$
 $S = 1.03$
 4250 reflections
 228 parameters
 6 restraints
 Primary atom site location: structure-invariant
 direct methods
 Secondary atom site location: difference Fourier
 map

Hydrogen site location: inferred from
 neighbouring sites
 H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0853P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.82 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.28 \text{ e \AA}^{-3}$
 Absolute structure: Flack (1983), 1799 Friedel
 pairs
 Absolute structure parameter: 0.014 (13)

Special details

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

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

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

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
Br1	0.46181 (8)	0.58731 (5)	0.00805 (3)	0.0720 (2)	
C1	0.2804 (4)	0.5002 (3)	0.7564 (2)	0.0320 (8)	
C2	0.2503 (5)	0.4645 (3)	0.8289 (3)	0.0468 (11)	
H2	0.2722	0.5050	0.8705	0.056*	
C3	0.1873 (6)	0.3679 (4)	0.8386 (3)	0.0570 (13)	
H3	0.1690	0.3433	0.8868	0.068*	
C4	0.1527 (5)	0.3098 (3)	0.7778 (3)	0.0502 (13)	
H4	0.1075	0.2466	0.7850	0.060*	
C5	0.1833 (5)	0.3426 (3)	0.7054 (3)	0.0463 (12)	

H5	0.1607	0.3011	0.6644	0.056*	
C6	0.2478 (4)	0.4378 (3)	0.6943 (2)	0.0371 (9)	
H6	0.2694	0.4602	0.6458	0.045*	
C7	0.3207 (5)	0.6876 (3)	0.6626 (2)	0.0340 (9)	
C8	0.4105 (6)	0.7668 (3)	0.6355 (3)	0.0493 (12)	
H8	0.5032	0.7770	0.6564	0.059*	
C9	0.3631 (8)	0.8296 (4)	0.5783 (3)	0.0737 (18)	
H9	0.4228	0.8822	0.5604	0.088*	
C10	0.2238 (10)	0.8130 (5)	0.5476 (3)	0.082 (2)	
H10	0.1893	0.8561	0.5098	0.098*	
C11	0.1386 (7)	0.7354 (6)	0.5719 (3)	0.0754 (18)	
H11	0.0478	0.7239	0.5491	0.090*	
C12	0.1835 (5)	0.6726 (4)	0.6299 (3)	0.0498 (12)	
H12	0.1223	0.6205	0.6471	0.060*	
C13	0.5706 (4)	0.5878 (3)	0.7380 (2)	0.0312 (8)	
C14	0.6312 (5)	0.5673 (3)	0.6682 (3)	0.0430 (10)	
H14	0.5765	0.5766	0.6245	0.052*	
C15	0.7767 (5)	0.5321 (3)	0.6648 (3)	0.0538 (14)	
H15	0.8199	0.5187	0.6183	0.065*	
C16	0.8552 (5)	0.5173 (3)	0.7293 (4)	0.0567 (14)	
H16	0.9514	0.4930	0.7264	0.068*	
C17	0.7943 (5)	0.5379 (4)	0.7989 (3)	0.0569 (14)	
H17	0.8493	0.5277	0.8425	0.068*	
C18	0.6508 (4)	0.5737 (3)	0.8037 (3)	0.0446 (11)	
H18	0.6089	0.5880	0.8503	0.054*	
C19	0.3403 (5)	0.7009 (3)	0.8260 (2)	0.0352 (9)	
H19	0.3655	0.6618	0.8713	0.042*	
C20	0.4348 (6)	0.7974 (3)	0.8257 (3)	0.0538 (13)	
H20A	0.4054	0.8408	0.7847	0.081*	
H20B	0.5365	0.7789	0.8198	0.081*	
H20C	0.4220	0.8334	0.8724	0.081*	
C21	0.1779 (5)	0.7266 (4)	0.8303 (3)	0.0530 (12)	
H21A	0.1593	0.7669	0.8745	0.079*	
H21B	0.1216	0.6644	0.8327	0.079*	
H21C	0.1498	0.7648	0.7864	0.079*	
O1	0.4288 (9)	0.4940 (6)	0.5103 (4)	0.0650 (19)	0.50
H1O1	0.3405	0.4745	0.5027	0.097*	0.50
H2O1	0.4722	0.4469	0.4855	0.097*	0.50
O2	0.6608 (8)	0.3426 (5)	0.4736 (4)	0.0566 (18)	0.50
H1O2	0.6370	0.2797	0.4777	0.085*	0.50
H2O2	0.7522	0.3365	0.4835	0.085*	0.50
P1	0.37744 (10)	0.61951 (7)	0.74506 (6)	0.0269 (2)	

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Br1	0.0904 (5)	0.0823 (4)	0.0433 (3)	0.0170 (3)	-0.0098 (3)	0.0009 (3)
C1	0.0245 (18)	0.0290 (17)	0.042 (2)	0.0013 (15)	0.0014 (18)	0.0014 (17)

C2	0.054 (3)	0.039 (2)	0.047 (3)	-0.008 (2)	0.000 (2)	0.005 (2)
C3	0.062 (3)	0.048 (3)	0.061 (3)	-0.016 (3)	0.000 (3)	0.019 (2)
C4	0.035 (3)	0.032 (2)	0.084 (4)	-0.0057 (18)	-0.004 (2)	0.004 (2)
C5	0.035 (2)	0.035 (2)	0.068 (3)	-0.001 (2)	-0.006 (2)	-0.018 (2)
C6	0.033 (2)	0.037 (2)	0.041 (2)	0.0025 (18)	0.0034 (18)	-0.0061 (18)
C7	0.033 (2)	0.032 (2)	0.036 (2)	0.0060 (17)	0.0020 (18)	0.0009 (17)
C8	0.059 (3)	0.040 (2)	0.049 (3)	0.000 (2)	0.006 (2)	0.006 (2)
C9	0.111 (5)	0.050 (3)	0.061 (4)	0.020 (3)	0.026 (4)	0.021 (3)
C10	0.125 (6)	0.077 (4)	0.042 (3)	0.043 (4)	-0.001 (4)	0.021 (3)
C11	0.071 (4)	0.110 (5)	0.045 (3)	0.026 (4)	-0.017 (3)	0.006 (3)
C12	0.043 (3)	0.064 (3)	0.043 (3)	0.014 (2)	-0.006 (2)	0.005 (2)
C13	0.0247 (17)	0.0274 (17)	0.042 (2)	0.0007 (14)	0.0065 (16)	-0.0011 (17)
C14	0.041 (2)	0.036 (2)	0.053 (3)	0.000 (2)	0.009 (2)	-0.0011 (19)
C15	0.041 (3)	0.042 (3)	0.079 (4)	0.000 (2)	0.028 (3)	-0.007 (3)
C16	0.029 (2)	0.038 (2)	0.104 (5)	0.0034 (19)	0.016 (3)	0.006 (3)
C17	0.035 (3)	0.053 (3)	0.083 (4)	0.003 (2)	-0.007 (3)	0.016 (3)
C18	0.030 (2)	0.049 (3)	0.055 (3)	0.006 (2)	0.0002 (19)	0.004 (2)
C19	0.037 (2)	0.032 (2)	0.036 (2)	0.0005 (17)	0.0023 (18)	-0.0023 (16)
C20	0.053 (3)	0.037 (2)	0.071 (3)	-0.009 (2)	0.008 (3)	-0.010 (2)
C21	0.036 (2)	0.061 (3)	0.061 (3)	0.009 (2)	0.006 (2)	-0.012 (2)
O1	0.072 (5)	0.077 (5)	0.045 (4)	-0.015 (4)	-0.011 (4)	0.001 (3)
O2	0.062 (4)	0.059 (4)	0.049 (4)	-0.009 (3)	0.008 (3)	0.013 (3)
P1	0.0222 (4)	0.0273 (4)	0.0313 (5)	0.0001 (4)	0.0016 (4)	0.0009 (4)

Geometric parameters (Å, °)

C1—C2	1.395 (6)	C13—C18	1.387 (6)
C1—C6	1.402 (6)	C13—P1	1.806 (4)
C1—P1	1.800 (4)	C14—C15	1.400 (6)
C2—C3	1.394 (6)	C14—H14	0.9300
C2—H2	0.9300	C15—C16	1.363 (8)
C3—C4	1.355 (7)	C15—H15	0.9300
C3—H3	0.9300	C16—C17	1.381 (8)
C4—C5	1.384 (7)	C16—H16	0.9300
C4—H4	0.9300	C17—C18	1.386 (6)
C5—C6	1.387 (6)	C17—H17	0.9300
C5—H5	0.9300	C18—H18	0.9300
C6—H6	0.9300	C19—C21	1.514 (6)
C7—C12	1.388 (6)	C19—C20	1.523 (6)
C7—C8	1.400 (6)	C19—P1	1.818 (4)
C7—P1	1.789 (4)	C19—H19	0.9800
C8—C9	1.374 (7)	C20—H20A	0.9600
C8—H8	0.9300	C20—H20B	0.9600
C9—C10	1.394 (10)	C20—H20C	0.9600
C9—H9	0.9300	C21—H21A	0.9600
C10—C11	1.345 (10)	C21—H21B	0.9600
C10—H10	0.9300	C21—H21C	0.9600
C11—C12	1.377 (7)	O1—H1O1	0.8522

C11—H11	0.9300	O1—H2O1	0.8527
C12—H12	0.9300	O2—H1O2	0.8515
C13—C14	1.382 (6)	O2—H2O2	0.8512
C2—C1—C6	119.4 (4)	C13—C14—H14	120.7
C2—C1—P1	119.2 (3)	C15—C14—H14	120.7
C6—C1—P1	121.1 (3)	C16—C15—C14	120.2 (5)
C3—C2—C1	119.7 (5)	C16—C15—H15	119.9
C3—C2—H2	120.2	C14—C15—H15	119.9
C1—C2—H2	120.2	C15—C16—C17	121.1 (4)
C4—C3—C2	120.2 (5)	C15—C16—H16	119.5
C4—C3—H3	119.9	C17—C16—H16	119.5
C2—C3—H3	119.9	C16—C17—C18	119.7 (5)
C3—C4—C5	121.4 (4)	C16—C17—H17	120.1
C3—C4—H4	119.3	C18—C17—H17	120.1
C5—C4—H4	119.3	C17—C18—C13	119.2 (5)
C4—C5—C6	119.5 (4)	C17—C18—H18	120.4
C4—C5—H5	120.2	C13—C18—H18	120.4
C6—C5—H5	120.2	C21—C19—C20	111.4 (4)
C5—C6—C1	119.8 (4)	C21—C19—P1	110.5 (3)
C5—C6—H6	120.1	C20—C19—P1	112.1 (3)
C1—C6—H6	120.1	C21—C19—H19	107.6
C12—C7—C8	118.9 (4)	C20—C19—H19	107.6
C12—C7—P1	122.0 (3)	P1—C19—H19	107.6
C8—C7—P1	118.7 (3)	C19—C20—H20A	109.5
C9—C8—C7	120.8 (5)	C19—C20—H20B	109.5
C9—C8—H8	119.6	H20A—C20—H20B	109.5
C7—C8—H8	119.6	C19—C20—H20C	109.5
C8—C9—C10	118.7 (6)	H20A—C20—H20C	109.5
C8—C9—H9	120.7	H20B—C20—H20C	109.5
C10—C9—H9	120.7	C19—C21—H21A	109.5
C11—C10—C9	120.9 (5)	C19—C21—H21B	109.5
C11—C10—H10	119.6	H21A—C21—H21B	109.5
C9—C10—H10	119.6	C19—C21—H21C	109.5
C10—C11—C12	121.1 (6)	H21A—C21—H21C	109.5
C10—C11—H11	119.4	H21B—C21—H21C	109.5
C12—C11—H11	119.4	H1O1—O1—H2O1	98.0
C11—C12—C7	119.6 (5)	H1O2—O2—H2O2	98.0
C11—C12—H12	120.2	C7—P1—C1	112.38 (19)
C7—C12—H12	120.2	C7—P1—C13	109.67 (19)
C14—C13—C18	121.2 (4)	C1—P1—C13	106.56 (17)
C14—C13—P1	119.5 (3)	C7—P1—C19	107.68 (18)
C18—C13—P1	118.8 (3)	C1—P1—C19	108.99 (19)
C13—C14—C15	118.6 (5)	C13—P1—C19	111.62 (19)

Hydrogen-bond geometry (Å, °)

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
O1—H1O1 \cdots Br1 ⁱ	0.85	2.86	3.701 (8)	169
O1—H2O1 \cdots O2	0.85	2.20	2.960 (11)	149
O2—H1O2 \cdots Br1 ⁱⁱ	0.85	2.68	3.526 (7)	175
O2—H2O2 \cdots Br1 ⁱⁱⁱ	0.85	2.81	3.598 (8)	154
C19—H19 \cdots Br1 ^{iv}	0.98	2.76	3.723 (4)	169

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