

Isobutyltriphenylphosphonium bromide

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At 102 K, the isobutyl group of the title compound, $C_{22}H_{24}P^+ \cdot Br^-$, is in the extended conformation. However, the isopropyl moiety is in the eclipsed conformation with respect to the H atoms on the C atom attached to the P atom. The benzene rings are in the propeller configuration usually found in this family of triphenylphosphonium compounds.

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Key indicators

Single-crystal X-ray study

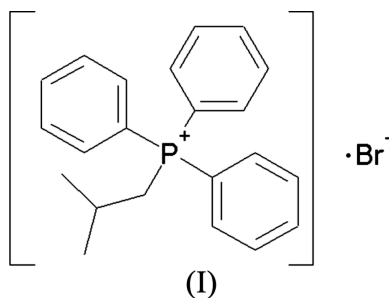
 $T = 102$ KMean $\sigma(C-C) = 0.004$ Å R factor = 0.039 wR factor = 0.112

Data-to-parameter ratio = 16.4

For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

Comment

The title compound, (I), is the seventh crystal structure in a series of alkyl-substituted triphenylphosphonium bromide compounds from this laboratory (Czerwinski, 1986, 2004; Ponnuswamy & Czerwinski, 1986; Czerwinski & Ponnuswamy, 1988*a,b*, 1989). The atom labelling is consistent with the previously reported structures.



Experimental

The title compound was obtained from Lancaster Synthesis Ltd. Suitable crystals were grown by evaporation of a methanol solution at 294 K.

Crystal data

 $C_{22}H_{24}P^+ \cdot Br^-$ $M_r = 399.29$ Monoclinic, $P2_1/c$ $a = 11.3309$ (3) Å $b = 9.8033$ (2) Å $c = 17.7075$ (3) Å $\beta = 106.487$ (1)° $V = 1886.08$ (7) Å³ $Z = 4$ $D_x = 1.406$ Mg m⁻³Cu $K\alpha$ radiation

Cell parameters from 19 309

reflections

 $\theta = 4.1$ – 70.1 ° $\mu = 3.75$ mm⁻¹ $T = 102$ (2) K

Prism, colorless

 $0.30 \times 0.20 \times 0.10$ mm

Data collection

Bruker AXS diffractometer

 ω scansAbsorption correction: multi-scan
(Blessing, 1995) $T_{\min} = 0.37$, $T_{\max} = 0.685$

18 578 measured reflections

3555 independent reflections

2997 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.089$ $\theta_{\text{max}} = 70.3$ ° $h = -13 \rightarrow 11$ $k = -11 \rightarrow 11$ $l = -21 \rightarrow 21$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.039$
 $wR(F^2) = 0.112$
 $S = 1.13$
 3555 reflections
 217 parameters

H atoms treated by a mixture of independent and constrained refinement
 $w = 1/[\sigma^2(F_o^2) + (0.0659P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.035$
 $\Delta\rho_{\max} = 0.67 \text{ e } \text{Å}^{-3}$
 $\Delta\rho_{\min} = -0.68 \text{ e } \text{Å}^{-3}$

Table 1
 Selected geometric parameters (Å, °) for (I).

C11—C12	1.541 (4)	C21—P	1.801 (3)
C11—P	1.791 (3)	C31—P	1.799 (2)
C12—C13	1.526 (3)	C41—P	1.798 (3)
C12—C14	1.539 (4)		
C12—C11—P	116.77 (18)	C46—C41—P	122.40 (19)
C13—C12—C14	110.0 (2)	C42—C41—P	117.4 (2)
C13—C12—C11	110.9 (2)	C11—P—C41	112.00 (12)
C14—C12—C11	110.2 (2)	C11—P—C31	111.26 (12)
C22—C21—P	117.8 (2)	C41—P—C31	109.71 (12)
C26—C21—P	122.20 (19)	C11—P—C21	110.30 (12)
C32—C31—P	121.77 (19)	C41—P—C21	107.68 (12)
C36—C31—P	118.6 (2)	C31—P—C21	105.65 (12)
P—C11—C12—C13	−102.2 (2)	C42—C41—P—C11	56.8 (2)
P—C11—C12—C14	135.71 (19)	C32—C31—P—C11	9.5 (3)
C12—C11—P—C41	52.7 (2)	C36—C31—P—C11	−168.3 (2)
C12—C11—P—C31	−70.5 (2)	C22—C21—P—C11	56.3 (2)
C12—C11—P—C21	172.64 (18)	C26—C21—P—C11	−128.0 (2)
C46—C41—P—C11	−126.9 (2)		

Following anisotropic refinement, all H atoms were placed in geometrically idealized positions. The C—H distances were constrained to ride on their parent atoms within the range of 0.95–1.00 Å and with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$.

Data collection: *PROTEUM* (Bruker, 2002); cell refinement: *SAINTE-Plus* (Bruker, 2001); data reduction: *SAINTE-Plus*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

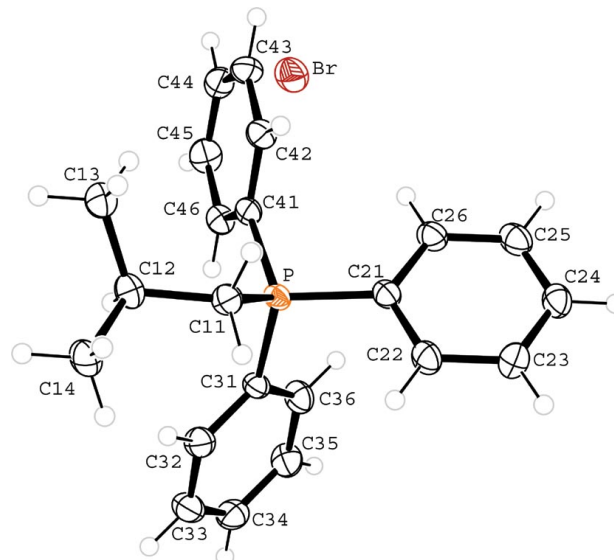


Figure 1
 View of the title compound, showing the atom-labelling scheme (50% probability displacement ellipsoids). H atoms are represented by small spheres.

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