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Key indicators

Single-crystal X-ray study T = 120 KMean $\sigma(C-C) = 0.003 \text{ Å}$ R factor = 0.026 wR factor = 0.057 Data-to-parameter ratio = 16.2

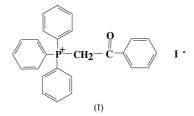
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. (Benzoylmethyl)triphenylphosphonium iodide: supramolecular zigzag chains *via* C—H···O hydrogen bonding stabilized by C—H···I interactions

The title phosphonium salt, $C_{26}H_{22}OP^+ \cdot I^-$, isomorphous with (benzoylmethyl)triphenylphosphonium nitrate, exhibits both intramolecular $C-H \cdot \cdot \cdot I$ and intermolecular $C-H \cdot \cdot \cdot O$ interactions. The latter interaction leads to supramolecular parallel zigzag chains along the *a* axis.

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Comment

Phosphonium keto ylides react with metal salts to form not only ylide complexes but also phosphonium salts (Antipin & Struchkov, 1984; Baby Marivatra et al., 2002) and phosphonium metalates (Albanese et al., 1989; Bart et al., 1982). The action of trimethyltin chloride on (benzoylmethylene)triphenylphosphorane led to the first O-coordinated ketoylide complex (Buckel *et al.*, 1972). The reaction of SnI_4 with the same ylide was used to prepare a six-coordinate complex, containing the ylide. However, the above reaction in benzene yielded the product of composition [(C₆H₅)₃PCH₂CO- $Ph_{2}[SnI_{6}]$, as revealed by its elemental analysis (analysis found: C 37.98, H 2.78%; calculated: C 38.01, H 2.70%). The product was recrystallized from methanol. Its ¹H NMR in CDCl₃ indicates that it is a phosphonium salt, different from the initial phosphonium metalate. The investigation reported herein has been undertaken to assign the molecular and supramolecular structure of the title compound, (I).



Compound (I) is formed by the C protonation of the parent ylide, and its molecular structure is unexceptional. It is confirmed that compound (I) is an ion pair (Fig. 1), with a distance of 4.6231 (7) Å between the P^+ and I^- centres. The metrical parameters in (I) resemble those of the isomorphous nitrate salt (*Pbca* with Z = 8; Baby Mariyatra *et al.*, 2002), but differ from the parent ylide (Kalyanasundari et al., 1994). The crystal structure can best be described as a zigzag chain formed by $C-H \cdots O$ hydrogen-bonding interactions (Table 2 and Fig. 2). The H1-C1-C2-O1 torsion angle $[145.4 (14)^{\circ}]$ indicates that atoms H1 and O1 are considerably distorted from an antiperiplanar arrangement, as expected for the formation of a zigzag chain. The I⁻ ions are embedded in the spaces between the chains and contribute to the threedimensional arrangement via three C-H···I interactions (Table 2 and Fig. 3). The C···I distances [3.696 (2), 3.802 (2)

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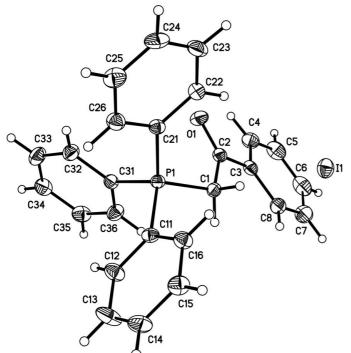
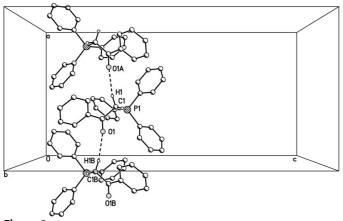


Figure 1

The molecular structure of the title compound, (I), showing 50% probability displacement ellipsoids.





View of the chain along the *a* axis, due to $C-H \cdots O$ interactions (dashed lines).

and 3.983 (2) Å] are comparable to those observed in the fulleride salt $[(C_6H_5)_4P]_2(C_{60})I_{0.35}$ (C···I⁻ = 3.67 Å; Penicaud et al., 1993) and those observed in (α -ferrocenylphenacyl)triphenylphosphonium iodide $[C \cdot \cdot \cdot I = 3.836(7), 3.932(7)]$ and 3.995 (7) Å; Ahmed et al., 1996].

The present results indicate that the phosphonium salt is present in the C-protonated form both in solution and in the solid state. The ylidic C atom is involved in hydrogen-bonding interactions, as observed in many similar structures (Yufit et al., 2000; Baby Mariyatra et al., 2002). We attribute the predominance of the C-protonated form to the considerable carbanionic character of the ylidic C atom in spite of resonance stabilization.

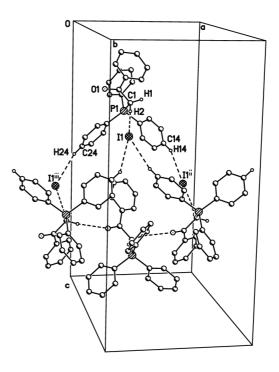


Figure 3

Diagram showing both the C-H···I and C-H···O interactions (dashed lines). The symmetry codes are as in Table 2.

Experimental

The title phosphonium salt, (I), was obtained when the product of the reaction between stannic iodide and the phosphonium ylide, (benzoylmethylene)triphenylphosphorane, was crystallized from dry methanol.

Crystal data

$C_{26}H_{22}OP^+ \cdot I^-$	Mo Ka radiation	
$M_r = 508.31$	Cell parameters from	
Orthorhombic, Pbca	reflections	
a = 10.870 (2) Å	$\theta = 10.2 - 23.6^{\circ}$	
b = 18.932 (2) Å	$\mu = 1.49 \text{ mm}^{-1}$	
c = 22.135 (3) Å	T = 120 (2) K	
$V = 4555.2 (11) \text{ Å}^3$	Block, light yellow	
Z = 8	$0.46 \times 0.32 \times 0.28$ m	
$D_x = 1.482 \text{ Mg m}^{-3}$		

Data collection

Bruker SMART 1K CCD diffractometer ω scans Absorption correction: by integration (XPREP in SHELXTL; Bruker, 1998) $T_{\rm min}=0.555,\;T_{\rm max}=0.684$ 29262 measured reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.026$ wR(F²) = 0.057 S = 1.055663 reflections 350 parameters All H-atom parameters refined

n 805 ım

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5663 independent reflections
4873 reflections with I > 2\sigma(I)
R_{\rm int}=0.030
\theta_{\rm max} = 28.3^{\circ}
h = -13 \rightarrow 14
k = -25 \rightarrow 24
l=-29\rightarrow23
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Table 1	
Selected geometric parameters (Å, °).	

P1-C1 O1-C2	1.807 (2) 1.225 (2)	C1-C2	1.518 (3)
C31-P1-C1 C2-C1-P1 O1-C2-C3	108.33 (9) 113.28 (14) 121.92 (18)	01-C2-C1 C3-C2-C1	119.90 (18) 118.17 (17)
P1-C1-C2-O1	22.6 (2)	P1-C1-C2-C3	-158.66 (14)

Table 2

Hydrogen-bonding geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$\begin{array}{c} C1 - H1 \cdots O1^{i} \\ C1 - H2 \cdots I1 \\ C14 - H14 \cdots I1^{ii} \\ C24 - H24 \cdots I1^{iii} \end{array}$	0.96 (2)	2.50 (2)	3.445 (3)	165.2 (18)
	0.92 (3)	2.78 (3)	3.696 (2)	177 (2)
	0.93 (3)	3.03 (3)	3.802 (2)	141 (2)
	0.92 (3)	3.09 (3)	3.983 (2)	164 (2)

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

All the H atoms were located from difference Fourier maps and their positional and $U_{\rm iso}$ parameters were refined. The C–H bond distances range from 0.91 (3) to 0.98 (2) Å.

Data collection: *SMART-NT* (Bruker, 1998); cell refinement: *SMART-NT*; data reduction: *SAINT-NT* (Bruker, 1998); program(s)

used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997) and *PLATON*97 (Spek, 1997); software used to prepare material for publication: *SHELXL*97.

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