

**O-Phenyl (triphenylphosphonio-
 methyl)phosphonate phenol solvate:
 supramolecular structure generated
 by O—H···O, C—H···O and
 C—H··· π (arene) hydrogen bonds**

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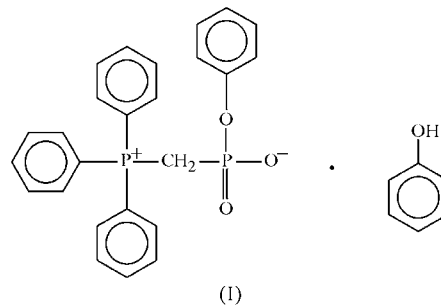
The title compound, C₂₅H₂₂O₃P₂·C₆H₆O, has a zwitterionic betaine-like structure and crystallizes as a phenol solvate. The two molecular components are held together by an almost linear intermolecular O—H···O hydrogen bond. The structure also contains three weak C—H···O and two C—H··· π (arene) interactions.

Comment

Betaine (*N,N,N*-trimethylglycine) acts as an important biochemical donor of the methyl group for tetrahydrofolic acid (*N*⁵,*N*¹⁰-methylenetetrahydrofolate synthesis) and homocysteine acid (methionine synthesis) (Stryer, 1995). The phosphonic analogue of betaine – trimethylammonium-methanophosphonic acid – was first synthesized over five decades ago (Medved & Kabachnik, 1951; Kabachnik & Medved, 1953; Myers & Jibril, 1957); more than a decade ago, its biological activity became recognized (Abdel-Ghany *et al.*, 1993). However, to our knowledge, no example of a betaine analogue containing two P atoms (phosphoniummethylene-phosphonic acid or its ester) has been described in the literature. We present here the synthesis and crystal and molecular structures of the first phosphoniophosphonic analogue of betaine, namely *O*-phenyl (triphenylphosphoniomethyl)phosphonate as a phenol solvate, (I).

The first example of phosphoniummethylene-phosphonic acid was synthesized in the form of the *O*-phenyl ester by controlled hydrolysis of [(diphenoxyphosphinyl)methylidene]triphenylphosphorane, (II), and obtained in the crystalline form as the phenol solvate, (I). Its structure was

tentatively assigned by MS, NMR and elemental analysis; however, an unequivocal confirmation was obtained from crystallographic analysis.



The molecular structure of (I), with the atom-numbering scheme, is shown in Fig. 1. The phosphonate molecule exists as a zwitterion. The positive charge is located on atom P1, while the negative charge is attributed formally to atom O2. Taking into account the P2—O2 and P2—O3 bond lengths (Table 1), partial delocalization of the charge between the two O atoms within the O2···P2···O3 fragment can be proposed. The P2—O3 bond is slightly shorter than the P2—O2 bond; however, both distances are longer than the P=O linkage within (C,N)(O)₂—P=O systems [1.457 (9) Å; Allen *et al.*, 1987] and close to the mean value of 1.483 (8) Å for (C—O)₂P(=O)₂[−] systems, indicating bond delocalization (Allen *et al.*, 1987).

In the molecule of (I), two P atoms are joined through a Csp³ atom. The P1—C1 and P2—C1 distances differ from each other significantly (Table 1). The coordination around P1 is nearly tetrahedral, with angles ranging from 106.98 (7) to 114.31 (7)°, whereas within the phosphonate group, the geometry around atom P2 indicates a considerable distortion of the tetrahedron, especially in the O2—P2—O3 and O1—P2—C1 bond angles (Table 1). The P⁺—C(phenyl) bond lengths are very close to the mean distance of 1.793 (10) Å

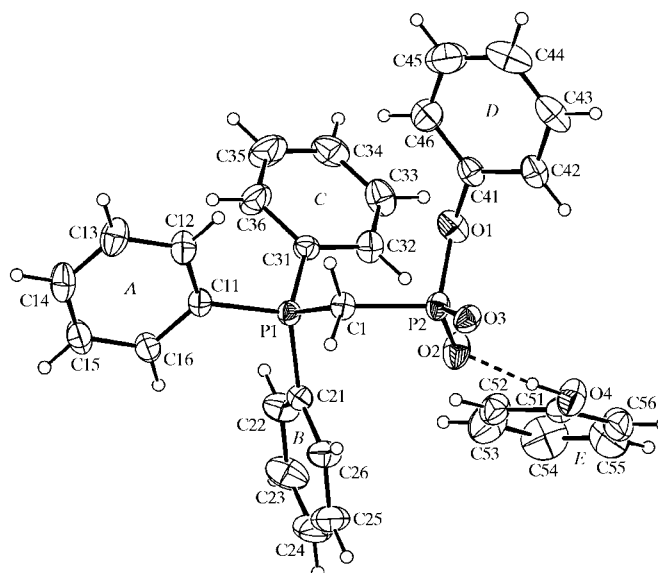


Figure 1
 The structure of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level and H atoms are shown as small spheres of arbitrary radii.

found for 118 $\text{Ph}_3\text{P}^+ - \text{Csp}^3\text{H}_2$ fragments ($3 \times 118 = 354$ values) in the Cambridge Structural Database (Version 5.27; Allen, 2002); the minimum and maximum distances are 1.752 (3) and 1.854 (3) Å, respectively.

All benzene rings, including the phenol ring, are planar; a maximum deviation of -0.010 (2) Å was observed for atom C12 from the least-squares plane of ring A. (The labelling scheme, A–E, applied for phenyl rings, is introduced in Fig. 1.) The C–C bond lengths and angles within these rings are typical for aromatic systems [1.384 (13) Å; Allen *et al.*, 1987], hence no detailed discussion is needed. The dihedral angles between the mean planes of the rings are summarized in Table 3. This spatial arrangement enables the formation of C–H \cdots O and C–H \cdots π (arene) interactions; however, aromatic π – π stacking interactions are absent in the structure.

Two molecular components are linked within the selected asymmetric unit by an almost linear O4–H4 \cdots O2 intermolecular hydrogen bond (Fig. 1 and Table 2).

There are also three short intermolecular C–H \cdots O contacts (Table 2), which on the basis of observed H \cdots O and C \cdots O distances can be regarded as weak hydrogen-bonding interactions (Taylor & Kennard, 1982). Atoms C1 and C12 act as hydrogen-bond donors to atom O3ⁱ [symmetry code: (i)

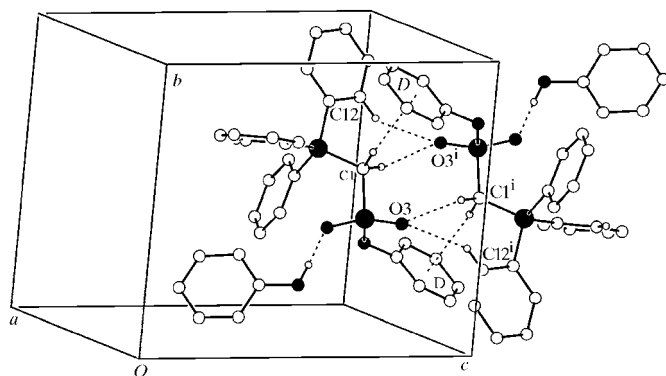


Figure 2
Part of the crystal structure of (I), showing the formation of a centrosymmetric dimer. For the sake of clarity, C-bound H atoms that are not involved in the motif shown have been omitted. [Symmetry code: (i) $-x + 1, -y + 1, -z + 2$.]

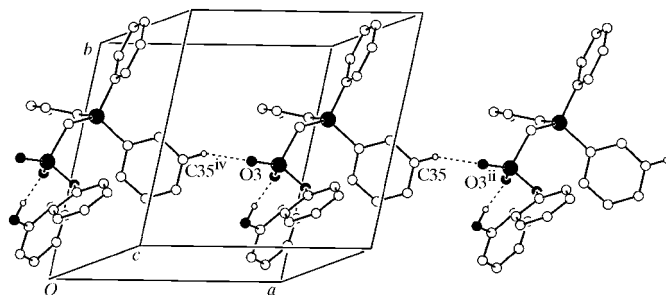


Figure 3
Part of the crystal structure of (I), showing the formation of an infinite C–H \cdots O chain running parallel to the [100] direction. For the sake of clarity, C-bound H atoms that are not involved in the motif shown have been omitted. [Symmetry codes: (ii) $x + 1, y, z$; (iv) $x - 1, y, z$.]

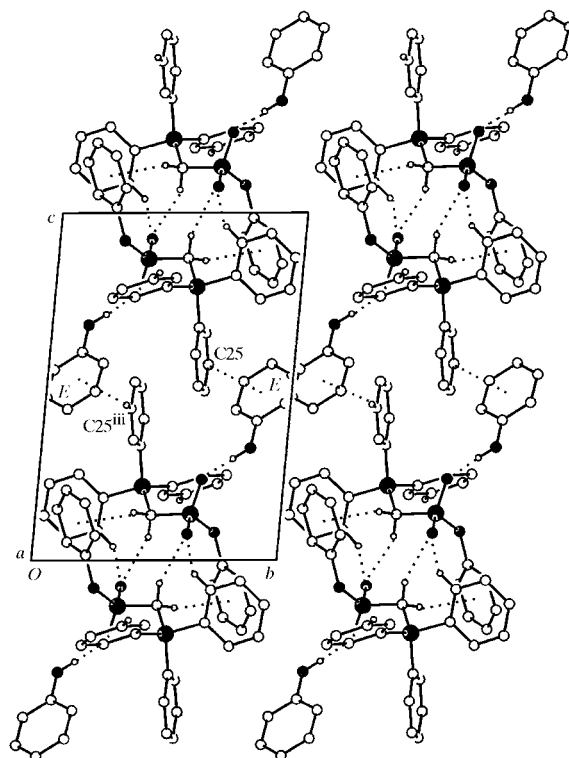


Figure 4
Part of the crystal structure of (I), showing (010) sheets generated by C–H \cdots π interactions between chains of dimers. For the sake of clarity, C-bound H atoms that are not involved in the motif shown have been omitted. [Symmetry code: (iii) $-x + 1, -y + 1, -z + 1$.]

$-x + 1, -y + 1, -z + 2$], forming a centrosymmetric dimer. The former of these hydrogen bonds generates a graph-set motif of $R_2^2(8)$, while the second generates an $R_2^2(14)$ ring (Bernstein *et al.*, 1995) (Fig. 2). Such a dimer is additionally stabilized by C1–H1A \cdots π (ring D) hydrogen bonds.

Fig. 3 shows another hydrogen-bonded motif that plays an important role in the supramolecular structure of (I). Atom C35, *via* atom H35, acts as a hydrogen-bond donor towards atom O3ⁱⁱ [symmetry code: (ii) $x + 1, y, z$], thereby producing an infinite $C(8)$ chain running parallel to the [100] direction.

In summary, the combination of all three weak C–H \cdots O interactions produces infinite chains of centrosymmetric dimers along the *a* axis. The aforementioned chains are linked into (010) sheets by C25–H25 \cdots π ⁱⁱⁱ interactions [ring E; symmetry code: (iii) $-x + 1, -y + 1, -z + 1$; Fig. 4]. Every sheet passes through the unit cell in the domain $-0.07 < y < 1.07$. There are no direction-specific interactions between adjacent sheets.

Experimental

The title compound, (I), was synthesized from [(diphenoxyphosphinyl)methylidene]triphenylphosphorane, (II), which was prepared exactly as described in our previous paper (Chęcińska *et al.*, 2003). For (II): ³¹P NMR (CD₃CN): δ 20.98 (*d*, $\text{Ph}_3\text{P}=\text{C}$), 28.03 [*d*, $(\text{PhO})_2\text{P}(\text{O})^-$, $^2J_{\text{PP}} = 46.4$ Hz]. The ylide (II) (1.08 g, 1 mmol) was dissolved in 20 ml of acetonitrile containing 2% water. The solution was kept for 48 h at 323 K in a closed flask and evaporated. The

residue was crystallized from ethyl acetate, yielding 0.61 g (58%) of phenol-solvated betaine (I) in the form of colorless crystals (m.p. 476–479 K). ^{31}P NMR (CDCl_3): δ -1.64 [d , (PhO) $\text{P}(\text{O})\text{O}^-$], 212.36 (d , Ph_3P^+ , $^2J_{\text{PP}} = 10.1$ Hz). ^1H NMR (CDCl_3): δ 3.565 (dd , 2H, $\text{P}-\text{CH}_2-\text{P}$, $^2J_{\text{HP}} = 15.7$ and 17.1 Hz), 6.7–7.2 (m , 10H, PhO H atoms), 7.45–7.8 (m , 15H, Ph_3P H atoms), 9.44 ($br\ s$, 1H, OH). MS FAB (Cs^+ , negative ions): m/z 431 (16%), 355 (100%), 278 (8%), 261 (28%). Elemental analysis found: C 70.49, H 5.25, P 11.58%; calculated for $\text{C}_{31}\text{H}_{28}\text{O}_4\text{P}_2$: 70.72, H 5.36, P 11.77%.

Crystal data

$\text{C}_{25}\text{H}_{22}\text{O}_3\text{P}_2 \cdot \text{C}_6\text{H}_6\text{O}$ $\gamma = 78.35$ (2) $^\circ$
 $M_r = 526.47$ $V = 1374.9$ (5) \AA^3
 Triclinic, $P\bar{1}$ $Z = 2$
 $a = 9.793$ (2) \AA Mo $K\alpha$ radiation
 $b = 10.170$ (2) \AA $\mu = 0.19$ mm^{-1}
 $c = 14.374$ (2) \AA $T = 293$ (2) K
 $\alpha = 82.78$ (2) $^\circ$ $0.5 \times 0.2 \times 0.1$ mm
 $\beta = 80.07$ (2) $^\circ$

Data collection

Rigaku AFC-5S diffractometer $R_{\text{int}} = 0.013$
 6644 measured reflections 3 standard reflections
 6293 independent reflections every 150 reflections
 4137 reflections with $I > 2\sigma(I)$ intensity decay: 1.6%

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.033$ H atoms treated by a mixture of
 $wR(F^2) = 0.082$ independent and constrained
 $S = 1.02$ refinement
 6293 reflections $\Delta\rho_{\text{max}} = 0.27$ e \AA^{-3}
 338 parameters $\Delta\rho_{\text{min}} = -0.24$ e \AA^{-3}

Table 1

Selected geometric parameters (\AA , $^\circ$).

P1–C1	1.7855 (14)	P2–O2	1.4815 (11)
P1–C21	1.7923 (14)	P2–O1	1.6189 (12)
P1–C11	1.7991 (15)	P2–C1	1.8202 (15)
P1–C31	1.8009 (15)	O1–C41	1.3809 (18)
P2–O3	1.4695 (10)	O4–C51	1.3523 (19)
C1–P1–C21	110.26 (7)	O3–P2–O1	111.10 (6)
C1–P1–C11	106.98 (7)	O2–P2–O1	105.09 (7)
C21–P1–C11	109.54 (7)	O3–P2–C1	107.50 (7)
C1–P1–C31	114.31 (7)	O2–P2–C1	107.85 (7)
C21–P1–C31	107.97 (7)	O1–P2–C1	101.60 (7)
C11–P1–C31	107.68 (7)	C41–O1–P2	124.97 (9)
O3–P2–O2	121.84 (7)	P1–C1–P2	120.06 (8)

Table 2

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

CgD and CgE are the centroids of rings D and E , respectively.

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{O4}-\text{H4}\cdots\text{O2}$	0.89 (2)	1.72 (2)	2.601 (2)	175 (2)
$\text{C1}-\text{H1B}\cdots\text{O3}^i$	0.97	2.30	3.142 (2)	145
$\text{C12}-\text{H12}\cdots\text{O3}^i$	0.93	2.57	3.269 (2)	133
$\text{C35}-\text{H35}\cdots\text{O3}^ii$	0.93	2.36	3.200 (3)	150
$\text{C1}-\text{H1A}\cdots\text{CgD}^i$	0.93	2.79	3.727 (2)	161
$\text{C25}-\text{H25}\cdots\text{CgE}^{iii}$	0.93	2.90	3.667 (2)	141

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

Table 3

The dihedral angles between the mean planes for the phenyl rings ($^\circ$).

Symbols $A-E$ refer to the phenyl rings as defined in Fig. 1.

Ring/ring	Dihedral angle
A/B	78.1 (1)
A/C	77.4 (1)
A/D	36.1 (1)
A/E	44.1 (1)
B/C	72.8 (1)
B/D	43.8 (1)
B/E	68.5 (1)
C/D	83.1 (1)
C/E	58.6 (1)
D/E	52.8 (1)

Atom H4, involved in the $\text{O4}-\text{H4}\cdots\text{O2}$ hydrogen bond, was located in a difference map and refined isotropically. All C-bound H atoms of methylene and phenyl groups were positioned geometrically ($\text{C}-\text{H} = 0.97$ and 0.93 \AA , respectively) and constrained to ride on their parent atoms [$U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$].

Data collection: *MSC/AFD Diffractometer Control Software* (Molecular Structure Corporation, 1989); cell refinement: *MSC/AFD Diffractometer Control Software*; data reduction: *TEXSAN* (Molecular Structure Corporation, 1989); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2003); software used to prepare material for publication: *PLATON*.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: SK3145). Services for accessing these data are described at the back of the journal.

References

Abdel-Ghany, Y. S., Ihnat, M. A., Miller, D. D., Kunin, C. M. & Tong, H. H. (1993). *J. Med. Chem.* **36**, 784–789.
 Allen, F. H. (2002). *Acta Cryst.* **B58**, 380–388.
 Allen, F. H., Kennard, O., Watson, D. G., Brammer, L., Orpen, A. G. & Taylor, R. (1987). *J. Chem. Soc. Perkin Trans. 2*, pp. S1–S3.
 Bernstein, J., Davis, R. E., Shimon, L. & Chang, N.-L. (1995). *Angew. Chem. Int. Ed. Engl.* **34**, 1555–1573.
 Chęcińska, L., Kudzin, Z. H., Malecka, M., Nazarski, R. B. & Okruszek, A. (2003). *Tetrahedron*, **59**, 7681–7693.
 Kabachnik, M. I. & Medved, T. Ya. (1953). *Izv. Akad. Nauk SSSR Ser. Khim.* pp. 1126–1127.
 Medved, T. Ya. & Kabachnik, M. I. (1951). *Izv. Akad. Nauk SSSR Ser. Khim.* pp. 620–623.
 Molecular Structure Corporation (1989). *MSC/AFD Diffractometer Control Software* and *TEXSAN* (Version 5.0). MSC, The Woodlands, Texas, USA.
 Myers, T. C. & Jibril, A. O. (1957). *J. Org. Chem.* **22**, 180–182.
 Sheldrick, G. M. (1997). *SHELXS97* and *SHELXL97*. University of Göttingen, Germany.
 Spek, A. L. (2003). *J. Appl. Cryst.* **36**, 7–13.
 Stryer, L. (1995). In *Biochemistry*. New York: W. H. Freeman and Company.
 Taylor, R. & Kennard, O. (1982). *J. Am. Chem. Soc.* **104**, 5063–5070.