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Key indicators

Single-crystal X-ray study
 $T = 293\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.002\text{ \AA}$
 R factor = 0.041
 wR factor = 0.109
Data-to-parameter ratio = 16.0For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

Tetraphenylphosphonium 2,4,5-tricarboxybenzoate

The title compound, $\text{C}_{24}\text{H}_{20}\text{P}^+\cdot\text{C}_{10}\text{H}_5\text{O}_8^-$, is a salt of singly-deprotonated 1,2,4,5-benzenetetracarboxylic acid (pyromellitic acid). The crystal packing is stabilized by intermolecular hydrogen bonds involving the anions, in addition to normal Coulombic forces.

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Comment

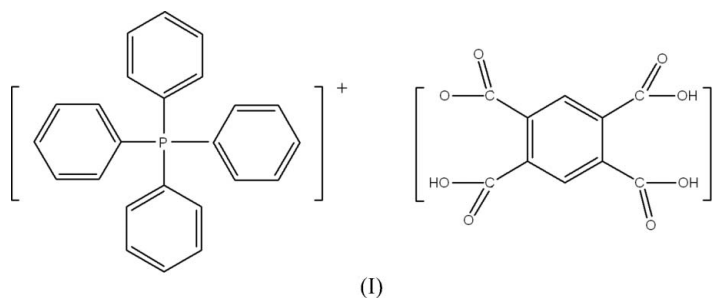
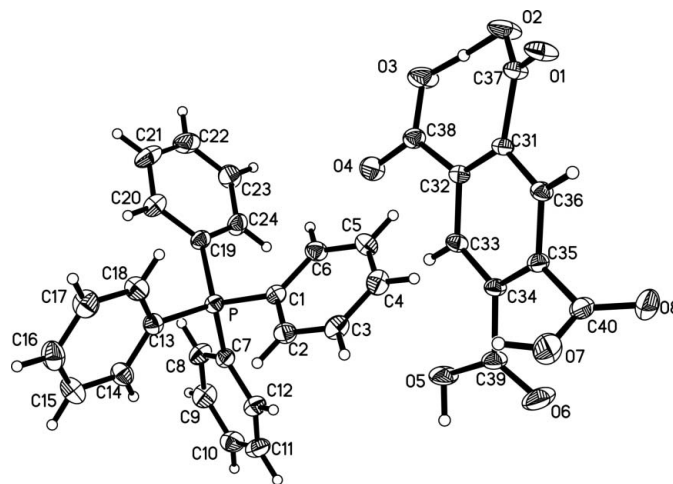
The title compound, (I), is an air-stable salt at ambient temperature. Its structure contains discrete Ph_4P^+ cations and H_3BTC^- anions (Fig. 1; $\text{H}_4\text{BTC} = 1,2,4,5$ -benzenetetracarboxylic acid or pyromellitic acid). All bond distances and angles in (I) are normal. H_3BTC^- anions are linked together by intermolecular hydrogen bonds between carboxyl and carboxylate groups (Fig. 2 and Table 2), generating a chain of centrosymmetric dimers along the a axis. There is also a strong intramolecular $\text{O}-\text{H}\cdots\text{O}$ hydrogen bond between one carboxyl group and the adjacent carboxylate group. The main interaction of the cations with the anions is normal Coulombic attraction.

Figure 1

The asymmetric unit of compound (I), with 30% probability displacement ellipsoids. The almost symmetric strong intramolecular hydrogen bond is shown with two full bonds.

Experimental

The title compound was obtained as hexagonal-prismatic colorless crystals in about 50% yield from the hydrothermal reaction of 1,2,4,5-benzenetetracarboxylic acid (0.5 mmol), tetraphenylphosphonium chloride (0.53 mmol) and water (5 ml) heated at 473 K for 5 d, then cooled at a rate of 2 K h⁻¹ to 423 K, and finally left to cool naturally to ambient temperature. The crystal structure is consistent with the IR spectrum of (I).

Crystal data

| | |
|--|---|
| C ₂₄ H ₂₀ P ⁺ ·C ₁₀ H ₅ O ₈ ⁻ | $V = 1412.3 (8) \text{ \AA}^3$ |
| $M_r = 592.51$ | $Z = 2$ |
| Triclinic, $P\bar{1}$ | $D_x = 1.393 \text{ Mg m}^{-3}$ |
| $a = 9.715 (3) \text{ \AA}$ | Mo $K\alpha$ radiation |
| $b = 10.920 (3) \text{ \AA}$ | $\mu = 0.15 \text{ mm}^{-1}$ |
| $c = 14.268 (5) \text{ \AA}$ | $T = 293 (2) \text{ K}$ |
| $\alpha = 90.218 (4)^\circ$ | Prism, colorless |
| $\beta = 106.435 (3)^\circ$ | $0.60 \times 0.35 \times 0.25 \text{ mm}$ |
| $\gamma = 102.776 (4)^\circ$ | |

Data collection

| | |
|--|--|
| Rigaku Mercury70 diffractometer | 11070 measured reflections |
| ω scans | 6387 independent reflections |
| Absorption correction: multi-scan (<i>CrystalClear</i> ; Rigaku, 2000) | 5443 reflections with $I > 2\sigma(I)$ |
| $T_{\min} = 0.897$, $T_{\max} = 0.963$ | $R_{\text{int}} = 0.013$ |
| | $\theta_{\text{max}} = 27.5^\circ$ |

Refinement

| | |
|--|--|
| Refinement on F^2 | $w = 1/[\sigma^2(F_o^2) + (0.0567P)^2 + 0.383P]$ |
| $R[F^2 > 2\sigma(F^2)] = 0.041$ | where $P = (F_o^2 + 2F_c^2)/3$ |
| $wR(F^2) = 0.109$ | $(\Delta/\sigma)_{\text{max}} < 0.001$ |
| $S = 1.01$ | $\Delta\rho_{\text{max}} = 0.29 \text{ e \AA}^{-3}$ |
| 6387 reflections | $\Delta\rho_{\text{min}} = -0.22 \text{ e \AA}^{-3}$ |
| 400 parameters | |
| H atoms treated by a mixture of independent and constrained refinement | |

Table 1

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

| $D-H\cdots A$ | $D-H$ | $H\cdots A$ | $D\cdots A$ | $D-H\cdots A$ |
|---------------------------------|----------|-------------|-------------|---------------|
| O5—H1 \cdots O1 ⁱ | 0.95 (2) | 1.64 (2) | 2.5832 (17) | 172 (2) |
| O7—H2 \cdots O4 ⁱⁱ | 0.90 (2) | 1.78 (3) | 2.682 (2) | 177 (2) |
| O3—H3 \cdots O2 | 1.16 (3) | 1.24 (3) | 2.3942 (18) | 174 (3) |

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

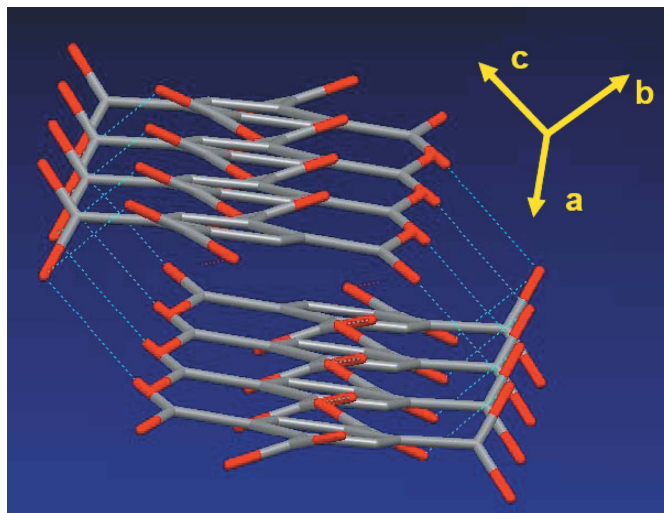


Figure 2

The hydrogen bonding (dashed lines) linking anions into a chain of centrosymmetric dimers. H atoms have been omitted.

C-bound H atoms were positioned geometrically and refined as riding, with C—H = 0.93 \AA and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$. Carboxyl H atoms were located in a difference map and refined freely.

Data collection: *CrystalClear* (Rigaku, 2000); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Sheldrick, 2000); software used to prepare material for publication: *SHELXTL*.

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