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## Key indicators

Single-crystal X-ray study $T=298 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.011 \AA$
$R$ factor $=0.056$
$w R$ factor $=0.138$
Data-to-parameter ratio $=16.6$
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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## [5-(Dibromomethyl)-3-methylbenzyl]triphenylphosphonium bromide

The asymmetric unit of the title structure, $\mathrm{C}_{27} \mathrm{H}_{24} \mathrm{Br}_{2} \mathrm{P}^{+} \cdot \mathrm{Br}^{-}$, comprises two $\left[\left(\mathrm{CH}_{3} \mathrm{C}_{6} \mathrm{H}_{3} \mathrm{CHBr}_{2}\right)\left(\mathrm{C}_{6} \mathrm{H}_{5}\right)_{3} \mathrm{P}\right]^{+}$cations and two bromide anions. The cations exhibit closely comparable conformations, with a slight distortion from tetrahedral geometry around $P$.

## Comment

The title compound includes two $\mathrm{C}_{27} \mathrm{H}_{24} \mathrm{Br}_{2} \mathrm{P}^{+}$cations and two bromide anions in the asymmetric unit (Fig. 1). The two independent cations have closely comparable conformations. The average $\mathrm{C}-\mathrm{P}$ bond length is 1.793 and $1.790 \AA$ in the two cations, comparable to the corresponding value of $1.792 \AA$ in the compound $\left[\mathrm{PPh}_{4}\right] \mathrm{Cl} \cdot \mathrm{H}_{2} \mathrm{O}$ (Blake et al., 2003), for example. The $\mathrm{C}-\mathrm{P}-\mathrm{C}$ angles span the range 107.5 (3) -111.5 (3) ${ }^{\circ}$, demonstrating a slight disortion from tetrahedral geometry around P .

(I)

## Experimental

A tetrahydrofuran solution ( 10 ml ) of 3-methyl-5-dibromomethylphenyl bromide $(0.17 \mathrm{~g}, 0.5 \mathrm{mmol})$ was added dropwise to a stirred tetrahydrofuran solution ( 10 ml ) of triphenylphosphine ( 0.12 g , 0.5 mmol ). Refluxing for 5 h gave a clear mixture from which yellow crystals of (I) were obtained in $65 \%$ yield after standing at room temperature for several days. Elemental analysis found: C $52.31, \mathrm{H}$ 3.77, Br 38.69 , P 4.86\%; calculated: C 52.37, H 3.91, Br 38.72, P 5.00\%.

## Crystal data

$\mathrm{C}_{27} \mathrm{H}_{24} \mathrm{Br}_{2} \mathrm{P}^{+} \cdot \mathrm{Br}^{-}$

$$
Z=8
$$

$M_{r}=619.16$
Monoclinic, $P 2_{b} / c$
$a=12.307$ (3) A
$b=21.789$ (4) $\AA$
$c=19.849$ (3) $\AA$
$\beta=91.387$ (2) ${ }^{\circ}$
$V=5321.2(17) \AA^{3}$

## Data collection

Bruker SMART CCD
diffractometer
$\varphi$ and $\omega$ scans
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
$T_{\text {min }}=0.182, T_{\text {max }}=0.262$

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## organic papers

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.056$
$w R\left(F^{2}\right)=0.138$
$S=1.01$
9287 reflections
559 parameters

H -atom parameters constrained
$w=1 /\left[\sigma^{2}\left(F_{\mathrm{o}}{ }^{2}\right)+(0.0549 P)^{2}\right]$
where $P=\left(F_{\mathrm{o}}{ }^{2}+2 F_{\mathrm{c}}{ }^{2}\right) / 3$
$(\Delta / \sigma)_{\max }<0.001$
$\Delta \rho_{\max }=1.12 \mathrm{e} \mathrm{A}^{-3}$
$\Delta \rho_{\min }=-0.84 \mathrm{e}^{-3}$

H atoms were placed in idealized positions and constrained to ride on their parent atoms, with $\mathrm{C}-\mathrm{H}=0.93 \AA$ for aromatic $\mathrm{H}, 0.98 \AA$ for $\mathrm{CHBr}_{2}, 0.97 \AA$ for $\mathrm{CH}_{2}$, and with $U_{\text {iso }}(\mathrm{H})=1.2 U_{\text {eq }}(\mathrm{C})$. For the methyl groups, $\mathrm{C}-\mathrm{H}=0.96 \AA$ and $U_{\text {iso }}(\mathrm{H})=1.5 U_{\text {eq }}(\mathrm{C})$. The highest peak in the residual electron density lies $1.66 \AA$ from the C36 methyl group.

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

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Figure 1
The asymmetric unit of (I), showing displacement ellipsoids at the $30 \%$ probability level. H atoms are omitted.

## References

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