

Tetramethylammonium dichlorodiphenylbismuthate(III)

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Key indicators

Single-crystal X-ray study
 $T = 173\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.007\text{ \AA}$
 R factor = 0.022
 wR factor = 0.052
Data-to-parameter ratio = 17.1

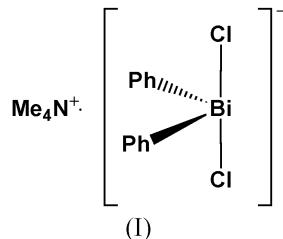
For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

The crystal structure of the title compound, $(\text{C}_4\text{H}_{12}\text{N})[\text{BiCl}_2(\text{C}_6\text{H}_5)_2]$, contains a $[\text{BiCl}_2\text{Ph}_2]^-$ anion with an equatorially vacant trigonal-bipyramidal geometry.

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Comment

A variety of anions of formula $[\text{ECl}_2\text{Ph}_2]^-$ are known for arsenic (Grewe *et al.*, 1998) and antimony (Sheldrick & Martin, 1992; Hall & Sowerby, 1988; Calderazzo *et al.*, 1991; Stark *et al.*, 1999). For bismuth, the $[\text{BiBr}_2\text{Ph}_2]^-$ (Stark *et al.*, 1999; Clegg *et al.*, 1992) and $[\text{BiI}_2\text{Ph}_2]^-$ (Clegg *et al.*, 1993) ions have been structurally characterized. This paper reports the structure of the $[\text{BiCl}_2\text{Ph}_2]^-$ anion as the tetramethylammonium salt, (I).



The structure of the anion in (I) is similar to that found for other diaryldihalobismuthate(III) anions. The geometry is based on a trigonal bipyramidal in which the formal lone pair and phenyl groups occupy equatorial positions and the halides reside in axial sites. For bismuth, the $\text{C}-\text{Bi}-\text{C}$ angle between the equatorial phenyl groups approaches 90° , which is typical of inter-bond angles in Bi^{III} structures (Clegg *et al.*, 1992, 1993).

Experimental

Crystals of the title compound were obtained from a reaction between BiCl_2Ph and $[\text{NMe}_4]\text{Cl}$ in tetrahydrofuran overlaid with hexane.

Crystal data

$(\text{C}_4\text{H}_{12}\text{N})[\text{BiCl}_2(\text{C}_6\text{H}_5)_2]$
 $M_r = 508.23$
Monoclinic, $P2_1/n$
 $a = 12.162 (3)\text{ \AA}$
 $b = 11.1693 (15)\text{ \AA}$
 $c = 14.241 (3)\text{ \AA}$
 $\beta = 110.642 (14)^\circ$
 $V = 1810.3 (7)\text{ \AA}^3$
 $Z = 4$

$D_x = 1.865\text{ Mg m}^{-3}$
Mo $K\alpha$ radiation
Cell parameters from 181 reflections
 $\theta = 2-20^\circ$
 $\mu = 10.03\text{ mm}^{-1}$
 $T = 173 (2)\text{ K}$
Block, colourless
 $0.30 \times 0.10 \times 0.05\text{ mm}$

Data collection

Bruker CCD area-detector diffractometer
 $0.3^\circ \omega$ scans
 Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)
 $T_{\min} = 0.305$, $T_{\max} = 0.605$
 9238 measured reflections

3185 independent reflections
 2787 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.026$
 $\theta_{\text{max}} = 25.0^\circ$
 $h = -11 \rightarrow 14$
 $k = -13 \rightarrow 13$
 $l = -16 \rightarrow 16$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.022$
 $wR(F^2) = 0.052$
 $S = 1.02$
 3185 reflections
 186 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0278P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.001$
 $\Delta\rho_{\text{max}} = 1.59 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -1.28 \text{ e } \text{\AA}^{-3}$
 Extinction correction: *SHELXL97*
 Extinction coefficient: 0.00272 (13)

Table 1

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

Bi1—C7	2.236 (5)	Bi1—Cl2	2.7310 (11)
Bi1—C1	2.252 (4)	Bi1—Cl1	2.7348 (11)
C7—Bi1—C1	94.74 (16)	C7—Bi1—Cl1	88.04 (10)
C7—Bi1—Cl2	87.37 (10)	C1—Bi1—Cl1	87.37 (10)
C1—Bi1—Cl2	89.82 (11)	Cl2—Bi1—Cl1	174.40 (4)

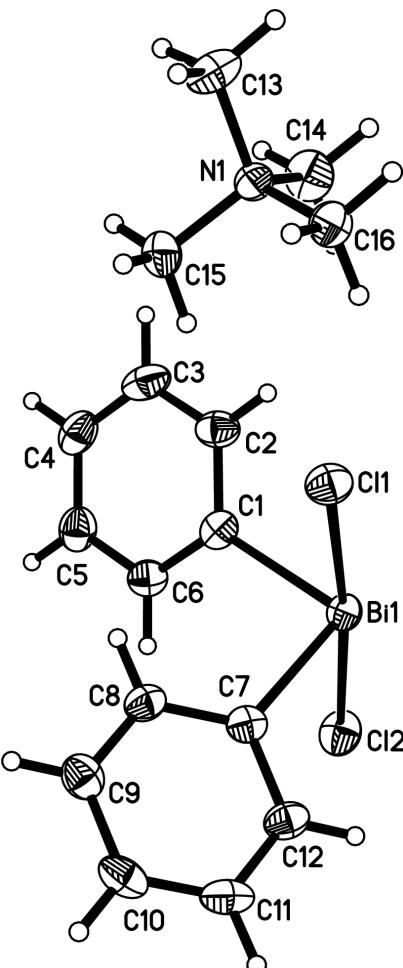
Methyl H atoms were positioned using a rotating-group refinement, with isotropic displacement parameters 1.5 times that of their adjacent C atom. The phenyl H atoms were constrained to ideal geometries and assigned isotropic displacement parameters 1.2 times that of their adjacent C atom. The two highest residual electron-density peaks (1.59 and 1.50 $\text{e } \text{\AA}^{-3}$) are found 0.97 and 0.98 \AA from the Bi atom. All other residual electron-density peaks have values less than 1 $\text{e } \text{\AA}^{-3}$.

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT* (Bruker, 1998); data reduction: *SHELXTL* (Bruker, 1998); program(s) used to solve structure: *SHELXTL*; program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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**Figure 1**

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