

## Sodium thiophenolate $N,N,N',N'$ -tetramethylethylenediamine

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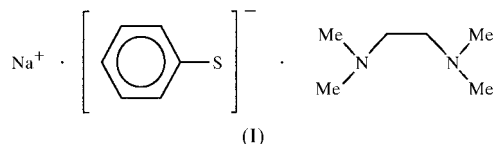
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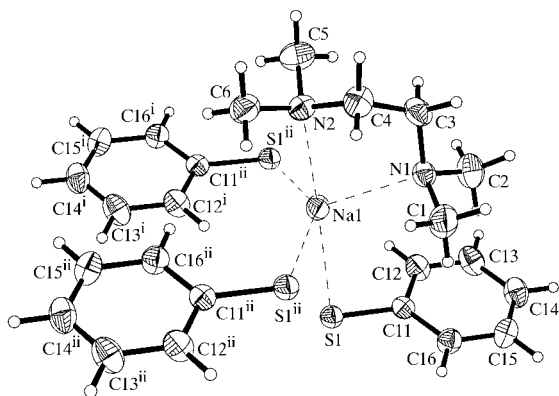
The title compound,  $\text{Na}^+ \cdot \text{C}_6\text{H}_5\text{S}^- \cdot \text{C}_6\text{H}_{16}\text{N}_2$ , forms lipophilically wrapped infinite sodium chains along the  $a$  axis. Each sodium cation is coordinated by three thiophenolate S atoms and two N centres of one tetramethylethylenediamine molecule.

### Comment

Strong phenyl–phenyl interactions stabilize the dimer of tris(phenylthio)methane, even in the gas phase (Bock *et al.*, 2001). An attempt to add a negative charge by reduction with sodium metal failed due to the decomposition of tris(phenylthio)methane to thiophenolate, which crystallizes in the



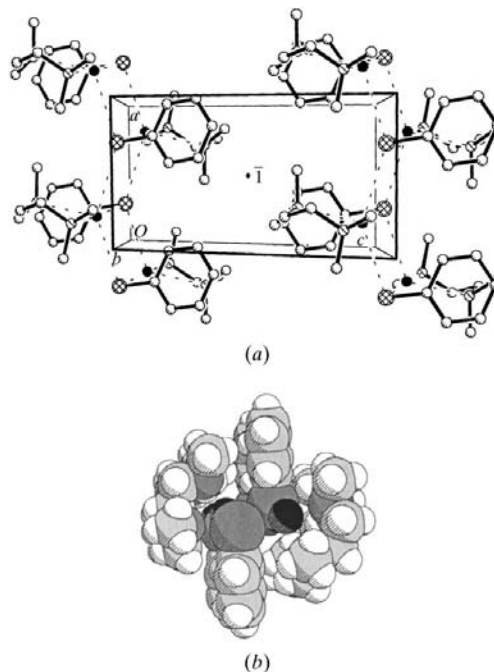
presence of a sodium metal mirror and tetramethylethylenediamine to give another lipophilically wrapped polyion aggregate, (I) (Bock, Hauck *et al.*, 1995; Bock, Schödel *et al.*, 1995; Bock *et al.*, 1998, 2000; Bock & Heigel, 2000*a,b*).



**Figure 1**

The coordination sphere of sodium in (I) with 50% probability displacement ellipsoids and the atom-numbering scheme. H atoms are drawn as spheres of arbitrary radii. The symmetry codes are as in Table 1.

The sodium cations form distorted trigonal bipyramids with sulfur centres from the thiophenolates and N centres from tetramethylethylenediamine. The axial positions are occupied



**Figure 2**

(a) The unit cell viewed along the  $b$  axis showing two infinite chains of (I) along the  $a$  axis and (b) a space-filled drawing of one chain (profile) illustrating the lipophilic wrapping of the polyionic chain inside (sodium: black; sulfur, nitrogen: dark grey; carbon: grey; hydrogen: white).

by S1 and N2, and the equatorial positions by S1A, S1B and N1 [Fig. 1; symmetry codes: (i)  $-x, 1 - y, -z$ ; (ii)  $1 - x, 1 - y, -z$ ]. These interatomic contacts agree with the sum of the van der Waals radii. Nevertheless, the interaction of the sodium cation with S1A and S1B decreases, but the geometry of a trigonal bipyramid is well defined (Fig. 1). The sodium cations are connected by  $\text{Na} \cdots \text{S} \cdots \text{Na}$  bridges to form infinite chains along the  $a$  axis with alternate  $\text{Na} \cdots \text{Na}$  distances of 4.132 (1) and 4.361 (1) Å (Fig. 2*a*). The C–S bond of the thiophenolate has a distance of 1.754 Å, which is in the mean band of 1.77 (3) calculated for the C–S distances in 472 crystal structures incorporating thiophenolate registered in the Cambridge Structural Database. The C–S bond of thiophenolates shows a broad variety in length obviously independent of the cation. Therefore, bis(tetra-*n*-butylammonium)tetrakis(benzene-thiolatoiron) containing only iron cations exhibits C–S distances with maximum differences of 0.04 Å in every published crystal structure (Guodong *et al.*, 1987; Gloux *et al.*, 1987; Excoffon *et al.*, 1991). The lipophilic wrapping by phenyl rings and methyl groups (Fig. 2*b*) is responsible for the kinetic stability of the polyion aggregate in analogy to the hexameric sodium tetraphenylimidodiphosphate (Bock, Schödel *et al.*, 1995).

## Experimental

From gentle heating of sodium (0.1 g, 4.3 mmol) at  $10^{-3}$  Pa, a mirror was generated on the wall of a dried Schlenk trap. Addition of tris(phenylthio)methane (100 mg, 0.2 mmol) in aprotic tetramethylethylenediamine (10 ml) produced a yellow solution, which after standing for 1 d at room temperature, was covered with a layer of aprotic hexane (10 ml). After one week, yellow crystals were collected from the trap wall near the phase border line.

### Crystal data

$\text{Na}^+ \cdot \text{C}_6\text{H}_5\text{S}^- \cdot \text{C}_6\text{H}_{16}\text{N}_2$	$Z = 2$
$M_r = 248.36$	$D_x = 1.187 \text{ Mg m}^{-3}$
Triclinic, $P\bar{1}$	Mo $K\alpha$ radiation
$a = 5.9440 (10) \text{ \AA}$	Cell parameters from 81 reflections
$b = 11.200 (2) \text{ \AA}$	$\theta = 8.5\text{--}18^\circ$
$c = 11.899 (2) \text{ \AA}$	$\mu = 0.241 \text{ mm}^{-1}$
$\alpha = 62.58 (1)^\circ$	$T = 180 (2) \text{ K}$
$\beta = 87.14 (2)^\circ$	Block, yellow
$\gamma = 81.32 (2)^\circ$	$0.60 \times 0.42 \times 0.25 \text{ mm}$
$V = 694.9 (2) \text{ \AA}^3$	

### Data collection

Siemens P4 four-circle diffractometer	$\theta_{\text{max}} = 28^\circ$
$\theta$ - $2\theta$ scans	$h = -1 \rightarrow 7$
3921 measured reflections	$k = -13 \rightarrow 13$
3066 independent reflections	$l = -15 \rightarrow 15$
2636 reflections with $I > 2\sigma(I)$	4 standard reflections
$R_{\text{int}} = 0.043$	every 100 reflections
	intensity decay: <5%

### Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F_o^2) + (0.0459P)^2 + 0.1435P]$
$R[F^2 > 2\sigma(F^2)] = 0.035$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.094$	$(\Delta/\sigma)_{\text{max}} < 0.001$
$S = 1.052$	$\Delta\rho_{\text{max}} = 0.34 \text{ e \AA}^{-3}$
3066 reflections	$\Delta\rho_{\text{min}} = -0.21 \text{ e \AA}^{-3}$
149 parameters	
H atoms constrained	

All H atoms were located from the difference map and placed in idealized positions using a riding model with distances  $\text{C}-\text{H}_{\text{phenyl}} = 0.95$ ,  $\text{C}-\text{H}_{\text{methylene}} = 0.99$  and  $\text{C}-\text{H}_{\text{methyl}} = 0.98 \text{ \AA}$ . The torsion of the methyl H atoms and the isotropic displacement parameters of all H atoms (groupwise for methyl and methylene H atoms) were refined.

Data collection: XSCANS (Siemens, 1994); cell refinement: XSCANS; data reduction: XSCANS; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: XP in

**Table 1**

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

S1—Na1	2.8219 (8)	Na1—S1 <sup>i</sup>	3.0310 (8)
Na1—N2	2.5375 (14)	Na1—S1 <sup>ii</sup>	3.1966 (8)
Na1—N1	2.5482 (13)	Na1—Na1 <sup>i</sup>	4.1323 (12)
N2—Na1—N1	73.60 (4)	S1—Na1—S1 <sup>i</sup>	90.249 (18)
N2—Na1—S1	174.95 (4)	N2—Na1—S1 <sup>ii</sup>	87.62 (3)
N1—Na1—S1	106.93 (3)	N1—Na1—S1 <sup>ii</sup>	93.31 (3)
N2—Na1—S1 <sup>i</sup>	93.77 (3)	S1—Na1—S1 <sup>ii</sup>	87.338 (19)
N1—Na1—S1 <sup>i</sup>	120.38 (3)	S1 <sup>i</sup> —Na1—S1 <sup>ii</sup>	145.28 (2)

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

SHELXTL/PC (Siemens, 1991); software used to prepare material for publication: SHELXL97 (Sheldrick, 1997).

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: AV1063). Services for accessing these data are described at the back of the journal.

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