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2,2′-(1,4-Phenylene)bis(propane-2,2-diyl) bis(benzodithioate)

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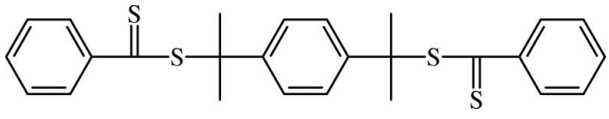
Key indicators: single-crystal X-ray study; T = 295 K; mean σ(C–C) = 0.003 Å; R factor = 0.032; wR factor = 0.091; data-to-parameter ratio = 15.9.

The title compound, $C_{26}H_{26}S_4$, shows a dihedral angle of 76.64 (15)° between the central and peripheral benzene rings. An inversion center is located at the centroid of the thiobenzoyl ring. In the crystal, weak C—H···S interactions form $C(5)$ chains along [001]. There are no classical hydrogen bonds.

Related literature

For control of the behavior of polymerization reactions, see: Patton *et al.* (2005); You *et al.* (2007); Pafiti *et al.* (2010). For radical polymerization with RAFT reactions, see: Le *et al.* (1998). For telechelic polymers, see: Tasdelen *et al.* (2011); Goethals (1989). For hydrogen bonding, see: Nardelli (1995). For graph-set motifs, see: Etter (1990). For standard bond lengths, see: Allen *et al.* (1987).



Experimental

Crystal data

$C_{26}H_{26}S_4$
M_r = 466.75
Monoclinic, $P2_1/c$
a = 8.6981 (6) Å
b = 11.7074 (7) Å
c = 12.5612 (6) Å
β = 107.626 (4)°
V = 1219.08 (13) Å^3
Z = 2
Mo $K\alpha$ radiation
μ = 0.40 mm^{-1}
T = 295 K
0.41 × 0.29 × 0.16 mm

Data collection

Bruker–Nonius KappaCCD diffractometer
Absorption correction: multi-scan (*SADABS*; Sheldrick, 2002)
T_{min} = 0.874, T_{max} = 0.939
4051 measured reflections
2160 independent reflections
1775 reflections with $I > 2\sigma(I)$
R_{int} = 0.021

Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.032
$wR(F^2)$ = 0.091
S = 1.03
2160 reflections
136 parameters
H-atom parameters constrained
$\Delta\rho_{max}$ = 0.19 e Å^{-3}
$\Delta\rho_{min}$ = −0.22 e Å^{-3}

Table 1
Hydrogen-bond geometry (Å, °).

D—H···A	D—H	H···A	D···A	D—H···A
C5—H5···S2^{i}	0.93	2.94	3.489 (2)	119

Symmetry code: (i) $x, -y+\frac{1}{2}, z+\frac{1}{2}$.

Data collection: *COLLECT* (Hooft, 2004); cell refinement: *SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *DENZO* (Otwinowski & Minor, 1997) and *SCALEPACK*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 2012) and *Mercury* (Macrae *et al.*, 2006); software used to prepare material for publication: *WinGX* (Farrugia, 2012).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: KJ2235).

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supporting information

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2,2′-(1,4-Phenylene)bis(propane-2,2-diyl) bis(benzodithioate)

Rodolfo Moreno-Fuquen, Carlos Grande, Rigoberto C. Advincula, Juan C. Tenorio and Javier Ellena

S1. Comment

The title compound belongs to a series of difunctional compounds that can be used to control the behavior of polymerization reactions to produce straight forward functional telechelic polymers in one pot (Patton *et al.*, 2005; You *et al.*, 2007; Pafiti *et al.*, 2010). They are also used in radical polymerization with RAFT (reversible addition fragmentation chain transfer) reactions (Le *et al.*, 1998). Telechelic polymers, defined as macromolecules with two reactive end groups, have been used for multiple purposes (Tasdelen *et al.*, 2011) including block copolymer synthesis (Goethals, 1989). A perspective view of the molecule of the title compund, showing the atomic numbering scheme, is given in Fig. 1. Bond lengths and angles in the title compound have normal values (Allen *et al.*, 1987). The molecular system has an inversion center and it is located at the center of the thiobenzoyl ring. The benzene rings bridged by the thio (C6—C7—S1—C8—C11) moiety are tilted to each other by a dihedral angle of 76.64 (15)°. The crystal packing shows no classical hydrogen bonds and it is stabilized by weak C—H···S intermolecular interactions, forming C(5) chains (Etter, 1990) along [001] (see Fig. 2; Etter, 1990). The C5 atom of the benzene ring at (*x*,*y*,*z*) acts as hydrogen-bond donors to S2 atom at (*x*, -*y* + 1/2, *z* - 1/2) (see Table 1; Nardelli, 1995).

S2. Experimental

The synthesis of the mentioned compound was accomplished following a procedure already reported (Le *et al.*, 1998; Patton *et al.*, 2005). A mixture of dithiobenzoic acid (5.00 g, 32.4 mmol) and 1,4-diisopropenylbenzene (2.44 g, 15.4 mmol) in carbon tetrachloride (40 ml) was heated at 348 K for 20 h. The volatiles were removed under reduced pressure and the oily product was mixed with 1:2 diethyl ether/hexane to isolate the product as a pink solid (40%).

S3. Refinement

All H-atoms were placed in calculated positions [C—H= 0.95 Å for aromatic and C—H= 0.96 Å for methyl group] and refined with U_{iso}(H) 1.2 and 1.5 times U_{eq} of the parent atom, respectively.

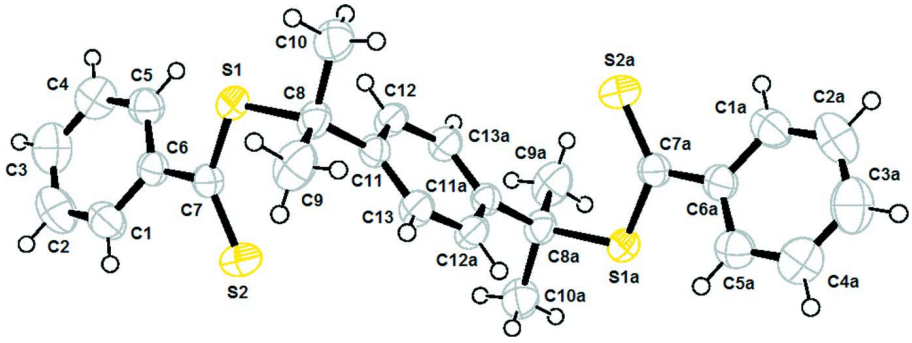

Figure 1

An *ORTEP-3* (Farrugia, 2012) plot of the title compound with displacement ellipsoids drawn at the 50% probability level. H atoms are shown as spheres of arbitrary radius. Atoms labelled with suffix "a" are generated by an inversion center via operation (1-x, 1-y, -z).

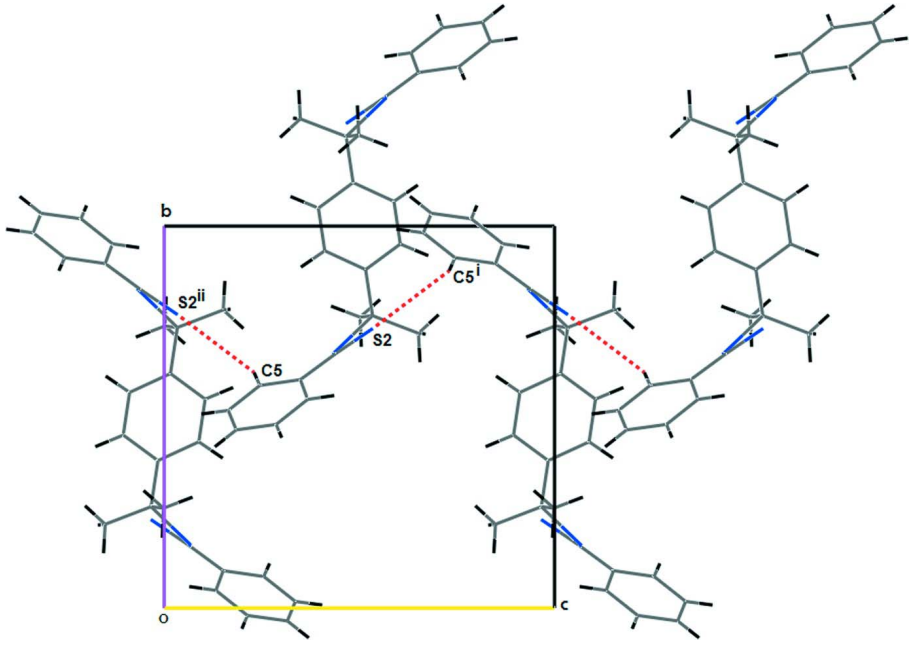

Figure 2

Part of the crystal structure of the title compound, showing the formation of chains of molecules running along [001]. Symmetry code: (i) *x*,-*y* + 1/2,*z* + 1/2. (ii) *x*,-*y* + 1/2,*z* - 1/2.

2,2′-(1,4-Phenylene)bis(propane-2,2-diyl) bis(benzodithioate)

Crystal data

$C_{26}H_{26}S_4$
M_r = 466.75
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b = 11.7074 (7) Å
c = 12.5612 (6) Å
β = 107.626 (4)°
V = 1219.08 (13) Å^3
Z = 2
$F(000)$ = 492
D_x = 1.272 Mg m^{-3}
Mo $K\alpha$ radiation, λ = 0.71073 Å
Cell parameters from 4051 reflections
μ = 0.40 mm^{-1}
T = 295 K
Block, pink
0.41 × 0.29 × 0.16 mm

Data collection

Bruker–Nonius KappaCCD diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
CCD rotation images, thick slices scans
Absorption correction: multi-scan (*SADABS*; Sheldrick, 2002)
T_{min} = 0.874, T_{max} = 0.939
4051 measured reflections
2160 independent reflections
1775 reflections with $I > 2\sigma(I)$
R_{int} = 0.021
θ_{max} = 25.1°, θ_{min} = 3.0°
h = −10→10
k = −12→13
l = −14→14

Refinement

Refinement on F^2
Least-squares matrix: full
$R[F^2 > 2\sigma(F^2)]$ = 0.032
$wR(F^2)$ = 0.091
S = 1.03
2160 reflections
136 parameters
0 restraints
Primary atom site location: structure-invariant direct methods
Secondary atom site location: difference Fourier map
Hydrogen site location: inferred from neighbouring sites
H-atom parameters constrained
$w = 1/[\sigma^2(F_o^2) + (0.0431P)^2 + 0.3633P]$ where $P = (F_o^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{max} < 0.001$
$\Delta\rho_{max}$ = 0.19 e Å^{-3}
$\Delta\rho_{min}$ = −0.22 e Å^{-3}

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R-factor wR and goodness of fit S are based on F^2, conventional R-factors R are based on F, with F set to zero for negative F^2. The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R-factors(gt) *etc*. and is not relevant to the choice of reflections for refinement. R-factors based on F^2 are statistically about twice as large as those based on F, and R- factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (Å²)

	x	y	z	U_{iso}*/U_{eq}
S1	0.50091 (6)	0.16376 (4)	0.06933 (4)	0.04958 (17)
S2	0.75578 (8)	0.23202 (6)	−0.03441 (5)	0.0685 (2)
C1	0.9423 (3)	0.0464 (2)	0.13247 (19)	0.0610 (6)
H1	0.9617	0.0565	0.0642	0.073*
C2	1.0454 (3)	−0.0191 (2)	0.2131 (2)	0.0754 (7)
H2	1.1332	−0.0541	0.1990	0.090*
C3	1.0198 (3)	−0.0335 (2)	0.3155 (2)	0.0754 (7)
H3	1.0904	−0.0778	0.3704	0.091*
C4	0.8906 (3)	0.0175 (2)	0.33591 (18)	0.0644 (6)
H4	0.8736	0.0081	0.4050	0.077*
C5	0.7852 (3)	0.08278 (17)	0.25504 (16)	0.0526 (5)
H5	0.6975	0.1171	0.2699	0.063*
C6	0.8086 (2)	0.09805 (15)	0.15112 (16)	0.0456 (4)
C7	0.6960 (2)	0.16592 (15)	0.06088 (16)	0.0462 (5)
C8	0.3740 (2)	0.26543 (15)	−0.03377 (16)	0.0462 (4)
C9	0.3421 (3)	0.21941 (19)	−0.15129 (18)	0.0668 (6)

H9A	0.2986	0.1436	−0.1554	0.100*
H9B	0.4413	0.2174	−0.1699	0.100*
H9C	0.2663	0.2681	−0.2030	0.100*
C10	0.2164 (3)	0.26165 (19)	−0.0017 (2)	0.0628 (6)
H10A	0.2361	0.2906	0.0727	0.094*
H10B	0.1789	0.1842	−0.0050	0.094*
H10C	0.1360	0.3078	−0.0529	0.094*
C11	0.4441 (2)	0.38607 (15)	−0.01603 (14)	0.0388 (4)
C12	0.4914 (2)	0.43533 (16)	0.08868 (15)	0.0446 (4)
H12	0.4861	0.3926	0.1499	0.054*
C13	0.4535 (2)	0.45299 (16)	−0.10490 (15)	0.0439 (4)
H13	0.4222	0.4225	−0.1766	0.053*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0527 (3)	0.0368 (3)	0.0662 (3)	0.0003 (2)	0.0284 (2)	0.0081 (2)
S2	0.0734 (4)	0.0679 (4)	0.0797 (4)	0.0051 (3)	0.0461 (3)	0.0187 (3)
C1	0.0580 (13)	0.0617 (13)	0.0683 (13)	0.0053 (11)	0.0264 (11)	−0.0067 (11)
C2	0.0560 (14)	0.0766 (17)	0.0882 (18)	0.0152 (12)	0.0138 (13)	−0.0131 (14)
C3	0.0697 (16)	0.0678 (16)	0.0714 (16)	0.0058 (13)	−0.0046 (13)	−0.0023 (13)
C4	0.0687 (15)	0.0667 (14)	0.0507 (12)	−0.0063 (12)	0.0077 (11)	−0.0045 (11)
C5	0.0566 (12)	0.0493 (11)	0.0532 (11)	−0.0037 (9)	0.0186 (10)	−0.0082 (9)
C6	0.0502 (10)	0.0342 (9)	0.0551 (11)	−0.0045 (8)	0.0200 (9)	−0.0074 (8)
C7	0.0551 (11)	0.0340 (9)	0.0564 (11)	−0.0011 (8)	0.0270 (9)	−0.0042 (8)
C8	0.0493 (11)	0.0349 (9)	0.0540 (11)	−0.0059 (8)	0.0150 (9)	−0.0007 (8)
C9	0.0871 (17)	0.0466 (12)	0.0601 (13)	−0.0201 (12)	0.0124 (12)	−0.0129 (10)
C10	0.0485 (12)	0.0508 (12)	0.0889 (16)	−0.0078 (9)	0.0207 (12)	0.0033 (11)
C11	0.0379 (9)	0.0333 (9)	0.0461 (10)	−0.0006 (7)	0.0140 (8)	−0.0014 (8)
C12	0.0551 (11)	0.0391 (10)	0.0414 (10)	−0.0035 (9)	0.0171 (8)	0.0040 (8)
C13	0.0524 (11)	0.0396 (10)	0.0391 (9)	−0.0037 (8)	0.0129 (8)	−0.0040 (8)

Geometric parameters (Å, º)

S1—C7	1.7325 (19)	C8—C9	1.516 (3)
S1—C8	1.857 (2)	C8—C11	1.528 (2)
S2—C7	1.6366 (19)	C8—C10	1.541 (3)
C1—C2	1.367 (3)	C9—H9A	0.9600
C1—C6	1.393 (3)	C9—H9B	0.9600
C1—H1	0.9300	C9—H9C	0.9600
C2—C3	1.380 (4)	C10—H10A	0.9600
C2—H2	0.9300	C10—H10B	0.9600
C3—C4	1.364 (3)	C10—H10C	0.9600
C3—H3	0.9300	C11—C12	1.380 (2)
C4—C5	1.376 (3)	C11—C13	1.386 (2)
C4—H4	0.9300	C12—C13^{i}	1.386 (3)
C5—C6	1.392 (3)	C12—H12	0.9300
C5—H5	0.9300	C13—C12^{i}	1.386 (3)

C6—C7	1.484 (3)	C13—H13	0.9300
C7—S1—C8	109.55 (9)	C9—C8—S1	110.25 (14)
C2—C1—C6	120.9 (2)	C11—C8—S1	111.34 (13)
C2—C1—H1	119.6	C10—C8—S1	100.81 (13)
C6—C1—H1	119.6	C8—C9—H9A	109.5
C1—C2—C3	120.3 (2)	C8—C9—H9B	109.5
C1—C2—H2	119.9	H9A—C9—H9B	109.5
C3—C2—H2	119.9	C8—C9—H9C	109.5
C4—C3—C2	119.8 (2)	H9A—C9—H9C	109.5
C4—C3—H3	120.1	H9B—C9—H9C	109.5
C2—C3—H3	120.1	C8—C10—H10A	109.5
C3—C4—C5	120.5 (2)	C8—C10—H10B	109.5
C3—C4—H4	119.8	H10A—C10—H10B	109.5
C5—C4—H4	119.8	C8—C10—H10C	109.5
C4—C5—C6	120.7 (2)	H10A—C10—H10C	109.5
C4—C5—H5	119.7	H10B—C10—H10C	109.5
C6—C5—H5	119.7	C12—C11—C13	117.27 (16)
C5—C6—C1	117.94 (19)	C12—C11—C8	121.06 (16)
C5—C6—C7	122.36 (17)	C13—C11—C8	121.55 (16)
C1—C6—C7	119.70 (18)	C11—C12—C13^{i}	121.62 (16)
C6—C7—S2	121.99 (14)	C11—C12—H12	119.2
C6—C7—S1	112.15 (13)	C13^{i}—C12—H12	119.2
S2—C7—S1	125.85 (13)	C11—C13—C12^{i}	121.12 (17)
C9—C8—C11	114.68 (16)	C11—C13—H13	119.4
C9—C8—C10	109.22 (18)	C12^{i}—C13—H13	119.4
C11—C8—C10	109.63 (16)		
C6—C1—C2—C3	−1.1 (4)	C7—S1—C8—C9	71.76 (16)
C1—C2—C3—C4	0.3 (4)	C7—S1—C8—C11	−56.71 (15)
C2—C3—C4—C5	0.3 (4)	C7—S1—C8—C10	−172.92 (13)
C3—C4—C5—C6	−0.1 (3)	C9—C8—C11—C12	−174.69 (18)
C4—C5—C6—C1	−0.7 (3)	C10—C8—C11—C12	62.0 (2)
C4—C5—C6—C7	178.79 (18)	S1—C8—C11—C12	−48.6 (2)
C2—C1—C6—C5	1.2 (3)	C9—C8—C11—C13	9.4 (3)
C2—C1—C6—C7	−178.2 (2)	C10—C8—C11—C13	−113.8 (2)
C5—C6—C7—S2	151.88 (16)	S1—C8—C11—C13	135.50 (16)
C1—C6—C7—S2	−28.7 (3)	C13—C11—C12—C13^{i}	−0.2 (3)
C5—C6—C7—S1	−28.9 (2)	C8—C11—C12—C13^{i}	−176.23 (17)
C1—C6—C7—S1	150.59 (16)	C12—C11—C13—C12^{i}	0.2 (3)
C8—S1—C7—C6	172.46 (12)	C8—C11—C13—C12^{i}	176.21 (17)
C8—S1—C7—S2	−8.33 (16)		

Symmetry code: (i) −*x*+1, −*y*+1, −*z*.

Hydrogen-bond geometry (Å, º)

D—H···*A*	*D*—H	H···*A*	*D*···*A*	*D*—H···*A*

C5—H5···S2^{ii}	0.93	2.94	3.489 (2)	119

Symmetry code: (ii) x, $-y+1/2$, $z+1/2$.