# organic papers

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# Hong Dae Choi,<sup>a</sup> Pil Ja Seo,<sup>a</sup> Byeng Wha Son<sup>b</sup> and Uk Lee<sup>b</sup>\*

<sup>a</sup>Department of Chemistry, Dongeui University, San 24 Kaya-dong Busanjin-ku, Busan 614-714, Republic of Korea, and <sup>b</sup>Department of Chemistry, Pukyong National University, 599-1 Daeyeon 3-dong Nam-ku, Busan 608-737, Republic of Korea

Correspondence e-mail: uklee@pknu.ac.kr

#### **Key indicators**

Single-crystal X-ray study T = 173 K Mean  $\sigma$ (C–C) = 0.003 Å R factor = 0.046 wR factor = 0.135 Data-to-parameter ratio = 17.9

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

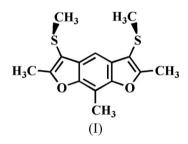
# 2,6,8-Trimethyl-3,5-bis(methylsulfanyl)benzo-[1,2-b:5,4-b']difuran

The title compound,  $C_{15}H_{16}O_2S_2$ , was prepared by the Lewis acid-catalysed reaction of 2-methylresorcinol with  $\alpha$ -chloro- $\alpha$ -(methylsulfanyl)acetone. The benzo[1,2-*b*:5,4-*b'*]difuran group is almost planar. The crystal structure is stabilized by aromatic  $\pi$ - $\pi$  stacking and  $-CH_2-H\cdots\pi$  interactions.

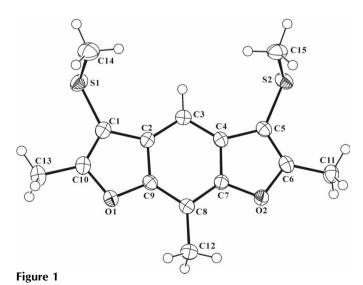
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#### Comment

This work follows on from our previous reports on the benzodifuran analogues, 2,5-dimethyl-3,4-bis(methylsulfanyl)benzo[1,2-b:4,3-b']difuran (Choi, Jang *et al.*, 2006) and 2,4,7trimethyl-3,8-bis(methylsulfanyl)benzo[1,2-b:5,6-b']difuran (Choi, Seo *et al.*, 2006). The isomeric title compound, (I) (Fig. 1), was obtained by the one-pot reaction of 2-methylresorcinol with an excess of  $\alpha$ -chloro- $\alpha$ -(methylsulfanyl)acetone in the presence of zinc chloride.



The bond lengths and angles in (I) are within normal ranges (Allen *et al.*, 1987; Choi, Jang *et al.*, 2006; Choi, Seo *et al.*, 2006). The benzo[1,2-*b*:5,4-*b*']difuran unit is nearly planar,



© 2007 International Union of Crystallography All rights reserved The molecular structure of (I), with 50% probability displacement ellipsoids for non-H atoms.

with a mean deviation of 0.014 Å from the least-squares plane defined by the 12 constituent C and O atoms. The methyl groups of the methylsulfanyl substituents point in the same direction with respect to the plane of the ring system. Atoms S1 and S2 lie 0.022 (2) and 0.088 (3) Å, respectively, out of the C1/C2/C9/O1/C10 and C4–C6/O2/C7 planes.

The crystal structure is stabilized by aromatic  $\pi$ - $\pi$  stacking and  $-CH_2-H\cdots\pi$  interactions (Fig. 2). The closest approach between the benzodifuran planes (BF) is BF $\cdots$ BF<sup>iii</sup> = 3.556 (5) Å, while the nearest ring-centroid separation,  $Cg2\cdots Cg3^{iii}$ , is 3.894 (4) Å (Cg2 and Cg3 are the centroids of the C2-C4/C7-C9 and C4-C6/O2/C7 rings, respectively; symmetry code as in Fig. 2). These data indicate the existence of a weak  $\pi$ - $\pi$  stacking interaction. The packing of the molecules is additionally stabilized by the  $-CH_2-H\cdots\pi$  interactions between adjacent furan rings (Table 1).

## **Experimental**

The title compound was prepared according to the method described for the previously reported isomer (Choi, Seo *et al.*, 2006), using 2methylresorcinol instead of 5-methylresorcinol. Single crystals suitable for X-ray analysis were obtained by slow evaporation of a tetrahydrofuran solution [yield 53%; m.p. 403–404 K;  $R_{\rm F} = 0.67$ (CCl<sub>4</sub>)].

## Crystal data

 $C_{15}H_{16}O_2S_2$   $M_r = 292.40$ Monoclinic,  $P2_1/c$  a = 7.895 (3) Å b = 13.008 (5) Å c = 14.498 (6) Å  $\beta = 104.479$  (7)° V = 1441.6 (10) Å<sup>3</sup>

#### Data collection

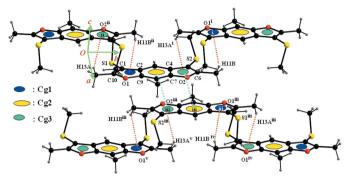
Bruker SMART CCD diffractometer  $\varphi$  and  $\omega$  scans Absorption correction: none 8031 measured reflections

#### Refinement

Refinement on  $F^2$   $R[F^2 > 2\sigma(F^2)] = 0.046$   $wR(F^2) = 0.135$  S = 1.053097 reflections 173 parameters H-atom parameters constrained Z = 4  $D_x = 1.347 \text{ Mg m}^{-3}$ Mo K $\alpha$  radiation  $\mu = 0.36 \text{ mm}^{-1}$ T = 173 (2) K Block, colorless 0.65 × 0.58 × 0.54 mm

3097 independent reflections 2702 reflections with  $I > 2\sigma(I)$  $R_{int} = 0.070$  $\theta_{max} = 27.0^{\circ}$ 

$$\begin{split} &w = 1/[\sigma^2(F_{\rm o}^2) + (0.0664P)^2 \\ &+ 1.1009P] \\ &where \ P = (F_{\rm o}^2 + 2F_{\rm c}^2)/3 \\ (\Delta/\sigma)_{\rm max} < 0.001 \\ \Delta\rho_{\rm max} = 0.29 \ {\rm e} \ {\rm \AA}^{-3} \\ \Delta\rho_{\rm min} = -0.37 \ {\rm e} \ {\rm \AA}^{-3} \end{split}$$



#### Figure 2

The C-H··· $\pi$  and  $\pi$ - $\pi$  interactions (dotted lines) in (I). *Cgn* denotes the centroid of each ring. [Symmetry codes: (i)  $1 - x, \frac{1}{2} + y, \frac{1}{2} - z$ ; (ii)  $1 - x, y - \frac{1}{2}, \frac{1}{2} - z$ ; (iii) 1 - x, 1 - y, -z; (iv)  $x, \frac{3}{2} - y, z - \frac{1}{2}$ ; (v)  $x, \frac{1}{2} - y, z - \frac{1}{2}$ .]

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

$\overline{D-\mathrm{H}\cdots A}$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$\begin{array}{c} \hline C11-H11B\cdots Cg1^{i}\\ C13-H13A\cdots Cg3^{ii} \end{array}$	0.96	2.75	3.480 (3)	133
	0.96	2.82	3.586 (3)	138

Symmetry codes: (i) -x + 1,  $y + \frac{1}{2}$ ,  $-z + \frac{1}{2}$ ; (ii) -x + 1,  $y - \frac{1}{2}$ ,  $-z + \frac{1}{2}$ . *Cg*1 is the centroid of the C1/C2/C9/O1/C10 ring and *Cg*3 is the centroid of the C4–C6/O2/C7 ring.

All H atoms were positioned geometrically (C–H = 0.93–0.96 Å) and refined as riding, with  $U_{iso}(H) = 1.2U_{eq}(C)$  or  $1.5U_{eq}(methyl C)$ .

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: *ORTEP-3* (Farrugia, 1997) and *DIAMOND* (Brandenburg, 1998); software used to prepare material for publication: *SHELXL97*.

## References

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