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Key indicators

Single-crystal X-ray study T = 297 KMean $\sigma(\text{C}-\text{C}) = 0.005 \text{ Å}$ R factor = 0.068 wR factor = 0.152 Data-to-parameter ratio = 15.8

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

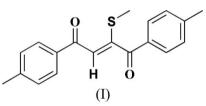
(*Z*)-2-Methylsulfanyl-1,4-di-*p*-tolylbut-2-ene-1,4-dione

The title compound, $C_{19}H_{18}O_2S$, displays a *trans* configuration with respect to the central C=C double bond. The dihedral angle between the two aromatic rings is 67.91 (5)°.

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Comment

1,4-Dicarbonyl compounds are widely used as synthetic building blocks for further elaboration into substituted cyclopentenones, such as jasmones, rethorolones, cuparenones and prostaglandins, and five-membered heterocyclic compounds, such as furans, pyrroles, thiophenes and pyridazines (Ellison, 1973; Yuguchi *et al.*, 2004). As a part of our ongoing investigation into 1,4-dicarbonyl compounds (Chen *et al.*, 2007), we present here the structure of the title compound, (I) (Fig. 1).



The dihedral angle between the two aromatic rings is $67.91 (5)^{\circ}$. The structure is stabilized by intermolecular C-H···O hydrogen bonds (Table 1).

Experimental

Compound (I) was prepared according to the method described by Yin *et al.* (2006). Single crystals suitable for X-ray analysis were obtained by slow evaporation of a dichloromethane–methanol (1:1, v/v) solution at 283 K.

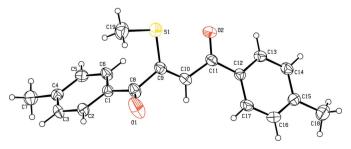


Figure 1

The molecular structure of (I) showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level and H atoms are represented by circles of arbitrary size.

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Crystal data

 $C_{19}H_{18}O_2S$ $M_r = 310.39$ Orthorhombic, $P2_12_12_1$ a = 5.7812 (6) Å b = 7.5573 (8) Å c = 37.412 (4) Å V = 1634.5 (3) Å³

Data collection

Bruker SMART 4K CCD areadetector diffractometer φ and ω scans Absorption correction: none 12645 measured reflections

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0684P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.068$	+ 0.2905P]
$wR(F^2) = 0.152$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.09	$(\Delta/\sigma)_{\rm max} = 0.017$
3199 reflections	$\Delta \rho_{\rm max} = 0.28 \text{ e} \text{ Å}^{-3}$
202 parameters	$\Delta \rho_{\rm min} = -0.16 \ {\rm e} \ {\rm \AA}^{-3}$
H-atom parameters constrained	Absolute structure: Flack (1983),
	1287 Friedel pairs

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$C16-H16\cdots O2^i$	0.93	2.43	3.318 (5)	159

Symmetry code: (i) x, y - 1, z.

Z = 4 $D_x = 1.261 \text{ Mg m}^{-3}$ Mo $K\alpha$ radiation $\mu = 0.20 \text{ mm}^{-1}$ T = 297 (2) K Block, yellow $0.20 \times 0.10 \times 0.10$ mm

3199 independent reflections 2450 reflections with $I > 2\sigma(I)$ $R_{\rm int} = 0.052$ $\theta_{\rm max} = 26.0^{\circ}$

Flack parameter: 0.07 (16)

All H atoms were refined using a riding model, with C-H = 0.93 Å and $U_{iso}(H) = 1.2U_{eq}(C)$ for aromatic H, and C-H = 0.96 Å and $U_{\rm iso}({\rm H}) = 1.5 U_{\rm eq}({\rm C})$ for methyl H atoms.

Data collection: SMART (Bruker, 1997); cell refinement: SAINT (Bruker, 1999); 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, 2001); software used to prepare material for publication: SHELXTL.

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