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Key indicators

Single-crystal X-ray study T = 294 K Mean σ (C–C) = 0.004 Å R factor = 0.044 wR factor = 0.115 Data-to-parameter ratio = 14.1

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3-(4-*tert*-Butyl-2,6-dimethyl-3,5-dinitrobenzoyl)-1,1-bis(methylsulfanyl)prop-2-en-1-one

The bond lengths and molecular conformation of the title compound, $C_{17}H_{22}N_2O_5S_2$, can be correlated with electronic conjugation and steric effects.

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Comment

As precursors for the synthesis of heterocyclic ketene aminals (HKAs) (Huang & Wang, 1994, 2002), aroyl-substituted ketene dimethylmercaptals have been widely used to synthesize a variety of heterocycles. The title compound, (I) (Fig. 1), was obtained from the one-pot reaction of 1-(4-*tert*-butyl-2,6-dimethyl-3,5-dinitrophenyl)ethanone, carbon disulfide and methyl iodide.



The geometrical parameters for (I) are normal. The length of the C13–C14 bond is shorter than that of a normal C–C single bond, owing to the conjugation involving the adjacent C13=O5 and C14=C15 bonds (Table 1). The lengths of the C15–S1 and C15–S2 bonds are shorter than that of a normal C–S single bond for the same reason. The C13/C14/C15/O5/ S1/S2 assembly is almost planar [r.m.s. deviation from the mean plane of 0.0071 (16) Å] and close to perpendicular to the benzene ring mean plane [dihedral angle = 89.37 (6)°]. Both nitro groups are almost perpendicular to the benzene ring plane, probably for steric reasons to avoid unfavourable interactions with the C3 and C12 methyl groups and the C6 *tert*-butyl group.

Experimental

1-(4-*tert*-Butyl-2,6-dimethyl-3,5-dinitrophenyl)ethanone (58.86 g, 0.2 mol) was added dropwise to a suspension of sodium hydride (12 g, 80%, 0.4 mol) in toluene (200 ml) and dimethylformamide (50 ml). Carbon disulfide (15.23 g, 0.2 mol) was then added dropwise to the reaction mixture slowly. After stirring for 2 h, methyl iodide (56.78 g, 0.4 mol) was added dropwise while the reaction flask was cooled in an ice bath. When thin-layer chromatography indicated the reaction was complete, the mixture was poured into ice–water and the organic phase was separated from the aqueous phase. The aqueous phase was extracted with ethyl acetate (3 × 50 ml). The combined organic phases were dried over anhydrous sodium sulfate. After partial removal of the solvents, the title compound was crystallized from the solution: yellow needles of (I) were collected by filtration and washed with petroleum ether (Huang & Liu, 1989). (yield 51.81 g, 65%; m.p.



Figure 1

The molecular structure of (I), showing 50% displacement ellipoids (arbitrary spheres for the H atoms).

495–496 K). ¹H NMR (CDCl₃): δ 6.04 (*s*, 1H, ==CH), 2.52 (*s*, 3H, SCH₃), 2.47 (*s*, 3H, SCH₃), 2.12 (*s*, 6H, 2CH₃), 1.44 (*s*, 9H, C(CH₃)₃); ESI-MS: *m*/*z* 39 (*M*⁺ + 1); Analysis calculated for C₁₇H₂₂N₂O₅S₂: C 51.24, H 5.56, N 7.03%; found: C 50.87, H 5.59, N 7.03%.

Crystal data

$C_{17}H_{22}N_2O_5S_2$	V = 971.5 (4) Å ³	
$M_r = 398.49$	Z = 2	
Triclinic, P1	$D_x = 1.362 \text{ Mg m}^{-3}$	
a = 8.657 (2) Å	Mo $K\alpha$ radiation	
b = 10.193 (3) Å	$\mu = 0.30 \text{ mm}^{-1}$	
c = 12.561 (3) Å	T = 294 (2) K	
$\alpha = 67.272 \ (4)^{\circ}$	Prism, yellow	
$\beta = 71.840 \ (4)^{\circ}$	$0.24 \times 0.20 \times 0.16 \text{ mm}$	
$\gamma = 82.758 \ (4)^{\circ}$		
Data collection		

Data collection

Bruker SMART CCD diffractometer ω scans

Absorption correction: multi-scan (SADABS; Bruker, 2002) $T_{\min} = 0.767, T_{\max} = 1.000$ 4980 measured reflections 3403 independent reflections 2092 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.025$ $\theta_{\text{max}} = 25.0^{\circ}$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.044$	H-atom parameters constrained $w = 1/[\sigma^2(F^2) + (0.05P)^2]$
$vR(F^2) = 0.115$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.07	$(\Delta/\sigma)_{\rm max} = 0.003$
403 reflections	$\Delta \rho_{\rm max} = 0.25 \ {\rm e} \ {\rm \AA}^{-3}$
42 parameters	$\Delta \rho_{\rm min} = -0.26 \text{ e } \text{\AA}^{-3}$

Table 1 Selected bond lengths (Å).

C15-S1	1.740 (3)	C13-C14	1.436 (3)
C15-S2	1.753 (3)	C14-C15	1.348 (4)

The H atoms were located in difference maps, repositioned in idealized positions (C–H = 0.93–0.96 Å) and refined as riding, with $U_{\rm iso}({\rm H}) = 1.2 U_{\rm eq}({\rm C})$ or $1.5 U_{\rm eq}({\rm methyl~C})$.

Data collection: *SMART* (Bruker, 2002); cell refinement: *SAINT* (Bruker, 2002); 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, 2002); software used to prepare material for publication: *SHELXTL*.

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