

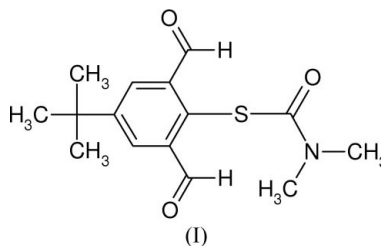
Christine McKenzie and  
Andrew D. Bond\*University of Southern Denmark, Department of  
Chemistry, Campusvej 55, 5230 Odense,  
Denmark

Correspondence e-mail: adb@chem.sdu.dk

## Key indicators

Single-crystal X-ray study  
 $T = 180$  K  
Mean  $\sigma(C-C) = 0.002$  Å  
 $R$  factor = 0.037  
 $wR$  factor = 0.112  
Data-to-parameter ratio = 25.4For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.5-*tert*-Butyl-2-(*N,N*-dimethylaminocarbonylthio)-  
isophthalaldehydeThe *N,N*-dimethylaminocarbonylthio moiety in the title  
compound,  $C_{15}H_{19}NO_3S$ , adopts a staggered conformation  
with respect to the *tert*-butyl substituent in the crystal  
structure at 180 K.Received 29 July 2004  
Accepted 5 August 2004  
Online 13 August 2004

## Comment

The crystal structure of the title compound, (I), is the first  
example of a *tert*-butyl-substituted isophthalaldehyde and one  
of very few structures to date that contain the *N,N*-di-  
methylaminocarbonylthio moiety bound to a benzene ring  
(Bennett *et al.*, 1999; Higgs & Carrano, 2002; Gibbs *et al.*,  
1995). The approximately planar *N,N*-dimethylamino-  
carbonylthio group adopts a dihedral angle of  $69.0(1)^\circ$  to the  
plane of the benzene ring so that it forms a 'staggered'  
orientation with respect to the *tert*-butyl group (Fig. 2). The  
benzene rings of molecules related by translation along [100]  
are close to coplanar and form pairs of  $C-H \cdots O$  contacts  
between the aldehyde groups (Fig. 3).

## Experimental

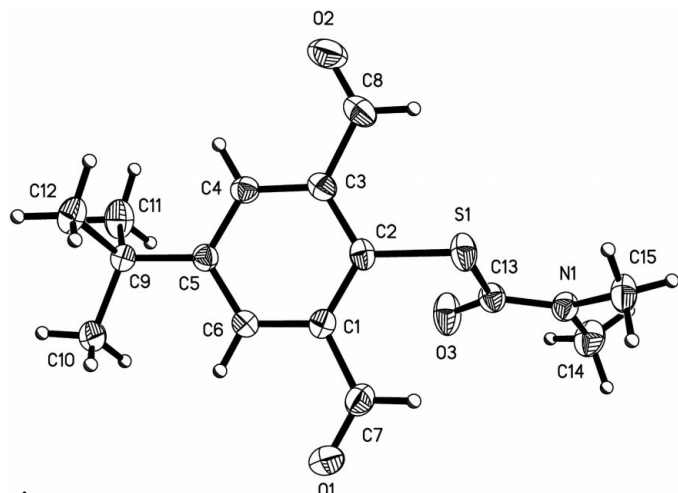
The title compound was prepared according to the method of  
Brooker *et al.* (2000).

## Crystal data

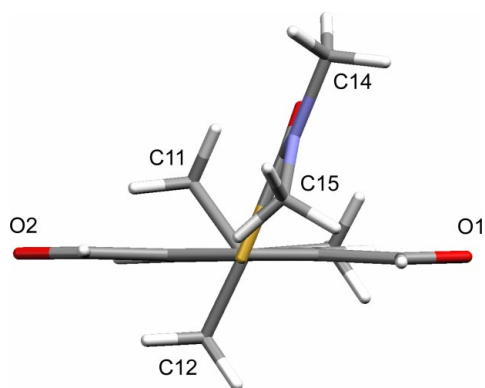
 $C_{15}H_{19}NO_3S$   
 $M_r = 293.37$   
Monoclinic,  $P2_1/c$   
 $a = 8.5830(2)$  Å  
 $b = 18.5242(4)$  Å  
 $c = 9.3889(2)$  Å  
 $\beta = 93.359(1)^\circ$   
 $V = 1490.21(6)$  Å<sup>3</sup>  
 $Z = 4$  $D_x = 1.308$  Mg m<sup>-3</sup>  
Mo  $K\alpha$  radiation  
Cell parameters from 7497  
reflections  
 $\theta = 2.4-29.8^\circ$   
 $\mu = 0.22$  mm<sup>-1</sup>  
 $T = 180(2)$  K  
Block, colourless  
 $0.40 \times 0.30 \times 0.10$  mm

## Data collection

Bruker-Nonius X8APEX-II CCD  
diffractometer  
Thin-slice  $\omega$  and  $\varphi$  scans  
Absorption correction: multi-scan  
(SADABS; Sheldrick, 2003)  
46 511 measured reflections  
4716 independent reflections3651 reflections with  $I > 2\sigma(I)$   
 $R_{int} = 0.031$   
 $\theta_{max} = 31.0^\circ$   
 $h = -12 \rightarrow 12$   
 $k = -26 \rightarrow 26$   
 $l = -13 \rightarrow 13$



**Figure 1**  
The molecular structure, showing displacement ellipsoids at the 50% probability level. H atoms are shown as spheres of arbitrary radii.



**Figure 2**  
Projection along the C2...C5 vector, showing the staggered arrangement of the *N,N*-dimethylaminocarbonylthio group with respect to the *tert*-butyl substituent.

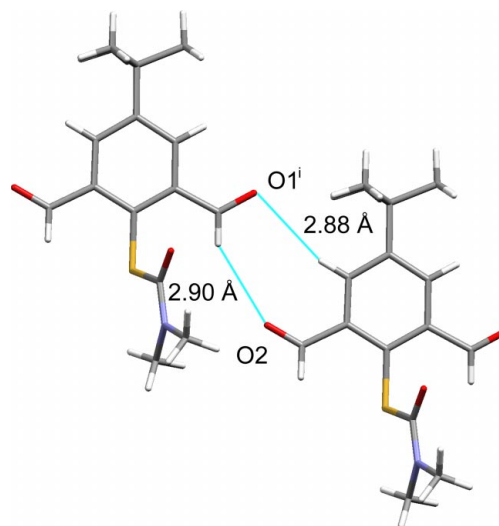
#### Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.037$   
 $wR(F^2) = 0.112$   
 $S = 1.07$   
 4716 reflections  
 186 parameters  
 H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0636P)^2 + 0.1853P]$$

where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} = 0.001$   
 $\Delta\rho_{\max} = 0.40 \text{ e } \text{\AA}^{-3}$   
 $\Delta\rho_{\min} = -0.17 \text{ e } \text{\AA}^{-3}$

H atoms were positioned geometrically and allowed to ride during subsequent refinement, with C–H = 0.95 Å and  $U_{\text{iso}}(\text{H}) = 1.2 U_{\text{eq}}(\text{C})$  for  $\text{Csp}^2$  atoms, and C–H = 0.98 Å and  $U_{\text{iso}}(\text{H}) = 1.5 U_{\text{eq}}(\text{C})$  for  $\text{Csp}^3$ . The methyl groups were allowed to rotate about their local threefold axes.



**Figure 3**  
Pairs of C–H...O contacts formed between aldehyde groups. [Symmetry code: (i) 1 + x, y, z.]

Data collection: *APEX2* (Bruker–Nonius, 2003); cell refinement: *SAINT* (Bruker, 2003); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 2000); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

The authors are grateful to the Danish Natural Science Research Council (SNF) and Carlsbergfondet for provision of the X-ray equipment.

#### References

- Bennett, S. M. W., Brown, S. M., Conole, G., Dennis, M. R., Fraser, P. K., Radojevic, A., McPartlin, M., Topping, C. M. & Woodward, S. (1999). *J. Chem. Soc. Perkin Trans. 1*, pp. 3127–3132.
- Brooker, S., Caygill, G. B., Croucher, P. D., Davidson, T. C., Clive, D. L. J., Magnuson, S. R., Cramer, S. P. & Ralston, C. Y. (2000). *J. Chem. Soc. Dalton Trans.* pp. 3113–3121.
- Bruker (2003). *SAINT*. Version 7.06a. Bruker AXS Inc., Madison, Wisconsin, USA.
- Bruker–Nonius (2003). *APEX2*. Version 1.0–8. Bruker Nonius BV, Delft, The Netherlands.
- Gibbs, C. G., Sujeth, P. K., Rogers, J. S., Stanley, G. G., Krawiec, M., Watson, W. H. & Gutsche, C. D. (1995). *J. Org. Chem.* **60**, 8394–8402.
- Higgs, T. C. & Carrano, C. J. (2002). *Eur. J. Org. Chem.* pp. 3632–3645.
- Sheldrick, G. M. (2000). *SHELXTL*. Version 6.10. Bruker AXS Inc., Madison, Wisconsin, USA.
- Sheldrick, G. M. (2003). *SADABS*. Version 2.10. Bruker AXS Inc., Madison, Wisconsin, USA.