Acta Crystallographica Section E Structure Reports Online ISSN 1600-5368 Editors: W. Clegg and D. G. Watson

3,5-Di-tert-butyl-2-hydroxybenzaldehyde

Duncan M. Tooke and Anthony L. Spek

Copyright © International Union of Crystallography

Author(s) of this paper may load this reprint on their own web site provided that this cover page is retained. Republication of this article or its storage in electronic databases or the like is not permitted without prior permission in writing from the IUCr.

organic papers

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

Duncan M. Tooke* and Anthony L. Spek

Bijvoet Center for Biomolecular Research, Department of Crystal and Structural Chemistry, Utrecht University, Padualaan 8, 3584 CH Utrecht, The Netherlands

Correspondence e-mail: d.m.tooke@chem.uu.nl

Key indicators

Single-crystal X-ray study T = 150 K Mean σ (C–C) = 0.002 Å Disorder in main residue R factor = 0.053 wR factor = 0.146 Data-to-parameter ratio = 16.2

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

3,5-Di-tert-butyl-2-hydroxybenzaldehyde

The title compound, $C_{15}H_{22}O_2$, crystallizes in the monoclinic space group $P2_1/c$, with two molecules in the asymmetric unit related by a non-crystallographic inversion centre. The O- $H \cdots O$ hydrogen bonding is exclusively intramolecular.

Comment

During the course of research into metal complexes with extended chelating ligands, the structure of the title compound, (I), was determined (Fig. 1).



The asymmetric unit contains two crystallographically independent molecules, one of which includes a disordered *tert*-butyl group, with relative occupancies of 69.6 (3) and 30.4 (3)% for the two components. Small (10 Å³) voids are found close to the disordered moieties.

A local pseudo-inversion centre is located between the two crystallographically independent molecules, and is broken by the disorder of the *tert*-butyl group involving atom C23. It is, in fact, the major component of the disorder that breaks the symmetry, and not the minor (Fig. 2).

The only strong hydrogen bonds are intramolecular, between H2 and O1, and between H4 and O3, and the molecules pack in layers parallel to the (101) plane in the crystal structure (Fig. 3).



Figure 1

The asymmetric unit of the title compound, with the atom-numbering scheme. Both disorder components are included. Dashed bonds represent hydrogen bonds. Displacement ellipsoids for non-H atoms are drawn at the 50% probability level.

 \odot 2004 International Union of Crystallography Printed in Great Britain – all rights reserved Received 3 March 2004

Accepted 5 April 2004

Online 17 April 2004

Experimental

The title compound, recovered unchanged from a reaction mixture in which it was one of the intended reagents, was recrystallized from hot methanol, giving large good-quality single crystals.

 $D_x = 1.115 \text{ Mg m}^{-3}$

Cell parameters from 56

Mo $K\alpha$ radiation

reflections

 $\mu = 0.07 \text{ mm}^{-1}$

Block, colourless

 $0.30 \times 0.10 \times 0.10 \ \mathrm{mm}$

 $\theta = 4.0 - 18.3^{\circ}$

T = 150 K

Crystal data

C15H22O2 $M_r = 234.33$ Monoclinic, $P2_1/c$ a = 11.627 (1) Åb = 10.1214 (7) Å c = 25.461 (2) Å $\beta = 111.326 \ (3)^{\circ}$ V = 2791.1 (4) Å³ Z = 8

Data collection

Nonius KappaCCD diffractometer	$R_{\rm int} = 0.063$
φ and ω scans	$\theta_{\rm max} = 26.5^{\circ}$
Absorption correction: none	$h = -14 \rightarrow 14$
28855 measured reflections	$k = -12 \rightarrow 12$
5782 independent reflections	$l = -31 \rightarrow 31$
4178 reflections with $L > 2\sigma(I)$	

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0694P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.053$	+ 1.0702P]
$wR(F^2) = 0.146$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.04	$(\Delta/\sigma)_{\rm max} < 0.001$
5782 reflections	$\Delta \rho_{\rm max} = 0.36 \text{ e } \text{\AA}^{-3}$
358 parameters	$\Delta \rho_{\rm min} = -0.20 \text{ e } \text{\AA}^{-3}$
H atoms treated by a mixture of	
independent and constrained	
refinement	

Table 1

Selected geometric parameters (Å, °).

O3-C16	1.227 (2)	01-C1	1.226 (2)
O4-C22	1.3560 (18)	02-C7	1.3584 (18)
O3-C16-C17	125.00 (16)	O1-C1-C2	125.03 (16)
O4-C22-C21	119.60 (14)	O2-C7-C6	119.89 (14)
O4-C22-C17	120.03 (14)	O2-C7-C2	120.05 (14)

Table 2



$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
O2−H2···O1	0.88 (2)	1.80 (2)	2.6122 (18)	151.5 (19)
$O4-H4\cdots O3$	0.88 (3)	1.79 (2)	2.6075 (18)	153.7 (19)

Atoms H2 and H4 were refined freely with individual isotropic displacement parameters. All remaining H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms, with C-H = 0.96 Å and $U_{iso}(H) = 1.5U_{eq}(C)$ for methyl H atoms, and C-H = 0.95 Å and $U_{iso}(H) = 1.2U_{eq}(C)$ for all other H atoms. The tert-butyl group containing atom C23 was rotationally disordered over two positions. The final occupancies of the two sites were 69.6 (3) and 30.4 (3)% for the major and minor components, respectively.



Figure 2

Plot of the fit between the two molecules related by a pseudo-inversion centre (one molecule has been inverted). It is the major component that breaks the symmetry, and not the minor.



Figure 3

Projection of the structure down the b axis, demonstrating the packing of the two crystallographically independent molecules in layers parallel to the (101) plane.

Data collection: COLLECT (Nonius, 1998); cell refinement: DIRAX (Duisenberg, 1992); data reduction: EVAL14 (Duisenberg et al., 2003); program(s) used to solve structure: SHELXS86 (Sheldrick, 1985); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: PLATON (Spek, 2003); software used to prepare material for publication: PLATON.

This work was supported in part (ALS) by the Council for the Chemical Sciences of the Netherlands Organization for Scientific Research (CW-NWO). The crystals were prepared and kindly donated by Dr K. Ramu, University of Leiden, The Netherlands.

References

- Duisenberg, A. J. M. (1992). J. Appl. Cryst. 25, 92-96.
- Duisenberg, A. J. M., Kroon-Batenburg, L. M. J. & Schreurs, A. M. M. (2003), J. Appl. Cryst. 36, 220-229.
- Nonius (1998). COLLECT. Nonius BV, Delft, The Netherlands.
- Sheldrick, G. M. (1985). SHELXS86. University of Göttingen, Germany. Sheldrick, G. M. (1997). SHELXL97. University of Göttingen, Germany. Spek, A. L. (2003). J. Appl. Cryst. 36, 7-13.