



Crystal structure of 5,5'-dibromo-3,3'-di-*tert*-butyl-6,6'-dimethylbiphenyl-2,2'-diol

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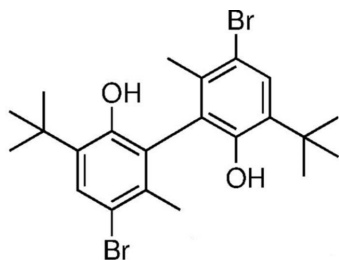
The whole molecule of the title compound, C₂₂H₂₈Br₂O₂, is generated by twofold rotation symmetry. The dihedral angle of the biphenyl moiety is 85.05 (11)°. The hydroxy groups show intramolecular O—H... π interactions without any other hydrogen-bond acceptors. In the crystal, there are no other significant intermolecular interactions present.

Keywords: crystal structure; biphenyl; axial chirality; O—H... π interactions.

CCDC reference: 1056738

1. Related literature

For the synthesis of the title compound using a transition-metal catalyst, see: Kubota *et al.* (2012). For the determination of the absolute configuration of the corresponding (+)-chloro derivative, *viz.* *S*, see: Gutierrez *et al.* (2010). For the crystal structure of a similar compound, *i.e.* 5,5'-dimethoxy-6,6'-dimethylbiphenyl-2,2'-diol dichloromethane solvate, see: Guo *et al.* (2011).



2. Experimental

2.1. Crystal data

C₂₂H₂₈Br₂O₂
M_r = 484.24
 Orthorhombic, *Pba*2
a = 7.3680 (5) Å
b = 22.4243 (14) Å
c = 6.6148 (4) Å

V = 1092.91 (12) Å³
Z = 2
 Mo *K*α radiation
 μ = 3.72 mm⁻¹
T = 299 K
 0.16 × 0.15 × 0.10 mm

2.2. Data collection

Bruker D8 VENTURE
 diffractometer
 Absorption correction: multi-scan
 (*SADABS*; Bruker, 2014)
T_{min} = 0.630, *T_{max}* = 0.773

9406 measured reflections
 1966 independent reflections
 1615 reflections with *I* > 2σ(*I*)
R_{int} = 0.036

2.3. Refinement

$R[F^2 > 2\sigma(F^2)] = 0.031$
 $wR(F^2) = 0.065$
S = 1.06
 1966 reflections
 123 parameters
 1 restraint
 H-atom parameters constrained
 $\Delta\rho_{\max} = 0.48 \text{ e \AA}^{-3}$

$\Delta\rho_{\min} = -0.21 \text{ e \AA}^{-3}$
 Absolute structure: Flack *x*
 determined using 621 quotients
 [(*I*⁺) − (*I*[−])] / [(*I*⁺) + (*I*[−])] (Parsons
et al., 2013)
 Absolute structure parameter:
 0.034 (9)

Table 1

Hydrogen-bond geometry (Å, °).

*C*_g is the centroid of benzene ring C3–C8.

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
O2—H2... <i>C</i> _g ⁱ	0.82	2.54	3.047 (5)	122

Symmetry code: (i) $-x + 2, -y + 2, z$.

Data collection: *APEX2* (Bruker, 2014); cell refinement: *SAINTE* (Bruker, 2014); data reduction: *SAINTE*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL2014* (Sheldrick, 2015); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXL2014* and *PUBLICIF* (Westrip, 2010).

Acknowledgements

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Supporting information for this paper is available from the IUCr electronic archives (Reference: SU5104).

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supporting information

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Crystal structure of 5,5'-dibromo-3,3'-di-*tert*-butyl-6,6'-dimethylbiphenyl-2,2'-diol

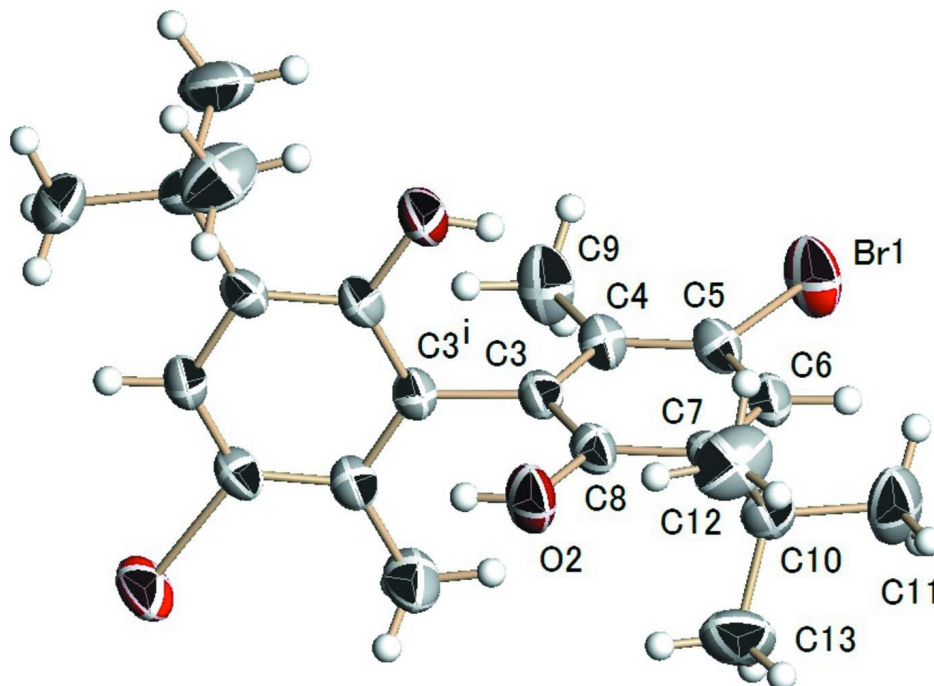
Rika Obata, Shigeru Ohba, Yasuaki Einaga and Shigeru Nishiyama

S1. Synthesis and crystallization

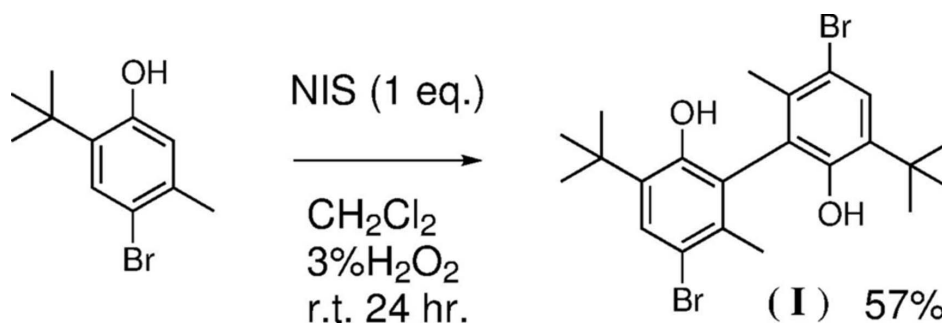
The synthesis of the title compound, (I), is described in Fig. 2. It was prepared using iodine-mediated coupling method from 4-bromo-2-*tert*-butyl-5-methylphenol. To the solution of 4-bromo-2-*tert*-butyl-5-methylphenol (0.242 g, 1 mmol) in dichloromethane (1 mL) was added *N*-iodosuccinimide (abbreviated to NIS, 0.225 g, 1 mmol) and 3% H₂O₂ (1 mL). After shaking (200 rpm) the reaction mixture for 24 h at room temperature, it was poured into saturated Na₂S₂O₃ solution, and extracted with chloroform. The organic layer was washed with saturated NaCl and dried over anhydrous Na₂SO₄. The mixture was evaporated and purified by silica-gel column chromatography to give title compound (I) as white solid (yield: 0.138 g, 57%). ¹H-NMR (400 MHz, CDCl₃) 1.32 (18H, s), 1.92 (6H, s), 4.80 (2H, s), 7.47 (2H, s). Tof-MS ES(-) Anal. 481.0357, Calcd. 481.0378 for C₂₂H₂₇O₂Br₂. The crystals were grown by slow evaporation from a toluene/*n*-hexane (1/4) solution.

S2. Refinement

Crystal data, data collection and structure refinement details are summarized in the experimental table. The hydroxyl H atom was located from a difference Fourier map but was refined as riding (AFIX 147) with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{O})$. C-Bound H atoms were included in calculated positions and refined as riding: C—H = 0.93–0.96 Å with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$ for methyl H atoms and $1.2U_{\text{eq}}(\text{C})$ for other H atoms.

**Figure 1**

The molecular structure of the title compound, with atom labelling. Displacement ellipsoids are drawn at the 50% probability level.

**Figure 2**

The synthesis of the title compound, (I).

5,5'-Dibromo-3,3'-di-*tert*-butyl-6,6'-dimethylbiphenyl-2,2'-diol

Crystal data

$C_{22}H_{28}Br_2O_2$

$M_r = 484.24$

Orthorhombic, *Pba2*

$a = 7.3680$ (5) Å

$b = 22.4243$ (14) Å

$c = 6.6148$ (4) Å

$V = 1092.91$ (12) Å³

$Z = 2$

$F(000) = 492$

$D_x = 1.472$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 4121 reflections

$\theta = 2.9$ – 23.6°

$\mu = 3.72$ mm⁻¹

$T = 299$ K

Prism, colourless

$0.16 \times 0.15 \times 0.10$ mm

Data collection

Bruker D8 VENTURE
diffractometer
Radiation source: fine-focus sealed tube
 ω scans
Absorption correction: multi-scan
(SADABS; Bruker, 2014)
 $T_{\min} = 0.630$, $T_{\max} = 0.773$
9406 measured reflections

1966 independent reflections
1615 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.036$
 $\theta_{\max} = 25.3^\circ$, $\theta_{\min} = 2.9^\circ$
 $h = -8 \rightarrow 8$
 $k = -26 \rightarrow 26$
 $l = -7 \rightarrow 7$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.031$
 $wR(F^2) = 0.065$
 $S = 1.06$
1966 reflections
123 parameters
1 restraint
Primary atom site location: structure-invariant
direct methods

Hydrogen site location: inferred from
neighbouring sites
H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0176P)^2 + 0.3778P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.48 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.21 \text{ e } \text{\AA}^{-3}$
Absolute structure: Flack x determined using
621 quotients $[(F^-)-(I)]/[(F^+)+(I)]$ (Parsons *et al.*,
2013)
Absolute structure parameter: 0.034 (9)

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
Br1	1.18668 (8)	0.81211 (2)	0.48288 (15)	0.0681 (2)
O2	0.7528 (4)	0.98204 (13)	0.9897 (9)	0.0576 (8)
H2	0.7777	1.0166	0.9607	0.086*
C3	0.9856 (6)	0.96676 (17)	0.7429 (7)	0.0352 (10)
C4	1.0903 (6)	0.9290 (2)	0.6223 (7)	0.0388 (11)
C5	1.0536 (6)	0.8689 (2)	0.6397 (7)	0.0399 (11)
C6	0.9212 (6)	0.84644 (19)	0.7649 (7)	0.0394 (11)
H6	0.9023	0.8054	0.7685	0.047*
C7	0.8150 (6)	0.88292 (19)	0.8859 (7)	0.0360 (10)
C8	0.8518 (6)	0.94415 (19)	0.8707 (7)	0.0370 (11)
C9	1.2322 (7)	0.9538 (2)	0.4852 (15)	0.0650 (14)
H9A	1.1993	0.9460	0.3473	0.098*
H9B	1.3467	0.9353	0.5143	0.098*
H9C	1.2417	0.9960	0.5058	0.098*
C10	0.6668 (5)	0.8582 (2)	1.0234 (8)	0.0428 (15)
C11	0.6564 (7)	0.7905 (2)	1.0111 (14)	0.074 (2)
H11A	0.6243	0.7789	0.8761	0.111*
H11B	0.5661	0.7762	1.1039	0.111*
H11C	0.7722	0.7737	1.0456	0.111*

C12	0.7053 (8)	0.8746 (3)	1.2449 (9)	0.0756 (18)
H12A	0.8260	0.8623	1.2802	0.113*
H12B	0.6195	0.8548	1.3311	0.113*
H12C	0.6944	0.9170	1.2619	0.113*
C13	0.4827 (6)	0.8829 (2)	0.9625 (15)	0.0760 (17)
H13A	0.4791	0.9249	0.9894	0.114*
H13B	0.3893	0.8632	1.0387	0.114*
H13C	0.4634	0.8760	0.8208	0.114*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Br1	0.0866 (4)	0.0458 (3)	0.0718 (4)	0.0118 (2)	0.0311 (4)	-0.0118 (4)
O2	0.0646 (19)	0.0366 (16)	0.072 (2)	0.0049 (14)	0.018 (3)	-0.009 (3)
C3	0.040 (3)	0.030 (2)	0.036 (3)	0.0016 (19)	-0.004 (2)	0.001 (2)
C4	0.048 (3)	0.033 (3)	0.035 (3)	0.003 (2)	0.002 (2)	0.000 (2)
C5	0.048 (3)	0.035 (3)	0.037 (3)	0.009 (2)	0.003 (2)	-0.003 (2)
C6	0.046 (3)	0.028 (2)	0.045 (3)	0.002 (2)	-0.002 (2)	-0.002 (2)
C7	0.036 (3)	0.034 (2)	0.038 (2)	0.008 (2)	-0.005 (2)	-0.002 (2)
C8	0.042 (3)	0.031 (3)	0.038 (2)	0.006 (2)	0.000 (2)	-0.005 (2)
C9	0.084 (3)	0.048 (3)	0.063 (3)	0.003 (2)	0.033 (5)	-0.003 (5)
C10	0.035 (3)	0.045 (3)	0.048 (4)	-0.0010 (19)	0.008 (3)	0.003 (2)
C11	0.070 (3)	0.048 (3)	0.104 (6)	-0.013 (2)	0.036 (5)	0.008 (4)
C12	0.078 (4)	0.099 (5)	0.049 (4)	-0.021 (4)	0.010 (3)	0.002 (3)
C13	0.039 (3)	0.086 (4)	0.103 (5)	0.005 (2)	0.012 (4)	0.020 (5)

Geometric parameters (Å, °)

Br1—C5	1.913 (4)	C9—H9B	0.9600
O2—C8	1.369 (6)	C9—H9C	0.9600
O2—H2	0.8200	C10—C13	1.520 (7)
C3—C8	1.394 (6)	C10—C11	1.521 (7)
C3—C4	1.396 (6)	C10—C12	1.537 (8)
C3—C3 ⁱ	1.506 (8)	C11—H11A	0.9600
C4—C5	1.379 (6)	C11—H11B	0.9600
C4—C9	1.491 (8)	C11—H11C	0.9600
C5—C6	1.375 (6)	C12—H12A	0.9600
C6—C7	1.386 (6)	C12—H12B	0.9600
C6—H6	0.9300	C12—H12C	0.9600
C7—C8	1.403 (6)	C13—H13A	0.9600
C7—C10	1.526 (6)	C13—H13B	0.9600
C9—H9A	0.9600	C13—H13C	0.9600
C8—O2—H2	109.5	C13—C10—C11	107.7 (4)
C8—C3—C4	121.1 (4)	C13—C10—C7	110.4 (5)
C8—C3—C3 ⁱ	117.3 (4)	C11—C10—C7	111.5 (4)
C4—C3—C3 ⁱ	121.6 (4)	C13—C10—C12	109.3 (5)
C5—C4—C3	115.9 (4)	C11—C10—C12	107.4 (5)

C5—C4—C9	123.5 (4)	C7—C10—C12	110.5 (4)
C3—C4—C9	120.5 (4)	C10—C11—H11A	109.5
C6—C5—C4	123.2 (4)	C10—C11—H11B	109.5
C6—C5—Br1	116.5 (3)	H11A—C11—H11B	109.5
C4—C5—Br1	120.3 (3)	C10—C11—H11C	109.5
C5—C6—C7	122.2 (4)	H11A—C11—H11C	109.5
C5—C6—H6	118.9	H11B—C11—H11C	109.5
C7—C6—H6	118.9	C10—C12—H12A	109.5
C6—C7—C8	115.3 (4)	C10—C12—H12B	109.5
C6—C7—C10	122.2 (4)	H12A—C12—H12B	109.5
C8—C7—C10	122.5 (4)	C10—C12—H12C	109.5
O2—C8—C3	120.0 (4)	H12A—C12—H12C	109.5
O2—C8—C7	117.6 (4)	H12B—C12—H12C	109.5
C3—C8—C7	122.4 (4)	C10—C13—H13A	109.5
C4—C9—H9A	109.5	C10—C13—H13B	109.5
C4—C9—H9B	109.5	H13A—C13—H13B	109.5
H9A—C9—H9B	109.5	C10—C13—H13C	109.5
C4—C9—H9C	109.5	H13A—C13—H13C	109.5
H9A—C9—H9C	109.5	H13B—C13—H13C	109.5
H9B—C9—H9C	109.5		
C8—C3—C4—C5	-0.5 (6)	C3 ⁱ —C3—C8—O2	0.0 (6)
C3 ⁱ —C3—C4—C5	178.5 (4)	C4—C3—C8—C7	0.1 (7)
C8—C3—C4—C9	179.6 (5)	C3 ⁱ —C3—C8—C7	-179.0 (4)
C3 ⁱ —C3—C4—C9	-1.4 (7)	C6—C7—C8—O2	-178.9 (4)
C3—C4—C5—C6	0.9 (7)	C10—C7—C8—O2	1.9 (6)
C9—C4—C5—C6	-179.3 (6)	C6—C7—C8—C3	0.1 (6)
C3—C4—C5—Br1	179.9 (3)	C10—C7—C8—C3	-179.1 (4)
C9—C4—C5—Br1	-0.2 (7)	C6—C7—C10—C13	-118.6 (5)
C4—C5—C6—C7	-0.8 (7)	C8—C7—C10—C13	60.5 (6)
Br1—C5—C6—C7	-179.8 (3)	C6—C7—C10—C11	1.0 (6)
C5—C6—C7—C8	0.3 (7)	C8—C7—C10—C11	-179.8 (5)
C5—C6—C7—C10	179.4 (4)	C6—C7—C10—C12	120.4 (5)
C4—C3—C8—O2	179.0 (4)	C8—C7—C10—C12	-60.5 (6)

Symmetry code: (i) $-x+2, -y+2, z$.

Hydrogen-bond geometry (\AA , $^\circ$)

C_g is the centroid of benzene ring C3—C8.

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
O2—H2 \cdots C _g ⁱ	0.82	2.54	3.047 (5)	122

Symmetry code: (i) $-x+2, -y+2, z$.