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1-Bromo-2,3,6-trichloro-4,5-dimethoxybenzene

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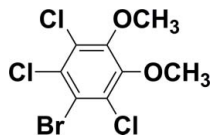
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 Key indicators: single-crystal X-ray study; $T = 90$ K; mean $\sigma(\text{C}-\text{C}) = 0.005$ Å; disorder in main residue; R factor = 0.038; wR factor = 0.103; data-to-parameter ratio = 13.2.

The halogen atoms of the title compound, $\text{C}_8\text{H}_6\text{BrCl}_3\text{O}_2$, are located within the plane of the benzene ring [r.m.s. deviation = 0.036 (11) Å]. The two methoxy groups are twisted out of this plane, with dihedral angles of 84.7 (3) and 68.5 (3)°, and point in opposite directions. The structure is disordered by a non-crystallographic twofold rotation which superimposes Cl and Br at two of the halogen sites. The refined occupancies for the major and minor components are 0.517 (2) and 0.483 (2).

Related literature

For similar structures of halogenated methoxy benzenes, see: Iimura *et al.* (1984); Rissanen *et al.* (1987, 1988*a,b*); Song *et al.* (2008, 2010); Telu *et al.* (2008); Weller & Gerstner (1995); Wieczorek (1980). For background to halogenated methoxy benzenes, see: Brownlee *et al.* (1993); Curtis *et al.* (1972); Pereira *et al.* (2000); Vlachos *et al.* (2007); Zhang *et al.* (2006).



Experimental

Crystal data

$\text{C}_8\text{H}_6\text{BrCl}_3\text{O}_2$
 $M_r = 320.39$
 Triclinic, $P\bar{1}$

$a = 7.7885$ (7) Å
 $b = 8.8600$ (7) Å
 $c = 9.1523$ (8) Å

$\alpha = 62.256$ (3)°
 $\beta = 75.358$ (4)°
 $\gamma = 75.133$ (4)°
 $V = 533.64$ (8) Å³
 $Z = 2$

Cu $K\alpha$ radiation
 $\mu = 11.94$ mm⁻¹
 $T = 90$ K
 $0.20 \times 0.15 \times 0.07$ mm

Data collection

Bruker X8 Proteum diffractometer
 Absorption correction: multi-scan
 (SADABS; Bruker, 2006)
 $T_{\min} = 0.193$, $T_{\max} = 0.434$

6588 measured reflections
 1828 independent reflections
 1720 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.038$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.038$
 $wR(F^2) = 0.103$
 $S = 1.12$
 1828 reflections
 138 parameters

6 restraints
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.35$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.52$ e Å⁻³

Data collection: APEX2 (Bruker, 2006); cell refinement: SAINT (Bruker, 2006); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: XP in SHELXTL (Sheldrick, 2008); software used to prepare material for publication: SHELXL97 and local procedures.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: OM2317).

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supporting information

Acta Cryst. (2010). E66, o487 [https://doi.org/10.1107/S1600536810002977]

1-Bromo-2,3,6-trichloro-4,5-dimethoxybenzene

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S1. Comment

Chlorinated methoxy benzenes are a group of persistent organic pollutants that are associated with off-flavors in water, fish, chicken and wine (Brownlee *et al.*, 1993; Curtis *et al.*, 1972; Pereira *et al.*, 2000; Vlachos *et al.*, 2007; Zhang *et al.*, 2006). Their biological properties depend, at least in part, on the conformation of the methoxy group relative to the aromatic ring system. The two methoxy groups of the title compound are twisted out of the plane of the benzene ring system due to the bulky *ortho* chlorine substituents and point in opposite directions. The respective dihedral angles were calculated between the plane of the benzene ring (C1 through C6) and the methoxy group and are 84.7 (3)° (atoms C1,O1,C7) and 68.5 (3)° (atoms C2,O2,C8), respectively. These dihedral angles are in agreement with the dihedral angles observed for other chlorinated methoxy benzenes with two *ortho* substituents (Iimura *et al.*, 1984; Rissanen *et al.*, 1987; Rissanen *et al.*, 1988b; Telu *et al.*, 2008; Weller & Gerstner, 1995; Wiczorek, 1980). In contrast, the methoxy group of structurally related compound with no or one substituent *ortho* to the methoxy group typically lie within the plane of the benzene ring system (Rissanen *et al.*, 1988a; Song *et al.*, 2010).

S2. Experimental

This title compound was synthesized by chlorination of 1-bromo-3,4-dimethoxy-benzene with HCl/H₂O₂ as described previously (Song *et al.*, 2008). Crystals suitable for X-ray diffraction were grown by slow evaporation of a saturated solution of the title compound in CHCl₃.

S3. Refinement

H atoms were found in difference Fourier maps and subsequently placed in idealized positions with constrained C-H distances of 0.98 Å and U_{iso}(H) values set to 1.5U_{eq} of the attached C atom. The structure is disordered by a non-crystallographic 2-fold rotation about an axis that bisects the midpoints of bonds C1—C2 and C4—C5. This disorder superimposes Cl and Br at the halogen sites bonded to C4 and C5. The refined occupancies for the major and minor components are 0.517 (2) and 0.483 (2). To ensure a physically/chemically sensible model in spite of the disorder, the bond distances for C—Cl were restrained to a refined variable, 1.713 (8). The C—Br bond distance was restrained to the same variable, but multiplied by 1.096, which is the ratio of C—Br:C—Cl for this type of bond. Six restraints in total were required. In addition, the displacement parameters of the superimposed atoms were constrained to be the same.

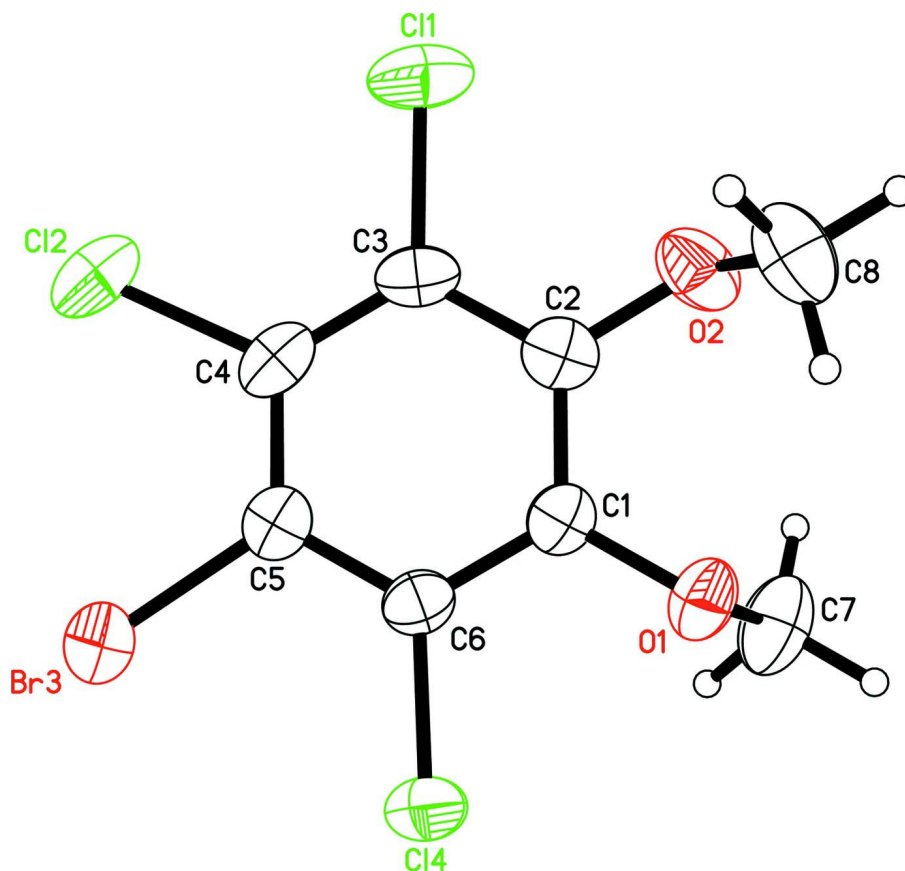


Figure 1

View of the title compound showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level.

1-Bromo-2,3,6-trichloro-4,5-dimethoxybenzene

Crystal data

$C_8H_6BrCl_3O_2$

$M_r = 320.39$

Triclinic, $P\bar{1}$

Hall symbol: -P 1

$a = 7.7885$ (7) Å

$b = 8.8600$ (7) Å

$c = 9.1523$ (8) Å

$\alpha = 62.256$ (3)°

$\beta = 75.358$ (4)°

$\gamma = 75.133$ (4)°

$V = 533.64$ (8) Å³

$Z = 2$

$F(000) = 312$

$D_x = 1.994$ Mg m⁻³

Cu $K\alpha$ radiation, $\lambda = 1.54178$ Å

Cell parameters from 5684 reflections

$\theta = 5.5$ – 66.1 °

$\mu = 11.94$ mm⁻¹

$T = 90$ K

Irregular plate, colourless

$0.20 \times 0.15 \times 0.07$ mm

Data collection

Bruker X8 Proteum
diffractometer

Radiation source: fine-focus rotating anode

Graded multilayer optics monochromator

Detector resolution: 18 pixels mm⁻¹

φ and ω scans

Absorption correction: multi-scan
(*SADABS*; Bruker, 2006)

$T_{\min} = 0.193$, $T_{\max} = 0.434$

6588 measured reflections

1828 independent reflections

1720 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.038$
 $\theta_{\text{max}} = 67.4^\circ$, $\theta_{\text{min}} = 5.5^\circ$
 $h = -9 \rightarrow 7$

$k = -10 \rightarrow 10$
 $l = -10 \rightarrow 10$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.038$
 $wR(F^2) = 0.103$
 $S = 1.12$
 1828 reflections
 138 parameters
 6 restraints
 Primary atom site location: structure-invariant
 direct methods
 Secondary atom site location: difference Fourier
 map

Hydrogen site location: inferred from
 neighbouring sites
 H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0466P)^2 + 0.726P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.004$
 $\Delta\rho_{\text{max}} = 0.35 \text{ e } \text{Å}^{-3}$
 $\Delta\rho_{\text{min}} = -0.52 \text{ e } \text{Å}^{-3}$
 Extinction correction: *SHELXL97* (Sheldrick,
 2008), $F_c^* = kFc[1 + 0.001x\text{Fc}^2\lambda^3/\sin(2\theta)]^{-1/4}$
 Extinction coefficient: 0.0232 (18)

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.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > 2\sigma(F^2)$ is used only for calculating R -factors(gt) *etc.* and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
C1	0.2391 (4)	0.6704 (4)	0.7165 (4)	0.0318 (6)	
C2	0.2512 (4)	0.7913 (5)	0.7681 (4)	0.0360 (7)	
C3	0.2598 (4)	0.9603 (4)	0.6519 (4)	0.0356 (7)	
C4	0.2536 (4)	1.0107 (4)	0.4851 (4)	0.0366 (7)	
C5	0.2393 (4)	0.8899 (4)	0.4345 (4)	0.0350 (7)	
C6	0.2314 (4)	0.7212 (4)	0.5504 (4)	0.0329 (7)	
O1	0.2295 (3)	0.5051 (3)	0.8291 (3)	0.0418 (6)	
C7	0.3984 (6)	0.3991 (5)	0.8622 (6)	0.0592 (11)	
H7A	0.4694	0.3956	0.7585	0.089*	
H7B	0.3800	0.2818	0.9444	0.089*	
H7C	0.4627	0.4464	0.9066	0.089*	
O2	0.2646 (3)	0.7442 (4)	0.9285 (3)	0.0486 (6)	
C8	0.1070 (5)	0.6975 (6)	1.0458 (5)	0.0558 (10)	
H8A	0.0096	0.7968	1.0188	0.084*	
H8B	0.1309	0.6619	1.1584	0.084*	
H8C	0.0717	0.6014	1.0412	0.084*	
Cl1	0.27710 (13)	1.10613 (13)	0.71869 (14)	0.0567 (3)	
Cl2	0.265 (3)	1.2206 (17)	0.326 (2)	0.0539 (7)	0.483 (3)
Br3	0.2388 (9)	0.9427 (6)	0.2147 (7)	0.0495 (7)	0.483 (3)
Br2	0.2599 (11)	1.2383 (6)	0.3376 (8)	0.0539 (7)	0.517 (3)

C13	0.228 (2)	0.9701 (14)	0.2211 (15)	0.0495 (7)	0.517 (3)
C14	0.20968 (13)	0.56783 (11)	0.49546 (11)	0.0485 (3)	

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0236 (14)	0.0338 (16)	0.0354 (16)	-0.0092 (12)	-0.0006 (11)	-0.0123 (13)
C2	0.0257 (15)	0.0442 (18)	0.0406 (17)	-0.0106 (13)	0.0000 (12)	-0.0203 (15)
C3	0.0255 (15)	0.0388 (17)	0.0513 (19)	-0.0071 (12)	-0.0013 (13)	-0.0280 (15)
C4	0.0242 (15)	0.0279 (15)	0.0501 (19)	-0.0039 (12)	0.0005 (13)	-0.0136 (14)
C5	0.0269 (15)	0.0348 (16)	0.0392 (16)	-0.0046 (12)	-0.0039 (12)	-0.0133 (14)
C6	0.0291 (15)	0.0323 (16)	0.0387 (16)	-0.0080 (12)	-0.0021 (12)	-0.0163 (13)
O1	0.0424 (13)	0.0343 (12)	0.0402 (12)	-0.0132 (10)	-0.0033 (10)	-0.0072 (10)
C7	0.054 (2)	0.040 (2)	0.060 (2)	0.0007 (17)	-0.0095 (19)	-0.0062 (18)
O2	0.0456 (14)	0.0687 (17)	0.0418 (13)	-0.0189 (12)	-0.0064 (10)	-0.0275 (13)
C8	0.050 (2)	0.076 (3)	0.0380 (19)	-0.009 (2)	-0.0024 (16)	-0.0245 (19)
C11	0.0559 (6)	0.0536 (6)	0.0797 (7)	-0.0177 (4)	0.0007 (5)	-0.0454 (5)
C12	0.0526 (6)	0.0296 (11)	0.0692 (11)	-0.0075 (10)	-0.0009 (7)	-0.0165 (7)
Br3	0.0579 (9)	0.0403 (16)	0.0456 (6)	-0.0037 (12)	-0.0149 (5)	-0.0133 (8)
Br2	0.0526 (6)	0.0296 (11)	0.0692 (11)	-0.0075 (10)	-0.0009 (7)	-0.0165 (7)
C13	0.0579 (9)	0.0403 (16)	0.0456 (6)	-0.0037 (12)	-0.0149 (5)	-0.0133 (8)
C14	0.0670 (6)	0.0407 (5)	0.0471 (5)	-0.0185 (4)	-0.0078 (4)	-0.0221 (4)

Geometric parameters (Å, °)

C1—O1	1.351 (4)	C5—Cl3	1.758 (12)
C1—C6	1.383 (5)	C5—Br3	1.842 (6)
C1—C2	1.388 (5)	C6—Cl4	1.710 (3)
C2—O2	1.350 (4)	O1—C7	1.417 (5)
C2—C3	1.382 (5)	C7—H7A	0.9800
C3—C4	1.387 (5)	C7—H7B	0.9800
C3—C11	1.711 (3)	C7—H7C	0.9800
C4—C5	1.386 (5)	O2—C8	1.418 (5)
C4—C12	1.756 (15)	C8—H8A	0.9800
C4—Br2	1.837 (6)	C8—H8B	0.9800
C5—C6	1.379 (5)	C8—H8C	0.9800
O1—C1—C6	120.3 (3)	C4—C5—Br3	122.6 (3)
O1—C1—C2	120.0 (3)	C5—C6—C1	120.8 (3)
C6—C1—C2	119.7 (3)	C5—C6—Cl4	121.6 (3)
O2—C2—C3	119.5 (3)	C1—C6—Cl4	117.6 (2)
O2—C2—C1	120.9 (3)	C1—O1—C7	114.5 (3)
C3—C2—C1	119.5 (3)	O1—C7—H7A	109.5
C2—C3—C4	120.7 (3)	O1—C7—H7B	109.5
C2—C3—C11	118.5 (3)	H7A—C7—H7B	109.5
C4—C3—C11	120.8 (3)	O1—C7—H7C	109.5
C5—C4—C3	119.5 (3)	H7A—C7—H7C	109.5
C5—C4—C12	115.5 (6)	H7B—C7—H7C	109.5

C3—C4—Cl2	125.0 (7)	C2—O2—C8	115.4 (3)
C5—C4—Br2	121.8 (3)	O2—C8—H8A	109.5
C3—C4—Br2	118.7 (3)	O2—C8—H8B	109.5
C6—C5—C4	119.7 (3)	H8A—C8—H8B	109.5
C6—C5—Cl3	125.0 (5)	O2—C8—H8C	109.5
C4—C5—Cl3	115.3 (4)	H8A—C8—H8C	109.5
C6—C5—Br3	117.6 (3)	H8B—C8—H8C	109.5
O1—C1—C2—O2	4.0 (5)	Br2—C4—C5—Cl3	0.1 (7)
C6—C1—C2—O2	-177.9 (3)	C3—C4—C5—Br3	-177.6 (3)
O1—C1—C2—C3	-179.6 (3)	Cl2—C4—C5—Br3	1.8 (9)
C6—C1—C2—C3	-1.5 (5)	Br2—C4—C5—Br3	3.7 (5)
O2—C2—C3—C4	177.4 (3)	C4—C5—C6—C1	-0.6 (5)
C1—C2—C3—C4	1.0 (5)	Cl3—C5—C6—C1	-179.2 (7)
O2—C2—C3—Cl1	-2.9 (4)	Br3—C5—C6—C1	177.2 (3)
C1—C2—C3—Cl1	-179.3 (2)	C4—C5—C6—Cl4	178.8 (2)
C2—C3—C4—C5	-0.3 (5)	Cl3—C5—C6—Cl4	0.2 (8)
Cl1—C3—C4—C5	-179.9 (2)	Br3—C5—C6—Cl4	-3.4 (4)
C2—C3—C4—Cl2	-179.6 (9)	O1—C1—C6—C5	179.4 (3)
Cl1—C3—C4—Cl2	0.7 (10)	C2—C1—C6—C5	1.3 (5)
C2—C3—C4—Br2	178.5 (4)	O1—C1—C6—Cl4	0.0 (4)
Cl1—C3—C4—Br2	-1.2 (5)	C2—C1—C6—Cl4	-178.1 (2)
C3—C4—C5—C6	0.0 (5)	C6—C1—O1—C7	96.2 (4)
Cl2—C4—C5—C6	179.4 (8)	C2—C1—O1—C7	-85.7 (4)
Br2—C4—C5—C6	-178.7 (4)	C3—C2—O2—C8	113.2 (4)
C3—C4—C5—Cl3	178.8 (6)	C1—C2—O2—C8	-70.5 (4)
Cl2—C4—C5—Cl3	-1.8 (10)		
