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### Key indicators

Single-crystal X-ray study T = 173 K Mean  $\sigma$ (C–C) = 0.007 Å R factor = 0.031 wR factor = 0.073 Data-to-parameter ratio = 10.9

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

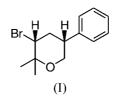
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# *cis*-(3*R*,5*R*)-3-Bromo-2,2-dimethyl-5-phenyl-tetrahydropyran

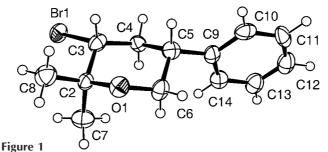
The heterocyclic ring in the title compound,  $C_{13}H_{17}BrO$ , adopts a  $_1C^4$  conformation, with the phenyl and bromo substituents located in equatorial positions.

## Comment

Positions 3 and 5 in *cis*-3-bromo-2,2-dimethyl-5-phenyltetrahydropyran, (I), are stereogenic centres. The racemate of (I) was prepared (Hartung *et al.*, 2003) and investigated by X-ray diffraction, in order to establish reference data for an upcoming conformational analysis of multiply substituted tetrahydropyrans.



Atoms O1 and C4 are displaced in opposite direction [-0.649 (6) Å for O1 and 0.690 (7) Å for C4] from the plane defined by atoms C2, C3, C5, and C6 [deviation of C6 = 0.01 (1) Å], thus leading to a  ${}_{1}C^{4}$  arrangement of the tetrahydropyran ring (Fig. 1). This arrangement corresponds to the major conformer of (I) in CDCl<sub>3</sub> solution, according to an analysis of the vicinal proton-proton coupling constants (<sup>1</sup>H NMR) (Hartung et al., 2003). The phenyl and bromo substituents in (I) are equatorially arranged [Br1-C3-C2-O1 = $-179.8 (3)^{\circ}$  and C9-C5-C4-C3 = 176.4 (4)°]. The absolute value of the six endocyclic torsion angles sum to  $349 (3)^{\circ}$  for the tetrahydropyran subunit in (I). The racemate of 3bromosubstituted tetrahydropyran (I) crystallizes as an enantiomorphic conglomerate in  $P2_12_12$  (orthorhombic). The absolute 3R, 5R configuration of (I) in the investigated crystal was established with the aid of the Flack (1983) parameter [-0.009 (19)]. No significant close contacts were found in the crystal structure of the brominated tetrahydropyran (I).



The molecular structure of (I). Displacement ellipsoids are plotted at the 50% probability level.

## Experimental

Crystals suitable for X-ray diffraction were obtained from a saturated solution of (I) in diethyl ether, which was kept in an atmosphere of petroleum ether.

## Crystal data

 $\begin{array}{l} C_{13} H_{17} BrO \\ M_r = 269.18 \\ Orthorhombic, \ P2_1 2_1 2 \\ a = 13.269 \ (2) \ \text{\AA} \\ b = 16.085 \ (2) \ \text{\AA} \\ c = 5.7278 \ (4) \ \text{\AA} \\ V = 1222.5 \ (3) \ \text{\AA}^3 \\ Z = 4 \\ D_x = 1.462 \ \mathrm{Mg \ m^{-3}} \end{array}$ 

### Data collection

Enraf-Nonius CAD-4 diffractometer  $\omega/2\theta$  scans Absorption correction:  $\psi$  scan (North *et al.*, 1968)  $T_{\min} = 0.402, T_{\max} = 0.513$ 2915 measured reflections 1598 independent reflections 1360 reflections with  $I > 2\sigma(I)$ 

### Refinement

Refinement on  $F^2$   $R[F^2 > 2\sigma(F^2)] = 0.031$   $wR(F^2) = 0.073$  S = 1.071598 reflections 146 parameters H atoms treated by a mixture of independent and constrained refinement Mo  $K\alpha$  radiation Cell parameters from 25 reflections  $\theta = 10-15^{\circ}$  $\mu = 3.34 \text{ mm}^{-1}$ T = 173 (2) K Block, colourless  $0.30 \times 0.20 \times 0.20 \text{ mm}$ 

 $\begin{aligned} R_{\rm int} &= 0.036\\ \theta_{\rm max} &= 22.5^{\circ}\\ h &= -14 \rightarrow 8\\ k &= 0 \rightarrow 17\\ l &= -6 \rightarrow 6\\ 3 \text{ standard reflections}\\ \text{every 100 reflections}\\ \text{intensity decay: none} \end{aligned}$ 

 $w = 1/[\sigma^{2}(F_{o}^{2}) + (0.0348P)^{2} + 0.2533P]$ where  $P = (F_{o}^{2} + 2F_{c}^{2})/3$  $(\Delta/\sigma)_{max} < 0.001$  $\Delta\rho_{max} = 0.39 \text{ e} \text{ Å}^{-3}$  $\Delta\rho_{min} = -0.27 \text{ e} \text{ Å}^{-3}$ Absolute structure: Flack (1983), 628 Friedel pairs Flack parameter = -0.009 (19) H atoms attached to C3 and C5 were located in a difference Fourier map and their positions were refined freely with isotropic displacement parameters. All other H atoms were placed in geometrically idealized positions (C-H = 0.95–0.99 Å), and refined as riding, with  $U_{iso}(H) = 1.2$  or 1.5 times  $U_{eq}(C)$ .

Data collection: CAD-4 Diffractometer Control Software (Enraf-Nonius, 1993); cell refinement: CAD-4 Diffractometer Control Software; data reduction: CAD-4 Diffractometer Control Software; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: PLATON (Spek, 2002) and ORTEP-3 (Farrugia, 1997); software used to prepare material for publication: SHELXL97.

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