

MOLECULAR STRUCTURE OF *r*-2-(*p*-TRIFLUOROMETHYLPHENYL)-TRANS-4,
TRANS-6-DIMETHYL-1,3-DIOXANE

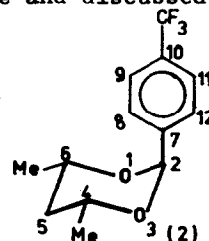
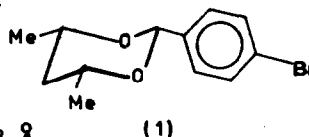
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In an earlier communication¹ the problem concerning conformational equilibria of 2-substituted 1,3-dioxanes have been discussed and the crystal and molecular structure of the equatorially aryl-substituted 1,3-dioxane (1), studied in this context, was reported. The X-ray analysis of an axially aryl-substituted analogue, the title compound (2)², is reported here and discussed in the context of the preceding paper.

Single crystals of (2), obtained by slow evaporation of petroleum ether solutions, were triclinic with space group $P\bar{1}$ and cell dimensions: $a=13.798 \text{ \AA}$, $b=4.758 \text{ \AA}$, $c=11.120 \text{ \AA}$, $\alpha=85.73^\circ$, $\beta=112.35^\circ$, $\gamma=101.66^\circ$. X-ray diffraction data were



collected on an automatic three-circle Siemens diffractometer controlled by a Siemens 305 on-line computer. The structure was solved by direct methods using a multiresolution technique³. The CF_3 group appeared to be statically disordered. Altogether nine distinct fluorine peaks appeared in successive difference maps from which, also, the locations of all hydrogen atoms were obtained. Refinement with anisotropic temperature factors for all heavy atoms, occupancy factors of .4, .4, and .3 for three sets of CF_3 groups, and isotropic temperature factors for hydrogen atoms with 2748 independent reflections yielded a final agreement factor of $R=.065$.

The 1,3-dioxane ring which has a non-crystallographic C_2 symmetry - as expected - appears in the chair conformation with both methyl groups occupying equatorial positions. The axial phenyl group is oriented in a perfect "perpendicular" conformation (see Fig. 1, nomenclature of ref. 4).

The crystal structure of (2) confirms the qualitative results in solution, namely that despite its relatively low ΔG° value, axial phenyl in the 2-position causes substantial distortions of the 1,3-dioxane ring geometry². Thus (2) appears to be initially sterically more strained than its 2-methyl analogue which nevertheless has the higher ΔG° value.

Deformations in the 1,3-dioxane ring⁵ which minimize the existing strain in (2) are apparent from ring torsional angles (cf. Fig. 1), the orientation

of the phenyl group and from changes in bond angles (cf. Fig. 2).

While in (1) the dioxane ring consists of a puckered ($O-C-O$: $\bar{\omega} = 61.4^\circ$) and a flattened ($C-C-C$: $\bar{\omega} = 53.3^\circ$) moiety, this distinction is lost in (2) which shows about equal torsional angles ranging from 53.7° ($\bar{\omega}_{C-C-C}$) to 54.4° (all ω_{C-O}). The major change is seen in the $O-C-O$ region; the torsional angles in the $C-C-C$ moiety remain virtually unaffected by the axial substitution.

The phenyl group is bent outward appreciably as was already suggested by dipole moment calculations², but the bending is not as extensive as was assumed earlier. If one takes the position of H-1 in (1) as the true and undistorted axial position at C-2 of the 1,3-dioxane chair then the bending-out amounts to about 7° . This bending-out is reflected in the widening of valency angles $O-C-2-C-7$ from 108° in (1) to 112.5° in (2). Such bending has previously been observed in r-2-(p-bromophenyl)-2,4,4,trans-6-tetramethyl-1,3-dioxane⁷. Over and above this effect the carbon atom C-2 is displaced significantly out of the plane formed by the six coplanar carbon atoms of the phenyl ring (maximal deviation .01 Å). This deviation corresponds to a 6.5° angle between the $C-2-C-7$ bond vector and the plane of the phenyl so that the total bending amounts to ca. $13-14^\circ$.

The experimental bond angles of (2) (cf. Fig. 2) compare well with those of (1) as far as the $C-C-C$ moiety and the methyl substituents are concerned. The only significant changes are observed for the $C-O-C$ angles which are widened from 111.1° in (1) to 113.8° which clearly is connected with the flattening in this part of the dioxane ring, as is the earlier discussed change in the $O-C-2-C-7$ angles.

The experimental bond lengths (cf. Fig. 2) compare well with those found for (1) with the exception of the $C-2-C-7$ bond. In the axial isomer (2) this bond is longer by .2-.3 Å (1.532 compared to 1.511 in (1) and 1.50 Å in p-chlorophenyl-1,3-dioxane⁵). The accepted average value for a $C_{sp^3}-C_{sp^2}$ bond reported by Bastiansen⁶ is 1.504 Å. The deviation for the axial isomer seems to be well beyond the limit of error and therefore is probably of significance⁺.

+ In r-2-(p-chlorophenyl)-trans-4-t-butyl-1,3-dioxane (yet unpublished results) the corresponding $C-C_{aryl}$ bond is 1.521 Å, again significantly longer than the value of Bastiansen⁶.

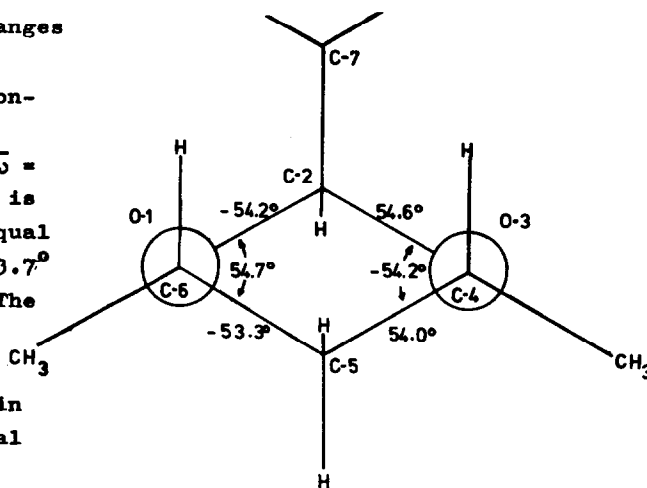


Fig. 1. Ring torsional angles of (2).

action. Experimentally this is clearly seen in the crystal structure of cis-p-bromophenyl-4-t-butyl-cyclohexane⁸. Since o-hydrogen/equatorial hydrogen interactions are absent in 2-phenyl-1,3-dioxane or, rather, supplanted by far less severe o-hydrogen/oxygen contacts (O-H...O 2.55 Å in (2)), it is reasonable that the resulting loss in interaction energy about balances the increase in syn-axial repulsion.

By the same token, the absence of the C-H/e-H interaction allows the phenyl ring to bend out substantially and with little resistance, thus further minimizing syn-axial interactions.

Besides the above steric argument we may have to consider electronic factors. The finding of a lengthening of the C-2 - C-7 bond in (2) may indicate the presence of an anomeric effect of the phenyl group which favours the axial position. Similar bond lengthening has been found for 2-halo-1,4-dioxanes by Altona⁹ and in 2-halo-glycopyranoses by Jeffrey and Pople¹⁰ and was related to the anomeric effect observed in these compounds¹¹.

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