CRYSTAL STRUCTURE OF A FUSED CYCLOHEXANE WITH AN AXIAL $t-B U T Y L$ GROUP: $8 \beta-t$-BUTYL-trans-DECAHYDROQUI NOLINE<br>Karl D. Hargrave and Ernest L. Eliel*<br>William R. Kenan, Jr. Laboratories of Chemistry University of North Carolina, Chapel Hill, N.C. 27514 USA

Summary: 8p-t-Butyl-trans-decahydroquinoline picrate crystallizes as a double chair with an axial t-butyl group. Except for a $16.2^{\circ}$ torsion of the t-butyl group away from the staggered conformation, the molecule displays only minor conformational deformations.

A t-butyl group substituted on a cyclohexane ring tends to avoid the axial position and thus serves as a "holding group" for the cyclohexane chair, preventing ring reversal. The first cyclohexane with a potentially axial t-butyl group - trans-1,3-di-t-butylcyclohexane was synthesized in $1960,^{2}$ but this species, ${ }^{3}$ as well as the corresponding cis-1,4-di-tbuty $^{3,4}$ and cis, trans-1,3,5-tri-t-butyl ${ }^{5}$ analogs appear to exist predominantly in the twist-boat form. A twist-boat was also originally assumed for the cis-1,2-di-t-butylcyclohexane, ${ }^{6}$ but later work ${ }^{7}$ suggests that this compound in fact exists as a chair with one axial t-butyl. The first reported axial t-butyl group is the one in cis-2-alkyl-5-t-butyl-1,3dioxane ( 2 alkyl group Me, Et, $i-\operatorname{Pr}$ or $t-B u$ ), ${ }^{8}$ but this case is anomalous in that the $t$-butyl group is syn-axial with two lone pairs of electrons rather than with two axial hydrogen atoms. traris-1,2-Di-t-butylcyclohexane and the corresponding $\mathrm{C}\left(\mathrm{CH}_{3}\right)_{2} \mathrm{CH}_{2} \mathrm{OH}$ and $\mathrm{C}\left(\mathrm{CH}_{3}\right)_{2} \mathrm{CO}$ Me substituted compounds in solution appear to be mixtures of diaxially substituted chairs and twistboats; ${ }^{7}$ the compound with two $\mathrm{CMe}_{2} \mathrm{CO}_{2} \mathrm{Me}$ groups exists as a diaxially substituted flattened chair in the crystalline state. ${ }^{9}$ A crystal structure determination has also been carried out of a 2 -t-butyl-l-alkylidenecyclohexane, ${ }^{10}$ in which the t-butyl group is axial because of $A^{(1,3)}$ stra1n.

To the best of our knowledge, no structure determination has yet been performed on a fully saturated cyclohexane with an axial t-butyl group. Recently we have synthesized ${ }^{11}$ such a structure in $8 \beta-t$-butyl-trans-decahydroquinoline (1). In relation to the argument concerning the position of the lone pair in this compound, ${ }^{11}$ it was important to be sure that 1 in fact exists in a chair form. This appeared plausible from the ${ }^{13} \mathrm{C}$ NMR data obtained; ${ }^{11}$ it has now been confirmed by $X$-ray structure analysis of the corresponding picrate (l. HPic). (Since vide infra - the salt exists as an only slightly distorted chair, the chair conformation may, a fortioni, be assumed for the free base.)


Crystal Data: $C_{19} \mathrm{H}_{28} \mathrm{~N}_{4} \mathrm{O}_{7}, M=424.45$, Monoclinic, $\underline{a}=9.241(4), \underline{b}=10.484(6), \underline{c}=22.584(9) \AA$,

 $\ell \neq 2 \underline{n}$, Ok0 when $\underline{k} \neq 2 \underline{n}$.
Crystallographic Measurements: A single crystal of dimensions ca. $0.16 \times 0.21 \times 0.88 \mathrm{~mm}$ was oriented on a Picker four-circle diffractometer equipped with graphite monochromater (Mo- $K_{\alpha}$ radiation; $1^{\circ}$ take-off angle). Refined unit-cell parameters were obtained by least-squares treatment of the $2 \theta, \chi$ and $\phi$ angles for 12 accurately centered high order reflections widely separated in reciprocal space. Intensities for 4692 unique reflections having $3^{\circ}<2 \theta<55^{\circ}$ were collected by the $\theta-2 \theta$ scanning technique. The data reduction procedure of Corfield et al. 12 was followed, yielding 2122 reflections used in the structure analysis.

Structure Analysis: The structure was solved by direct phase-determining methods with the MULTAN 76 series of programs ${ }^{13}$ and 350 reflections. The E-map based on that set of phases having the lowest $R$ and one of the highest absolute figures of merit revealed the positions of all the non-hydrogen atoms. Atomic positions 14 and non-hydrogen-atom anisotropic thermal parameters ${ }^{14}$ were refined by full-matrix least-square calculations to convergence at $R 0.057$ $\left(R=\Sigma| | F_{0}\left|-\left|F_{\mathrm{C}}\right|\right| / \Sigma\left|F_{0}\right|\right)$. Bond lengths $(\AA)$ and valency angles $\left({ }^{\circ}\right)$ and torsional angles ${ }^{\circ}$ ) for ${\underset{\sim}{l}}^{\circ} . \mathrm{HPic}$ are shown in Figs. 1 and 2, respectively; e.s.d.'s for the bond lengths are shown in parentheses, those for the valency and torsional angles are in the range $0.3-0.4^{\circ}$. Discussion: The A ring of $\underset{\sim}{1}$. HPic is clearly in the chair form with axial t-butyl; the overall distortion of this ring is surprisingly small. The mean endocyclic torsion angle in the $A$ ring, $54.6^{\circ}$ is intemediate between that (53.0 $)$ of the more flattened chair predicted by empirical force field calculation (SFF) for axial t-butylcyclohexane ${ }^{15}$ and the more normal value of $55.9^{\circ}$ for cyclohexane. 16 Flattening may be restrained by the adjacent trans-fused Bring.

In order to minimize severe non-bonded interactions between the t-butyl group and the axial hydrogens on $N(1), C(6)$ and $C(10)$, the $t$-butyl substituent deviates from a perfectly staggered arrangement relative to the ring (Fig. 3) exhibiting a twist of $16.2^{\circ}$. This value agrees perfectly with the corresponding twist of $16.3^{\circ}$ calculated (SFF) ${ }^{15}$ for the diaxial chair conformer of trans-1,2-di-t-butylcyclohexane and closely with the value of $17.4^{\circ}$ found in the crystal structure ${ }^{9}$ of the diaxial trans-1,2-bis[(2-methoxycarbonyl)-2-propyl]cyclohexane ( $\mathrm{CMe}_{2} \mathrm{CO}_{2} \mathrm{Me}$ analog), but is somewhat smaller than the $20.9^{\circ}$ calculated (SFF) ${ }^{15}$ for axial $t$-butyl. cyclohexane and a value of $24.2^{\circ}$ recently calculated (SFF) for $1-\alpha$ (axial)-t-butyl-transdecalin. 24



$$
\begin{array}{lr}
C(7)-C(8)-C(9)-N(1) & -175.9 \\
C(11)-C(8)-C(9)-C(10) & -78.4 \\
C(7)-C(8)-C(11)-C(12) & 157.4 \\
C(7)-C(8)-C(11)-C(14) & -81.0 \\
C(9)-C(8)-C(11)-C(13) & 170.4 \\
C(9)-C(8)-C(11)-C(14) & 47.4 \\
N(1)-C(9)-C(10)-C(5) & 176.3 \\
C(8)-C(9)-C(10)-C(4) & -174.5
\end{array}
$$

Figure 2

Figure 3

Further relief from steric crowding is accomplished by opening of the exocyclic valency angles at $C(8)$ to $121.6^{\circ}$ and $115.8^{\circ}$, with concomitant diminution of the endocyclic valency angle [at $C(8)]$ to $104.4^{\circ}$, and by enlargement of the $C(14)-C(q u a t e r n a r y)-C(r i n g)$ valency angle ${ }^{17}$ to $115.7^{\circ}$ with corresponding reduction of the $C(12)-C(q u a t e r n a r y)-C(13)$ valency angle to $104.9^{\circ}$. Such large valency angle deformations are normally not found in cases ${ }^{18-22}$ where the t-butyl group is equatorial. However, the $C(r i n g)-C$ (quaternary) bond length of 1.564 A , while longer than the normal $C\left(s p^{3}\right)-C\left(s p^{3}\right)$ distance ${ }^{23}$ of 1.537 A , is comparable to values predicted (SFF) ${ }^{15}$ and observed ${ }^{18-22}$ for rings bearing equatorial t-butyl groups. of note also is the large endocyclic valency angle at $C(9)\left(118.1^{\circ}\right)$; the mean carbon-carbon bond length ( 1.526 A ) and the mean endocyclic valency angle at carbon (111.2 ${ }^{\circ}$ ) are normal.

Interestingly, the deformation in $l_{2}$ appears to be substantially different from that in the previously studied ${ }^{10}$ 2-t-butylalkylidenecyclohexane in which strain is mainly relieyed by flattening of the ring and outward bending of the $t$-butyl group plus lengthening of the $C_{\text {ring }}{ }^{-}$ $\hat{C}_{\text {quat }}$ and shortening of the $C_{\text {quat }}-C_{\text {Me }}$ bonds. This may in part be a reflection of the much greater deformability of the alkylidenecyclohexane structure as compared to ${\underset{\sim}{l}}_{1}$ plus a low stretching force constant in the allylic bond.

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