Conformation Involving gauche t-Butyl Groups

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THE bulky t-butyl-group might be expected to impose severe conformational restraints in acyclic molecules¹, *cf.* cyclohexanes.² However, in n.m.r. studies of many acyclic pairs of diastereomers, very low vicinal coupling constants were found for t-butyl substituents quite unlike those for isopropyl or other large substituents (Table).³



* Ar s 2,4-dinitrophenyl. The spectra have been duplicated by computer simulation.

For the *erythro* isopropyl compound, quite high conformational purity seems likely, with vicinal protons predominately *trans* (the phenyl and isopropyl groups must likewise be *trans*).⁴ For the *erythro* t-butyl compound, the much lower *J*-value might indicate a substantial population of a conformer in which the t-butyl is *gauche* to another large group, in this case, phenyl. Possible angle deformations, involving the t-butyl group and the vicinal protons, could render the vicinal coupling constants a poor guide as to conformation.⁵ Alternatively, it was possible that the equilibrium dihedral angles in the various conformers were grossly different from 60° .⁶ In any case, other types of evidence are needed to corroborate the n.m.r. findings. In one system, 4,4-dimethyl-2,3-dibromopentane, dipole moment studies showed that both isomers possessed predominately gauche halogens; this is consistent with n.m.r. findings.^{7,8}



For di-t-butyl systems, Fahey⁹ reported a *J*-value of 0.8 c./sec. for (\pm) -3,4-dichloro-2,2,5,5-tetramethylhexane (very likely gauche protons and trans t-butyl groups). However, in the meso isomer, J_{AB} was 5.2 c./sec.,⁹ much lower than the value (>10) expected for trans protons (and hence trans t-butyl groups). Again it seemed possible that a second conformer, with gauche t-butyl groups, was

substantially populated (see, however, our previous reservations). If conformer B† were indeed populated, the dipole moment should be non-zero, due to vector addition of the C-Cl group moments.⁸ We found a dipole moment of 1.9 D (cyclohexane), § in reasonable agreement with the n.m.r. data.⁹ The dipole moment of the (\pm) -isomer was 2.7 D, more or less as expected, \ddagger and also in agreement with n.m.r. data.

The reason that seemingly unfavourable conformations such as B are populated is obscure. However, conformer A is itself far from strain free, as the extended conformational diagram shows,† Two 1,3 non-bonded interactions are present between chlorine and methyl, somewhat analogous to 1,3 diaxial interactions in a cyclohexane system. We suggest that some angle widening has occurred. Alternatively, some lateral movement of the methyl groups

Dihedral angles are shown as 60°, purely for convenience.

A. L. McClellan, "Tables of Experimental Dipole Moments," W. H. Freeman, San Francisco, 1965, p. 206. A check using fresh material and a slightly different technique gave a value of 2.20. The lower value is probably more reliable.

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may occur to reduce 1,3 interactions. In either case the gauche interaction between t-butyl groups would also be reduced, allowing an increased population of conformer B. However, these tentative ideas await more positive evidence, e.g. X-ray analysis.



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