THEORETICAL QUESTIONS ASSOCIATED WITH UTILIZING AB INITIO RESULTS TO AID IN INTERPRETA-TION OF VIBRATION-TORSION-ROTATION SPECTRA

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This talk will discuss several questions which arise when trying to connect ab initio results with molecular parameters in phenomenological torsion-rotation and vibration-torsion-rotation fitting Hamiltonians, and for which we now have varying degrees of understanding: (1) How are the various definitions of the torsional angle (intrinsic reaction coordinate, average of three dihedral angles, γ 's in torsional Hamiltonians in the literature, etc.) related? Short answer: they differ by a Fourier series in $\sin 3n\gamma$. (2) How can we determine the rho axis system in a flexible internal rotor molecule? Short answer: use an instantaneous rho system determined from the moment of inertia and the angular momentum generated at each point along the internal rotation path. (3) How exactly are partial derivatives (of atom positions, moment of inertia tensor, etc.) with respect to 3N - 6 internal coordinates to be related to partial derivatives with respect to one large amplitude coordinate and 3N - 7 small amplitude vibrational coordinates at various points along the internal rotation path? How exactly are the $(3N-6) \times (3N-6)$ F (force constant), G (reduced mass), and GF matrices to be converted to $(3N-7) \times (3N-7)$ quantities when the torsional degree of freedom is removed from the small-amplitude vibrational problem? Short answer: project out the torsional angle direction in the 3N-6 space and throw it away. (4) How are the C_{3v} permutation-inversion symmetry species A_1, A_2, E to be used in the vibrational problem after motion along the internal rotational path causes the C_s point group symmetry species A', A''to be lost? Short answer: work in progress. (5) How does setting certain terms to zero in the fitting Hamiltonian, which in turn implies certain reduction schemes, change the values of other phenomenological Hamiltonian parameters of lower, the same, or higher order, so that they are no longer directly comparable with ab initio output? Short answer: work in progress. Concrete illustrations of numerical answers to some of these questions for the methanol molecule will be given in the following talk.