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### ADVERTISEMENT



## Ab initio calculations and analysis of the torsional spectra of dimethylamine and dimethylphosphine

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In the present paper, *ab initio* calculations at MP2/RHF level are performed with different basis sets 6-31G(d,p), 6-311G(d,p), and 6-311(df,p) to determine the potential energy functions, the kinetic parameters, and the dipole moment components as a function of the double methyl rotation in dimethylamine (DMA) and dimethylphosphine (DMP). From the potential energy and kinetic parameters, the torsional energy levels and torsional functions are determined, and from the dipole moment variations, the far infrared spectra are synthesized by calculating both the frequencies and the intensities. The results are in relatively good agreement with experimental spectra. Calculations confirm the assignments performed with the experimental potentials fitted with only five terms. The calculations, however, allow to reassign the observed band at 239.8 cm<sup>-1</sup> in DMA and at 177.2 cm<sup>-1</sup> in DMP to the superimposition of two different transitions: the  $03 \rightarrow 04$  third sequence and an  $10 \rightarrow 11$  vibrationally excited fundamental. © *1996 American Institute of Physics*. [S0021-9606(96)03231-X]

#### I. INTRODUCTION

The analysis of the far infrared patterns of dimethylamine<sup>1</sup> (DMA) and dimethylphosphine<sup>2,3</sup> (DMP) shows an unusual complexity since the two torsional modes a' and a'' are active in infrared. Both sets of transitions lie in the same region of the spectrum giving rise to bc-hybrid and a-type bands. These bands show a prominent Q branch and their assignments require the comparative study of the Raman and FIR spectra. In DMA, furthermore, the bands due to the bending mode appear in the same region as those of two torsional overtones. In exchange, several torsional bands of vibrationally excited states of the CNC bending mode appear also in the same zone.

The torsional frequencies, structures and barriers of DMA<sup>1,4-15</sup> and DMP<sup>2-3,16-21</sup> have been considered in many papers. In 1967, Fateley *et al.*<sup>6</sup> and Möller *et al.*<sup>7</sup> have first observed the torsional spectrum of DMA. They have found barrier heights of 1150 and 1266 cm<sup>-1</sup>. Later, in 1971, Wollrab *et al.*<sup>9,10</sup> have obtained a barrier of 1126.2 cm<sup>-1</sup> from microwave data. The most relevant analysis of the IR and Raman structures of DMA is due to Durig, Griffin, and Groner,<sup>1</sup> who have evaluated, in 1977, a torsional barrier of 1053.8 cm<sup>-1</sup>. As is expected, the barrier height was found to possess approximately an intermediate value between those of methylamine<sup>22</sup> (714.6 cm<sup>-1</sup>) and trimethylamine<sup>23</sup> (1538.9 cm<sup>-1</sup>). Finally, Consalvo *et al.*<sup>15</sup> have also considered the Raman spectra of DMA.

The most relevant experiments for DMP are due to Durig, Griffin, and Natter,<sup>2</sup> and Durig, Groner, and Li.<sup>3</sup> Durig *et al.*<sup>2,3</sup> have evaluated a barrier height of 700.8 cm<sup>-1</sup> from IR and Raman spectroscopies and of 811.0 cm<sup>-1</sup> from MW data. This barrier shows also an intermediate value between those of methylphosphine<sup>24</sup> and trimethylphosphine.<sup>23</sup>

Fully and partially optimized *ab initio* calculations can be applied to verify the assignment of IR and Raman spectra. This technique was first employed for a set of molecules

showing the  $G_{36}$  symmetry and two  $C_{3v}$  symmetrical rotors thioacetone,<sup>25</sup> acetone,<sup>26</sup> biacetyl,<sup>27</sup> such as dimethyl-ether,<sup>28,29</sup> dimethyl-sulphide,<sup>30</sup> and butenes.<sup>31</sup> For that purpose, two and three dimension models were used. In the present paper, we applied the same technique to study the internal rotation in DMA and DMP, assuming that the torsional coordinates could be separated from those of the others vibration modes. The CNC (or CPC) bending and the hydrogen inversion modes, however, are expected to interact strongly with the two torsional modes in the amine or phosphine groups. The potential energy interactions are partially introduced in the geometry optimization. With this restriction, the two molecules can be classified according to the  $G_{18}$ nonrigid group.<sup>32-34</sup>

The minimal analytic expression<sup>32–34</sup> of the restricted torsional potentials of the  $G_{18}$  molecules is a ten term symmetry adapted Fourier expansion. Experimental data, however, allows only the fitting of five term functions into the  $G_{36}$  symmetry. The incidence of the remaining terms on the frequencies could be evaluated by performing *ab initio* calculations. In order to estimate these effects, the torsional levels were evaluated theoretically by using ten and seven termed expansions. For these purposes, different basis sets and approximations for the electronic correlation were resorted.

#### **II. THEORY**

By assuming the separability of large amplitude vibrations, the molecules of DMA and DMP may be described as a rigid  $C_s$  frame and two  $C_{3v}$  symmetric tops. Thus, the two-dimensional structure may be classified according to the  $G_{18}$  r-NRG group. At the lowest energy levels, where the tops undergo torsional oscillations, two vibrational normal modes, in which the torsion occurs in the same or opposite sense, may be defined. They can be classified into the a' and a'' representations of the  $C_s$  point group. Both modes are

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FIG. 1. The molecular structure of dimethylamine, the symmetry axes and the torsional angles  $\theta_1$  and  $\theta_2$ .

active in IR. Figure 1 shows the molecule of DMA, as well as the symmetry axis x, y, and z. The sense of the rotation of both methyl groups is defined clockwise.

The torsional Hamiltonian may be written as<sup>26</sup>

$$\hat{H}(\theta_1, \theta_2) = -\frac{\partial}{\partial \theta_1} B_{11} \frac{\partial}{\partial \theta_1} - \frac{\partial}{\partial \theta_1} B_{12} \frac{\partial}{\partial \theta_2}$$
$$-\frac{\partial}{\partial \theta_2} B_{12} \frac{\partial}{\partial \theta_1} - \frac{\partial}{\partial \theta_2} B_{22} \frac{\partial}{\partial \theta_2} + V(\theta_1, \theta_2)$$

where  $B_1(\theta_1, \theta_2)$ ,  $B_2(\theta_1, \theta_2)$ , and  $B_{12}(\theta_1, \theta_2)$  are the kinetic parameters and  $V(\theta_1, \theta_2)$  is the potential energy function. Both types of parameters depend on the two torsional angles. The analytic expression for them is a ten term double Fourier expansion.<sup>32–34</sup> The first seven terms are totally symmetric with respect to the exchange of the two torsional angles:

$$V(\theta_1, \theta_2)$$

$$= \sum_{L>K}^{2} \sum_{K=0}^{1} A_{KL}^{cc} [\cos 3K\theta_1 \cos 3L\theta_2 + \cos 3L\theta_1 \cos 3K\theta_2] + \sum_{K}^{2} (A_{KK}^{cc} \cos 3K\theta_1 \times \cos 3K\theta_2) + \sum_{K}^{1} A_{KK}^{ss} (\sin 3K\theta_1 \sin 3K\theta_2) + \sum_{K=0}^{2} A_{K}^{cs} [\cos 3K\theta_1 \sin 3\theta_2 - \sin 3\theta_1 \cos 3K\theta_2].$$

TABLE I. Character table of the G<sub>18</sub> r-NRG.<sup>a</sup>

The last three are antisymmetric. Both types of terms, however, are invariant with respect to the double-switchexchange operation, WV.

The r-NRG<sup>34</sup>  $G_{18}$  contains all the symmetry operations that commute with the torsional Hamiltonian. This group can be defined by the products

$$G_{18} = [C_3^I \otimes C_{3'}^I] \wedge [WV^I]$$

where the subgroups are defined as

$$C_{3}^{I} = [\hat{E} + \hat{C}_{3} + \hat{C}_{3}^{2}], \quad WV^{I} = [\hat{E} + \hat{W}\hat{V}]$$

and  $\hat{C}_3$  and  $\hat{W}\hat{V}$  are the rotation and double-switch-exchange operators, respectively. This last operation is defined by the expression:

$$\hat{W}\hat{V}f(\theta_1,\theta_2) = f(-\theta_2,-\theta_1).$$

Table I shows the character table of the  $G_{18}$  group.<sup>33</sup> This group contains six symmetry species. The nondegenerate  $A_1$  and  $A_2$  are symmetric and antisymmetric with respect to the double-switch-exchange operation. The pseudodegenerate representations  $E_1$  and  $E_2$  include a complex conjugate pair of one-degenerate representations ( $E_{1a}$  and  $E_{1b}$ ,  $E_{2a}$  and  $E_{2b}$ ).  $E_3$  is a two-degenerate representation and G contains a complex conjugate pair of two-degenerate representations ( $G_{1a}$ ,  $G_{1b}$ ,  $G_{2a}$ , and  $G_{2b}$ ).

The nuclear Hamiltonian is solved variationally by expanding the solutions onto the basis of the symmetry eigenvectors, which factorize the Hamiltonian matrix in eight boxes. Six correspond to the  $A_1, A_2, G_1, G_2, E_1$ , and  $E_2$ representations and two to the  $E_3$  two-degenerate specie. Boxes containing pseudo-degenerate representations cannot be factorized without including the wagging coordinate as a third degree of freedom. The symmetry eigenvectors for the  $G_{18}$  group may be determined from those of  $G_{36}$  group developed for the analysis of the torsional spectra of acetone.<sup>35</sup> Thus, the set of eigenvectors for the  $A_1$  representations contains the  $A_1$  and  $A_2$  ones of the  $G_{36}$  group. In the same way, the  $G_{18}$  representations  $A_2$  and  $E_3$  can be related to the  $A_3$ and  $A_4$ , and  $E_{3x}$  and  $E_{4y}$  representations of the  $G_{36}$  group. The vectors for the pseudo-degenerate representations  $G_1$ ,  $G_2, E_1$ , and  $E_2$  can be obtained from the  $G_1$  and  $G_2, G_3$  and  $G_4$ ,  $E_{1x}$  and  $E_{1y}$ , and  $E_{2x}$  and  $E_{2y}$  of the  $G_{36}$  group.

	Ê	$\begin{array}{c} \hat{C}_{3'}^2 \\ C_3 \end{array}$	$\hat{C}_3^2$ $\hat{C}_{3'}$	$2\hat{C}_3\hat{C}_{3'}$	$\hat{C}_3\hat{C}_{3'}^2$	$\hat{C}_3^2 \hat{C}_{3'}$	$\hat{W}\hat{V} \\ \hat{W}\hat{V}\hat{C}_{3}^{2}\hat{C}_{3}^{2}, \\ \hat{W}\hat{V}\hat{C}_{3}\hat{C}_{3}, \end{cases}$	$ \begin{array}{c} \hat{W}\hat{V}C_{3}C_{3}^{2}, \\ \hat{W}\hat{V}\hat{C}_{3}^{2}, \\ \hat{W}\hat{V}\hat{C}_{3}^{2}, \\ \hat{W}\hat{V}\hat{C}_{3} \end{array} $	$\hat{W}\hat{V}\hat{C}_{3}^{2}\hat{C}_{3'},\ \hat{W}\hat{V}\hat{C}_{3}^{2},\ \hat{W}\hat{V}\hat{C}_{3'},\ \hat{W}\hat{V}\hat{C}_{3'},$
$A_1$	1	1	1	1	1	1	1	1	1
$A_2$	1	1	1	1	1	1	-1	-1	-1
$E_{1a}$	1	$\omega^2$	ω	1	$\omega^2$	ω	1	$\omega^2$	ω
$E_{1b}$	1	ω	$\omega^2$	1	ω	$\omega^2$	1	ω	$\omega^2$
$E_{2a}$	1	$\omega^2$	ω	1	$\omega^2$	ω	-1	$-\omega^2$	$-\omega$
$E_{2b}$	1	ω	$\omega^2$	1	ω	$\omega^2$	-1	$-\omega$	$-\omega^2$
$E_3$	2	-1	-1	-1	2	2	0	0	0
$G_{1a}$	2	$-\omega^2$	$-\omega$	-1	$2\omega^2$	$2\omega$	0	0	0
$G_{1b}$	2	$-\omega$	$-\omega^2$	-1	2ω	$2\omega^2$	0	0	0

 $a\overline{\omega} = \exp(2\pi i/3).$ 

#### J. Chem. Phys., Vol. 105, No. 7, 15 August 1996

TABLE II. Selection rules for DMA and DMP.

FIR Raman	$\begin{array}{c} \mu_x \ \mu_y \\ \alpha_{xx}, \alpha_{yy}, \alpha_{zz} \end{array}$	$\mu_z$
	$A_1 \rightleftharpoons A_1$ $A_2 \rightleftharpoons A_2$	$A_1 \rightleftharpoons A_2$
	$G \rightleftharpoons G$	$G \rightleftharpoons G$
	$E_1 \rightleftharpoons E_1 \\ E_2 \rightleftharpoons E_2$	$E_1 \rightleftharpoons E_2$
	$E_3 \overleftrightarrow E_3$	$E_3 \rightleftharpoons E_3$

The intensities of each type of bands are determined into the electric dipole moment variation approximation. In this aim, the intensity equation adapted for a two symmetric top problem<sup>36</sup> is used:

$$f_{fi} = \frac{g}{3B} \left( E_f - E_i \right) \left( C_f - C_i \right) \left\langle \psi_i \right| \frac{\mu(\theta_1, \theta_2)}{\text{Re}} \left| \psi_j \right\rangle^2.$$

In this expression,  $E_f$ ,  $E_i$ ,  $C_f$ ,  $C_i$ ,  $\psi_f$ , and  $\psi_i$  are the energies, populations, and torsional wave functions of the final and initial states, respectively.  $\mu(\theta_1, \theta_2)$  is the dipole moment vector expressed as a function of the rotation angles. *g* is the nuclear statistical weight, *R* and *B* are the average rotation radius and kinetic parameter of a methyl group, and *e* the elemental electric charge.

The electric dipole moment components  $\mu_z$ ,  $\mu_y$ , and  $\mu_x$ with respect to the principle axis *a*, *b*, and *c*, have to be deduced from the *ab initio* calculations. It is easy to verify that these components transform according to the  $A_2$ ,  $A_1$ , and  $A_1$  representations of the  $G_{18}$  group. In Table II, the selection rules obtained from the nonzero conditions of the transitions moments, are given. Because of the vectorial nature of the spin functions, the nuclear statistical weights for each symmetry coincide with those determined for the  $G_{36}$  molecules.<sup>26</sup> Finally, the populations are given by a Boltzmann statistics.

#### **III. CALCULATIONS**

Fully optimized ab initio calculations were performed at the MP2/RHF level with the program GAUSSIAN 92.37 Torsional energies, frequencies and intensities were determined with the program ROCA25 especially written for this paper. The potential energy functions were calculated by fitting the relative electronic energy values (with respect to the minima) of ten selected conformations to Eq. (2). In Table III, these relative electronic energies calculated with 6-31G(d,p), 6-311G(d,p), and 6-311G(df,p) are given. The angles  $\theta_1 = 0.0 \ y \ \theta_2 = 0.0$  correspond to the planar conformation in which two of the hydrogens are lying on the CNC (CPC) plane and pointing outward. The relative energies for these conformations were determined to be 33.588 cm<sup>-1</sup> in DMA and 149.776 cm<sup>-1</sup> in DMP [MP2/6-31G(df, p)]. Table III shows also the effective barrier height (saddle-point) and the maxima of the surface, as well as the conformational angles of the minima. In Table IV, the expansion coefficients of the potential energy functions are given.

The minimum energy conformation of DMA shows torsion angles of  $\theta_1$ =3.3° and  $\theta_2$ =-3.3° [MP2/6-311G(*df*,*p*)]. The wagging and bending coordinates at this equilibrium geometry were found to be equal to 55° and 111.6°, respectively. The C–H and N–H bond distances were found to be 1.449 and 1.012 Å. The minimum energy conformation of DMP possesses torsion angles of  $\theta_1$ =10.8° and  $\theta_2$ =-10.8°. The bending and wagging angles of this structure were found to be 99.9° and 67°. The values for the C–P and P–H bond distances are equal to 1.84 and 1.41 Å. The separation be-

TABLE III. Relative energies (in cm<sup>-1</sup>) with respect to the equilibrium geometry at the MP2/RHF approximation.

			DMA <sup>a</sup>		DMP <sup>a</sup>		
$ heta_1^b$	$ heta_2^v$	Ι	П	III	Ι	П	III
0.0	0.0	39.967	40.320	33.588	150.362	151.356	149.776
60.0	0.0	1259.180	1222.850	1197.378	770.059	731.396	747.963
60.0	60.0	2385.832	2372.815	2334.771	1506.754	1480.753	1536.449
30.0	0.0	766.030	743.204	710.012	704.215	677.945	684.424
60.0	30.0	1661.300	1626.821	1614.142	865.429	829.613	860.862
30.0	30.0	1213.572	1174.133	1148.663	760.542	717.255	731.962
30.0	-30.0	1418.349	1414.472	1356.711	1349.330	1331.405	1356.750
0.0	30.0	454.536	425.325	423.462	203.830	180.321	183.978
30.0	60.0	1951.405	1936.607	1884.283	1413.513	1381.757	1417.870
-30.0	30.0	861.653	836.060	848.160	297.289	277.826	293.521
$V_{\rm eff}$		1263.578	1228.356	1201.238	762.117	726.434	744.954
Maximum		2420.453	2412.709	2364.790	1690.390	1672.736	1727.413
Equilibrium structure	e						
$\theta_1 = -\theta_2$	$P_2$	3.6	4.0	3.3	10.9	11.1	10.8
Total energy	,c	-134.737 67	-134.829 33	-134.875 40	-420.974 01	-420.172 56	-420.220 11

<sup>a</sup>Basis set I(6-31G(d,p); basis set II=6-311G(d,p); basis set III=6-311G(df,p).

<sup>b</sup>In degrees.

<sup>c</sup>In a.u.

TABLE IV. Calculated expansion coefficients for the potential energy in dimetylamine and dimethylphosphine<sup>a</sup>.

		$DMA^b$			$\mathrm{DMP}^{\mathrm{b}}$	
	<i>V</i> <sub>1</sub>	$V_2$	V <sub>3</sub>	$V_1$	$V_2$	$V_3$
$A_{00}^{cc}$	1207.112	1181.297	1157.792	796.492	767.277	786.656
$A_{10}^{cc}$	-609.461	-609.682	-597.945	-427.789	-424.612	-437.577
$A_{11}^{cc}$	-26.676	-11.001	-9.422	41.539	54.967	59.376
$A_{20}^{cc}$	15.612	18.061	17.091	4.298	7.436	9.124
$A_{21}^{cc}$	4.347	6.505	6.747	5.192	6.428	5.723
$A_{22}^{cc}$	-1.434	-0.064	-0.157	-1.432	-0.748	-0.573
$A_{11}^{ss}$	34.433	21.328	21.202	-20.057	-30.558	-33.680
$A_{01}^{cs}$	-30.237	-25.673	-30.894	-37.482	-35.024	-35.068
$A_{11}^{cs}$	-7.334	-5.032	-6.496	4.806	5.872	5.686
$A_{21}^{cs}$	-6.960	-8.127	-7.676	-1.113	-2.267	-1.196

 $^{a}$ In cm<sup>-1</sup>.

 ${}^{b}V_{1}$  from MP2/6-31G(d,p) energies;  $v_{2}$  from MP2/6-311G(d,p);  $v_{3}$  from MP2/6-311G(df,p) energies.

tween the methyl hydrogen atoms appears to be larger in DMP than in DMA. The variations of the coordinates with the torsion are weaker in DMP than in DMA, since the steric interactions are smaller.

The kinetic parameters were determined for each conformation from the optimized geometries. For this purpose, the derivatives of the Cartesian coordinates with respect to the torsional coordinates were calculated numerically. In Table V, the expansion coefficients of the kinetic functions, obtained by fitting the parameters for each structure to a symmetry adapted Fourier expansion, are given.

The torsional energy levels were determined by using six different potentials of  $G_{36}$  and  $G_{18}$  symmetries obtained by using three different basis set in the electronic calculations. The  $G_{36}$  potential and the  $G_{36}$  kinetic parameters were determined from the  $G_{18}$  functions of Tables IV and V by dropping the nonsymmetric terms with respect to the exchange operator. An accurate variational calculation of the levels requires 37×37 torsional basis functions. Thus, the Hamiltonian matrix factorizes in eight boxes of dimensions:  $A_1(91), A_2(78), G(2 \times 312), E_1(156), E_2(132), \text{ and}$   $E_3(2 \times 144)$ . Boxes G,  $E_1$ , and  $E_2$  contain two pseudodegenerate and inseparable representations. Table VI shows the levels for both molecules obtained with four of these potentials and classified according to the vibrational quanta and the symmetry species of the  $G_{36}$  and  $G_{18}$  groups. In Tables VII, the calculated and observed band positions are given.

For the intensities,<sup>36</sup> the values of the dipole moment components, obtained for each conformation at the RHF/6-311(df,p) level, were retained and fitted to  $A_2, A_1$ , and  $A_1$ symmetry adapted Fourier expansions. Intensities for the inseparable degenerate species were the sums of those of the separate allowed transitions. The G-G transitions were found to be the most intense because of the effect of the nuclear statistical weights. These intensities are given in Table VIII.

The rotational contours<sup>38</sup> of the bands were simulated from the expectation values of the A, B, and C rotational constants at the two first  $A_1$  levels as well as at the first  $A_2$ . Figures 2 and 3 show a plot of the calculated rotational structures corresponding to bc-hybrid and a-type bands. Both

TIDDD IT Dipulsion coefficients for the innere energy.	TABLE V.	Expansion	coefficients	for the	kinetic	energy.a
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	MP2/6-3	MP2/6-31G( <i>d</i> , <i>p</i> )		11G( <i>d</i> , <i>p</i> )	MP2/6-31	MP2/6-311G( <i>df</i> , <i>p</i> )	
	$B_1$	<i>B</i> <sub>12</sub>	$B_1$	<i>B</i> <sub>12</sub>	<i>B</i> <sub>1</sub>	<i>B</i> <sub>12</sub>	
DMA <sup>b</sup>							
$A_{00}^{cc}$	6.696	-1.111	6.626	-1.090	6.651	-1.110	
$A_{10}^{cc}$	-0.054	0.051	-0.051	0.048	-0.053	0.050	
$A_{11}^{cc}$	0.020	-0.024	0.016	-0.021	0.019	-0.023	
$A_{11}^{ss}$	-0.027	0.032	-0.025	0.030	-0.027	0.032	
$A_{01}^{cs}$	-0.020	0.021	-0.016	0.016	-0.019	0.019	
DMP <sup>c</sup>							
$A_{00}^{cc}$	5.856	-0.259	5.805	-0.255	5.812	-0.262	
$A_{10}^{cc}$	-0.011	0.009	-0.012	0.010	-0.013	0.010	
$A_{11}^{cc}$	0.006	-0.006	0.006	-0.006	0.006	-0.007	
$A_{11}^{ss}$	-0.007	0.008	-0.007	0.009	-0.083	0.008	
$A_{01}^{cs}$	0.003	-0.006	0.003	-0.005	0.031	-0.005	

 $^{a}$ In cm<sup>-1</sup>.

<sup>b</sup>Experimental values are  $B_1 = B_2 = 6.622 \text{ cm}^{-1}$  and  $B_{12} = -1.115 \text{ cm}^{-1}$ . <sup>c</sup>Experimental values are  $B_1 = B_2 = 5.728 \text{ cm}^{-1}$  and  $B_{12} = -0.255 \text{ cm}^{-1}$ .

TABLE VI. (a) Dimethylamine and (b) dimethylphosphine energy levels.<sup>a</sup>

		$V_3(G_{36})$		$V_1(G_{18})$	$V_2(G_{18})$	$V_3(G_{18})$
(a)						
$\begin{pmatrix} a \\ 0 \\ 0 \end{pmatrix}$	Α.	240 392	Α.	248 791	240,454	237.541
0 0	G	240.392	G	248.791	240.454	237.541
	Ē,	240.392	Ē	248.791	240.454	237.541
	$E_{2}$	240.392	$E_{2}$	248.791	240.454	237.541
1 0	$A_2$	461.562	$A_{2}$	481.603	463.867	458.767
	Ğ	461.561	Ĝ	481.602	463.867	458.766
	$E_{2}$	461.560	$E_{2}$	481.602	463.866	458.765
	$\tilde{E_3}$	461.560	$\tilde{E_3}$	481.602	463.866	458.765
0 1	$A_2$	491.478	$A_1$	509.064	494.859	490.330
	G	491.477	G	509.064	494.859	490.329
	$E_1$	491.476	$E_1$	509.064	494.858	490.328
	$E_4$	491.476	$E_3$	509.064	494.858	490.328
2 0	$A_1$	677.741	$A_1$	707.107	682.433	675.153
	G	677.759	G	707.123	682.448	675.170
	$E_1$	677.778	$E_1$	707.138	682.464	675.188
	$E_3$	677.778	$E_3$	707.138	682.464	675.188
1 1	$A_4$	701.276	$A_2$	729.945	707.152	700.143
	G	701.304	G	729.968	707.176	700.171
	$E_2$	701.332	$E_2$	729.991	707.200	700.198
0.0	$E_4$	701.332	$E_3$	729.991	707.200	700.198
0 2	$A_1$	739.400	$A_1$	763.525	745.222	738.909
	G	739.409	G	763.532	745.230	/38.918
	$E_1$	739.418		763.539	745.238	728.927
2.0		/39.418		703.339	743.238	130.921
30	G A3	886 907	A <sub>2</sub>	923.697	894.329	884.663
	E.	886 729	Е.	923.755	894.182	884.003
	$E_2$ $E_2$	886 729	L <sub>2</sub> F.	923.613	894.040	884 493
2 1	$A_{3}$	903 181	A.	940 348	911 629	902 170
2 1	G	902.854	G	940.092	911.357	901.849
	Ē,	902.523	Ē,	939.834	911.082	901.525
	$E_{4}$	902.523	$E_3$	939.834	911.082	901.525
1 2	$A_3$	937.734	$A_2$	972.710	946.199	937.202
	Ğ	937.571	Ğ	972.595	946.060	937.038
	$E_2$	937.407	$E_2$	972.479	945.918	936.872
	$E_3$	937.406	$E_3$	972.479	945.918	936.872
0 3	$A_2$	982.308	$A_1$	1011.478	990.273	982.045
	G	982.287	G	1011.470	990.256	982.024
	$E_1$	982.266	$E_1$	1011.463	990.239	982.003
	$E_4$	982.266	$E_3$	1011.463	990.239	982.003
(b)						
0 0	$A_1$	179.505	$A_1$	184.902	175.592	177.339
	G	179.505	G	184.902	175.592	177.340
	$E_1$	179.506	$E_1$	184.903	175.593	177.340
1.0	$E_3$	1/9.500	$E_3$	184.903	1/5.595	2/2 269
10	G A3	345.013	G A2	359.855	340.440	343.200
	Е.	344 993	Е.	359.837	340.418	343 249
	$E_2$ $E_2$	344 993	L <sub>2</sub> F.	359.837	340.418	343 249
0 1	A <sub>2</sub>	367.201	$A_1$	376.200	361.209	365.437
	G	367.190	G	376.191	361.196	365.427
	$\tilde{E_1}$	367.179	Ē	376.181	361.184	365.415
	$E_4$	367.179	$E_3$	376.181	361.184	365.415
2 0	$A_1$	510.287	$A_1$	530.442	504.459	508.916
	G	510.406	G	530.572	504.597	509.034
	$E_1$	510.530	$E_1$	530.712	504.743	509.157
	$E_3$	510.530	$E_3$	530.712	504.743	509.157
1 1	$A_4$	518.943	$A_2$	535.907	511.788	517.494
	G	519.146	G	536.107	512.018	517.695
	$E_2$	519.345	$E_2$	536.299	512.240	517.891
	$E_4$	519.345	$E_3$	536.299	512.240	517.891
0 2	$A_1$	552.637	$A_1$	564.212	544.443	551.170
	G	552.711	G	564.262	544.521	551.243
	$E_1$	552.785	$E_1$	564.313	544.599	551.316
	$E_3$	552.785	$E_3$	564.313	544.599	551.316

TABLE VI. (Continued.)

		$V_3(G_{36})$		$V_1(G_{18})$	$V_2(G_{18})$	$V_3(G_{18})$
2 1	$A_2$	669.439	$A_1$	689.951	660.753	668.235
	G	665.800	G	686.144	656.806	664.635
	$E_1$	665.576	$E_1$	685.854	656.402	664.396
	$E_4$	665.576	$E_3$	685.853	656.401	664.395
3 0	$A_3$	669.496	$A_2$	690.430	661.351	668.344
	G	669.470	G	690.210	661.086	668.295
	$E_2$	666.033	$E_2$	686.475	657.279	664.887
	$E_3$	666.034	$E_3$	686.475	657.280	664.888
1 2	$A_3$	697.148	$A_2$	718.157	688.600	696.055
	G	696.767	G	718.029	688.294	695.686
	$E_2$	696.385	$E_2$	717.901	687.989	695.318
	$E_3$	696.385	$E_3$	717.901	687.989	695.318
03	$A_2$	733.976	$A_1$	746.434	723.291	732.735
	G	733.784	G	746.378	723.114	732.547
	$E_1$	733.588	$E_1$	746.321	722.933	732.355
	$E_4$	733.588	$E_3$	746.321	722.933	732.355

 $^{\mathrm{a}}\mathrm{In}\ \mathrm{cm}^{-1}.$ 

contours show a prominent Q branch. The Q branch of the hybrid structure is due from the c-type component.

#### **IV. ASSIGNMENT AND DISCUSSION**

Assignments of the Q branches have to be accomplished from the calculations and from the comparative analysis of the Raman and FIR spectra.<sup>1-3</sup> From the rotational band profiles in Figs. 2 and 3, it is clear that the *bc*-hybrid and *a*-type bands have sharp Q branches. Transitions attached to the *a'* mode are active in Raman, whereas the *a''* transitions are nonactive. The pattern observed in the FIR spectrum of DMA between 235.0 and 256.3 cm<sup>-1</sup> (see Fig. 4) is also visible in the Raman spectrum. Thus, it has to be assigned to the *a'* mode. Furthermore, bands observed between 202 and 220 cm<sup>-1</sup> should be assigned to *a''* mode. In DMP spectrum (see Fig. 5), four of the six recorded bands can be assigned to the *a'* mode. Calculations support this assignment.

The fundamental transitions present the strongest intensities (see Table VIII). They correspond to the sharpest bands of each symmetry. In DMA, the most prominent bands of each pattern were observed at 256.3 and 219.4 cm<sup>-1</sup>. They were predicted theoretically at 252.788 and 221.225 cm<sup>-1</sup> into the MP2/6-311G(df,p) approximation. In DMP, the two bands were observed at 190.4 and 169.9 cm<sup>-1</sup> and they were predicted at 188.087 and 165.918 cm<sup>-1</sup> at the same level of approximation. The separation between the two fundamentals of DMA was observed and calculated to be 36.9 and 31.563 cm<sup>-1</sup>, respectively. In DMP, 20.5 and 22.169 cm<sup>-1</sup>.

Divergences between the experimental and calculated values for the fundamental frequencies arise from the potential energy parameters rather than the kinetic parameters, since they were calculated accurately and are in agreement with the experimental data.<sup>1–3</sup> In DMA, one of the calculated fundamental's is overestimated and the other is underestimated. In consequence, divergences are related to the sin  $\times$ sin gearing term rather than to the barrier. However, in DMP, the two fundamentals are slightly underestimated. The

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TABLE VII. The frequencies (in  $\mbox{cm}^{-1})$  for (a) dimethylamine and (b) dimethylphosphine. 

(a) Mode $r_{12}$ 0 0-0 1 A)-A <sub>2</sub> 251.086 A <sub>1</sub> -A <sub>1</sub> 260.273 254.405 252.788 G-G 251.085 G-G 260.273 254.404 252.787 E <sub>3</sub> -E <sub>4</sub> 251.084 E <sub>3</sub> -E <sub>3</sub> 260.273 254.404 252.787 O 1-0 2 A)-A <sub>1</sub> 247.922 A <sub>1</sub> -A <sub>1</sub> 254.461 250.363 248.579 G-G 247.932 G-G 254.468 250.371 248.589 E <sub>1</sub> -E <sub>1</sub> 247.942 E <sub>1</sub> -E <sub>1</sub> 254.475 250.380 248.599 O 2-0 3 A)-A <sub>2</sub> 247.942 E <sub>1</sub> -E <sub>1</sub> 254.475 250.380 248.599 O 2-0 3 A)-A <sub>2</sub> 242.908 A <sub>1</sub> -A <sub>1</sub> 247.953 245.051 243.136 G-G 242.878 G-G 247.938 245.026 243.106 E <sub>1</sub> -E <sub>1</sub> 242.848 E <sub>3</sub> -E <sub>3</sub> 247.924 245.001 243.076 O 3-0 4 A)-A <sub>1</sub> 235.977 A <sub>1</sub> -A <sub>1</sub> 240.257 238.099 236.021 G-G 235.943 G-G 240.293 238.078 235.985 E <sub>1</sub> -E <sub>1</sub> 235.910 E <sub>1</sub> -E <sub>1</sub> 240.259 238.058 235.951 E <sub>4</sub> -E <sub>3</sub> 235.906 E <sub>3</sub> -E <sub>3</sub> 240.257 238.054 235.985 E <sub>1</sub> -E <sub>1</sub> 235.910 E <sub>1</sub> -E <sub>1</sub> 248.342 243.285 241.376 G-G 239.743 G-G 248.362 243.309 241.405 E <sub>4</sub> -E <sub>3</sub> 239.772 E <sub>2</sub> -E <sub>2</sub> 248.389 243.334 241.433 H <sub>0</sub> -A <sub>1</sub> 239.771 A <sub>1</sub> -A <sub>2</sub> 232.812 223.413 221.226 G-G 239.743 G-G 248.362 243.309 241.405 E <sub>4</sub> -E <sub>3</sub> 239.772 E <sub>3</sub> -E <sub>3</sub> 248.389 243.334 241.433 H <sub>3</sub> -A <sub>4</sub> 229.772 E <sub>3</sub> -E <sub>3</sub> 232.811 223.413 221.226 G-G 221.169 G-G 232.811 223.413 221.225 E <sub>1</sub> -E <sub>2</sub> 221.168 E <sub>1</sub> -E <sub>2</sub> 232.811 223.412 221.224 H <sub>3</sub> -A <sub>4</sub> 120.16179 A <sub>2</sub> -A <sub>1</sub> 225.504 218.566 216.386 G-G 201.148 G-G 235.251 218.81 216.404 E <sub>3</sub> -E <sub>3</sub> 216.188 E <sub>3</sub> -E <sub>3</sub> 225.536 218.598 216.423 E <sub>1</sub> -E <sub>2</sub> 21.168 E <sub>3</sub> -E <sub>3</sub> 232.811 223.412 221.224 H <sub>3</sub> -A <sub>1</sub> 216.179 A <sub>1</sub> -A <sub>2</sub> 216.475 211.576 209.305 G/O Mode $\nu_{12}$ 00 -0 0 A <sub>1</sub> -A <sub>3</sub> 209.349 A <sub>1</sub> -A <sub>2</sub> 216.475 211.576 209.305 E <sub>3</sub> -E <sub>3</sub> 208.951 E <sub>1</sub> -E <sub>2</sub> 216.475 211.576 209.305 G/O Mode $\nu_{12}$ 00 -0 01 A <sub>1</sub> -A <sub>2</sub> 187.696 A <sub>1</sub> -A <sub>1</sub> 188.021 183.234 185.591 188.075 G-G 185.521 G-G 188.071 183.325 188.103 G-G 185.521 G-G 188.071 183.325 188.5901 A <sub>1</sub> -A <sub>2</sub> 185.606 E <sub>3</sub> -E <sub>3</sub> 188.102 185.591 188.075 E <sub>3</sub> -E <sub>4</sub> 180.503 E <sub>1</sub> -E <sub>1</sub> 191.990 185.591 188.075 E <sub>3</sub> -E <sub>4</sub> 180.503 E <sub>1</sub> -E <sub>1</sub> 191.990 185.591 188.075 E <sub>3</sub> -E <sub>4</sub> 180.503 E <sub>3</sub> -E <sub>3</sub> 182.008 178.334 181.039 C <sub>2</sub> -O 3 A <sub>1</sub> -A <sub>2</sub> 187.673 G-G 182.116 178.593 1183.045 F <sub>1</sub> -E <sub>1</sub> 185.506 E <sub>3</sub> -E <sub>3</sub>		$V'_{3}a(G_{36})$		$V_1(G_{18})$	$V_2(G_{18})$	$V_3(G_{18})$
	(a)					
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	Mode $\nu_{12}$ 0 0 $\rightarrow$ 0 1					
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	$A_1 - A_2$	251.086	$A_1 - A_1$	260.273	254.405	252.789
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	G-G	251.085	G-G	260.273	254.405	252.788
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	$E_{1} - E_{1}$	251.084	$E_{1} - E_{1}$	260.273	254.404	252.787
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$E_{3} - E_{4}$	251.084	$E_{3} - E_{3}$	260.273	254.404	252.787
$\begin{array}{llllllllllllllllllllllllllllllllllll$	$0 1 \rightarrow 0 2$		5 5			
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	$A_2 - A_1$	247.922	$A_1 - A_1$	254.461	250.363	248.579
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	G-G	247.932	G-G	254.468	250.371	248.589
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	$E_{1} - E_{1}$	247.942	$E_1 - E_1$	254.475	250.380	248.599
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$E_4 - E_3$	247.942	$E_{3} - E_{3}$	254.475	250.380	248.599
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$0 2 \rightarrow 0 3$					
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$A_1 - A_2$	242.908	$A_1 - A_1$	247.953	245.051	243.136
$ \begin{array}{llllllllllllllllllllllllllllllllllll$	G-G	242.878	G-G	247.938	245.026	243.106
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	$E_1 - E_1$	242.848	$E_{1} - E_{1}$	247.924	245.001	243.076
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$E_{3}-E_{4}$	242.848	$E_{3} - E_{3}$	247.924	245.001	243.076
$\begin{array}{llllllllllllllllllllllllllllllllllll$	$0 3 \rightarrow 0 4$					
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$A_2 - A_1$	235.977	$A_1 - A_1$	240.327	238.099	236.021
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	G-G	235.943	G-G	240.293	238.078	235.985
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$E_1 - E_1$	235.910	$E_{1} - E_{1}$	240.259	238.058	235.951
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$E_4 - E_3$	235.906	$E_{3}-E_{3}$	240.257	238.054	235.946
$\begin{array}{llllllllllllllllllllllllllllllllllll$	$1  0 \rightarrow 1  1$					
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$A_3 - A_4$	239.714	$A_2 - A_2$	248.342	243.285	241.376
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	G-G	239.743	G-G	248.366	243.309	241.405
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$E_2 - E_2$	239.772	$E_2 - E_2$	248.389	243.334	241.433
$\begin{array}{llllllllllllllllllllllllllllllllllll$	$E_3 - E_4$	239.772	$E_3 - E_3$	248.389	243.334	241.433
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	Mode $\nu_{24}$					
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$0 0 \rightarrow 1 0$	221 170	A A	222 012	222 412	221 226
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$A_1 - A_3$	221.170	$A_1 - A_2$	252.812	223.413	221.220
$\begin{array}{cccccccccccccccccccccccccccccccccccc$		221.109		232.011	223.413	221.223
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$E_1 - E_2$	221.108	$E_1 - E_2$	232.011	223.412	221.224
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$L_3 - L_3$	221.108	$L_3 - L_3$	232.811	223.412	221.224
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$1  0 \rightarrow 2  0$	216 179	4 - 4	225 504	218 566	216 386
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$G_{-}G$	216.198	$G_{-}G$	225.504	218.500	216.300
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$E_{2}-E_{1}$	216 218	$E_{\alpha} - E_{\alpha}$	225.521	218 598	216.404
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$E_2 = E_1$ $E_2 = E_2$	216 218	$E_2 = E_1$	225.536	218 598	216.423
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$2, 0 \rightarrow 3, 0$	210.210	23 23	223.330	210.570	210.125
	$A_1 - A_2$	209.349	$A_1 - A_2$	216.790	211.896	209.685
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	G - G	209.148	$\dot{G}-\dot{G}$	216.630	211.734	209.493
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	$E_1 - E_2$	208.951	$E_1 - E_2$	216.475	211.576	209.305
	$E_{3}^{1} - E_{3}^{2}$	208.951	$E_{3}^{1} - E_{3}^{2}$	216.475	211.576	209.305
	5 5		5 5			
$\begin{array}{llllllllllllllllllllllllllllllllllll$	(b)					
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	Mode $\nu_{12}$					
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	$0  0 \rightarrow 0  1$	107 101		101 000	105 115	100.000
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	$A_1 - A_2$	187.696	$A_1 - A_1$	191.298	185.617	188.098
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	G-G	187.685	G-G	191.289	185.604	188.087
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$E_1 - E_1$	187.673	$E_1 - E_1$	191.990	185.591	188.075
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	$E_3 - E_4$	187.673	$E_3 - E_3$	191.990	185.591	188.075
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$0 \rightarrow 0 2$	195 426		100.012	102 024	105 722
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	$A_2 - A_1$	185.430	$A_1 - A_1$	188.012	183.234	185./35
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$		185.521	6-6 E E	100.071	103.323	185.001
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	$E_1 - E_1$ E E	185.000	$E_1 - E_1$	100.132	103.415	185.001
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	$L_4 - L_3$ 0 2 - 0 3	100.000	$L_3 - L_3$	100.132	103.413	105.901
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	A = A	181 330	<b>A A</b>	182 222	178 849	181 565
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	$G_{-}G$	181.073	$G_{-}G$	182.222	178 503	181 30/
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	F = F	180.803	6-0 E E	182.110	178 22/	181.004
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$E_1 = E_1$ $F_2 = F_1$	180.803	$E_1 - E_1$ $E_2 - E_2$	182.008	178 334	181 039
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	$0 \xrightarrow{3} 0 \xrightarrow{4} 0 \xrightarrow{4} 0$	100.005	<b>L</b> 3-L3	102.000	170.004	101.037
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	$A_2 = A_1$	174 949	$A \cdot - A \cdot$	175 871	172 495	175 176
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	G-G	175.580	G-G	176.064	172.983	175 799
$E_4 - E_3$ 176.189 $E_3 - E_3$ 176.255 173.465 176.399	$E_1 - E_1$	176.199	$E_1 - E_2$	176.260	173.473	176.409
	$E_{4}^{1}-E_{3}^{1}$	176.189	$E_{3}^{1} - E_{3}^{1}$	176.255	173.465	176.399

ABLE VII.	(Continued.)	

	$V_3'a(G_{36})$		$V_1(G_{18})$	$V_2(G_{18})$	$V_3(G_{18})$
$1  0 \rightarrow 1  1$					
$A_3 - A_4$	173.930	$A_2 - A_2$	176.052	171.348	174.226
G-G	174.143	G-G	176.261	171.589	174.437
$E_2 - E_2$	174.352	$E_{2}-E_{2}$	176.462	171.822	174.642
$E_3 - E_4$	174.352	$E_3 - E_3$	176.462	171.822	174.642
Mode $\nu_{24}$					
$0  0 \longrightarrow 1  0$					
$A_1 - A_3$	165.508	$A_1 - A_2$	174.953	164.848	165.929
G-G	165.498	G-G	174.944	164.837	165.918
$E_1 - E_2$	165.487	$E_{1} - E_{2}$	174.934	164.825	165.909
$E_{3} - E_{3}$	165.487	$E_{3}-E_{3}$	174.934	164.825	165.902
$1  0 \rightarrow 2  0$					
$A_3 - A_1$	165.274	$A_2 - A_1$	170.587	164.019	165.648
G-G	165.403	G-G	170.726	164.168	165.776
$E_2 - E_1$	165.537	$E_{2} - E_{1}$	170.875	164.325	165.908
$E_{3}-E_{3}$	165.537	$E_{3}-E_{3}$	170.875	164.325	165.908

errors are due to the calculated barrier height which is underestimated even in the calculations with the largest basis set.

The torsional barriers of DMA and DMP were calculated to be 1201.2 and 745.0 cm<sup>-1</sup> with MP2/6-31G(df, p). Recently, Durig et al.<sup>1-3</sup> and Wolrab et al.<sup>9,10</sup> have calculated the barrier heights from MW and IR data. In both molecules, the barriers fitted from IR data were found to be approximately 100 cm<sup>-1</sup> lower than those obtained from MW spectroscopy.<sup>1</sup> In DMA,<sup>1,9,10</sup> this difference comes from the average of the inversion splitting performed with the MW data.<sup>1-3,10</sup> On the contrary, the error in DMP<sup>2,3</sup> derives from the geometry employed in the calculations of the kinetic energy parameters.<sup>1,2</sup> The improvements of the basis decrease the DMA barrier height  $(1201.2 \text{ cm}^{-1})$ , whereas they introduce only barrier fluctuations around 750  $cm^{-1}$  in DMP. Anyway, the ab initio DMP barrier values remain always between those obtained from MW and IR spectroscopies,<sup>2,3</sup> in particular, they are larger than the IR values given by Durig *et al.*<sup>2</sup> (700.8 and 733 cm<sup>-1</sup>).

Barriers of DMA and DMP may be compared with those of dimethylether<sup>29</sup> (DME) and dimethyl-sulfide<sup>30</sup> (DMS). The barrier shape and origin of both types of molecules may be correlated. The sin×sin gearing term in DMA shows a positive sign, as in DME. This term arises from the interactions between the bending and torsion modes. It is easy to verify, during the optimization procedure, that the bending angle of both molecules does open to evade the steric effects between the methyl hydrogens. In exchange, the N–C/P–C bond-length ratio is comparable to the O–C/S–C one. The nonbonding interactions may be thus expected to be small in DMP as in DMS. Smaller steric effects give rise to lower barriers and a negative or very small sin×sin gearing term for DMP when compared with DMA.

Figure 2 show the intensities derived from c-type bands of the bc-hybrid. Selection rules in Table II relate the c-type bands with the a' mode and the a-type with the a'' mode. It can be inferred than relative intensities are well reproduced by this model. The ratio between c and a fundamental inten-

TABLE VIII. Calculated frequencies<sup>a</sup> and intensities<sup>b</sup> [MP2/6-311G(df,p)].

Assign.	Freq.	Int.	Expt. <sup>c</sup>	Freq.	Int.	Expt. <sup>c</sup>
		DMA			DMP	
Mode $v_{12}$						
$0  0 \rightarrow 0  1$						
$A_1 - A_1$	252.789	3.690		188.098	0.404	
G-G	252.788	7.379	256.3 vs	188.087	0.807	190.4 vw
$E_{1} - E_{1}$	252.787	1.845		188.075	0.202	
$E_3 - E_3$	252.787	1.845		188.075	0.202	
$0 1 \rightarrow 0 2$						
$A_1 - A_1$	248.579	2.038		185.733	0.237	
G-G	248.589	4.077	250.8 s	185.816	0.476	188.6 vw
$E_1 - E_1$	248.599	1.019		185.901	0.120	
$E_{3} - E_{3}$	248.599	1.019		185.901	0.120	
$0 2 \rightarrow 0 3$						
$A_1 - A_1$	243.136	0.834		181.565	0.108	
$\dot{G-G}$	243.106	1.657	245.3 ms	181.304	0.213	183.0 vvw
$E_1 - E_1$	243.076	0.417		181.039	0.053	
$E_{3} - E_{3}$	243.076	0.417		181.039	0.053	
0 3→0 4						
$A_1 - A_1$	236.021	0.000		175.176	0.000	
G-G	235.985	0.000	239.8 s	175.799	0.000	177.2 vvw
$E_{1} - E_{1}$	235.951	0.000		176.409	0.000	
$E_{3}^{-} - E_{3}^{-}$	235.946	0.000		176.399	0.000	
$1 0 \rightarrow 1 1$						
$A_2 - A_2$	241.376	0.926		174.226	0.117	
$\tilde{G-G}$	241.405	1.854	239.8 s	174.437	0.234	177.2 vvw
$E_2 - E_2$	241.433	0.464		174.642	0.059	
$E_{3}^{2} - E_{3}^{2}$	241.433	0.464		174.642	0.059	
Mode $\nu_{24}$						
$0  0 \rightarrow 1  0$						
$A_1 - A_2$	221.226	0.101		165.929	0.006	
$\dot{G-G}$	221.225	0.203	219.4 m	165.918	0.012	169.9 vvw
$E_1 - E_2$	221.224	0.051		165.909	0.003	
$E_{3}^{1} - E_{3}^{2}$	221.224	0.051		165.902	0.003	
$1 0 \rightarrow 2 0$						
$A_2 - A_1$	216.386	0.023		165.929	0.003	
$\overline{G}-\overline{G}$	216.404	0.046	213.0 w	165.776	0.005	166.1 vvw
$E_2 - E_1$	216.423	0.012		165.908	0.002	
$E_{3} - E_{3}$	216.423	0.012		165.908	0.002	
5 5						

<sup>a</sup>In cm<sup>-1</sup>

 $^{b} \times 10^{-4}$ .

<sup>c</sup>vs=very strong; s=strong; ms=medium strong; w=weak; vw=very weak; vvw=very weak.

sities was found 36.44 for DMA and 34.94 for DMP. Intensities are stronger in DMA than in DMP.

The first and second sequences of the a' mode in DMA were calculated to be 248.589 and 243.106 cm<sup>-1</sup> with the largest basis set. The differences between calculated and experimental bands<sup>1</sup> are -2.2 cm<sup>-1</sup>. Corresponding values for the a' mode were determined to be 216.404 and 209.493 cm<sup>-1</sup>. Divergences are equal to +3.4 and +6.8 cm<sup>-1</sup>. In DMP, the calculated values were 185.816 and 181.304 cm<sup>-1</sup> for the a' mode and 165.918 and 165.776 cm<sup>-1</sup> for a''. It can be deduced that the main diagonal of the potential surface (where  $\theta_1 = \theta_2$ ) related to the a'' mode is well described in the calculations, whereas the secondary diagonal is not so well reproduced.

Durig *et al.*<sup>1</sup> have assigned the IR band of DMA at 239.8 cm<sup>-1</sup> to the third sequence of the a' mode. This assignment have to be supported by the intensities. The observed intensity of this band, however, is unusually stronger than the second sequence one, whereas the calculated intensity is relatively weak. If one remarks that the calculated a' third sequence and the 10 $\rightarrow$ 11 (a'' torsionally excited a' funda-



FIG. 2. The rotational contours for the single degenerate A component simulated for *a*-type  $A_1(00) \rightarrow A_2(10)$  and *bc*-hybrid  $A_1(00) \rightarrow A_1(01)$  bands of dimethylamine.



FIG. 3. The rotational contours for the single degenerate A component simulated for a-type  $A_1(00) \rightarrow A_2(10)$  and bc-hybrid  $A_1(00) \rightarrow A_1(01)$  bands of dimethylphosphine.

mental) transitions are found approximately at the same frequency (taking into account that the theoretical values for the a'' levels are found 2 cm<sup>-1</sup> too high), the 239.8 cm<sup>-1</sup> band could be due thus the superimposition of both transitions. The observed band at 235.0 cm<sup>-1</sup>, assigned to the 10 $\rightarrow$ 11 transition by Durig *et al.* could be reassigned to the 21 $\rightarrow$ 11 overtone. The Raman band observed at 476 cm<sup>-1</sup> attached with the 01 $\rightarrow$ 12 overtone (approx 239.8+235.0) does agree with this reassignment.

In DMP<sup>2</sup>, the band observed at 177.2 cm<sup>-1</sup> contains in the same way the a' third sequence and the 10 $\rightarrow$ 11 transi-



FIG. 4. Observed and calculated far infrared spectra of dimethylamine. Observed spectrum is from Ref. 1.



FIG. 5. Observed and calculated far infrared spectra of dimethylphosphine. Observed spectrum is from Ref. 2.

tions. The experimental relative intensities support also this assignment.

In the present paper, the *ab initio* calculations appear to be an useful tool to evaluate the effects of the extra potential energy (and kinetic) terms on the band locations. These terms change drastically the symmetry properties of the Hamiltonian operator. The energy levels of Tables IV and V are calculated from one  $G_{36}$  and three  $G_{18}$  different potential energy functions. It is shown that the effects on the energy levels are relatively small. Calculations confirm generally the assignments performed with the experimental potentials fitted with only five terms. The calculations, however, allow to reassign the observed band at 239.8 cm<sup>-1</sup> in DMA and 177.2 cm<sup>-1</sup> in DMP to the superimposition of two different transitions: the third sequence and a vibrationally excited fundamental.

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