

## Infrared Matrix Isolation and Theoretical Studies on Glutarimide

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Infrared spectra of glutarimide isolated in low-temperature Ar and N<sub>2</sub> matrixes are reported. The molecular structure, vibrational frequencies, and infrared intensities of glutarimide are calculated with *ab initio* Hartree–Fock and second-order Møller–Plesset perturbation (MP2) methods as well as with density functional theory (DFT) using the nonlocal gradient corrected functional (BP86). The best overall agreement between the calculated and experimental spectra has been obtained at the MP2/D95V\*\* level. Unequivocal assignment of the experimental infrared bands is performed on the basis of the potential energy distribution (PED). A striking similarity is noted for frequencies of the corresponding CO and NH vibrations in glutarimide and in uracil, thymine, and their methyl derivatives. Furthermore, the significant flattening of the glutarimide ring, predicted by calculations, indicates its structural resemblance to pyrimidine bases. It is suggested that some glutarimide drugs are able to intercalate between nucleic base pairs in the DNA helix or they may act as antagonists of uracil and thymine in biological processes.

### Introduction

Glutarimide, GI (2,6-piperidinedione), is a structural part of a number of molecules with interesting biochemical activity. Several glutarimide derivatives have been reported to possess significant anticancer activity. For example, PCNU [ $\lambda$ -(chloroethyl)-3-(2,6-piperidinedione)-1-nitrosourea] can easily cross the blood–brain barrier, and it shows activity against brain tumors;<sup>1</sup> aminoglutethimide is a strong inhibitor of steroid biogenesis and is used in the treatment of metastatic breast cancer;<sup>2</sup> and antineoplaston A10 (*N*-phenylacetyl- $\alpha$ -aminoglutarimide), which has recently been introduced into experimental chemotherapy, has a remarkable anticancer activity, particularly against brain tumors, and it shows very low toxicity.<sup>3,4</sup> *In vitro* studies employing the thermal denaturation of DNA as well as fluorescence spectroscopy have revealed that antineoplaston A10 may bind weakly and noncovalently with DNA.<sup>5</sup> Eriguchi et al.<sup>6</sup> suggested that the A10 molecule may enter into DNA since it showed significant chemopreventive effects against various carcinogens in mice.

Glutarimide is also a component of newly synthesized antibiotics which exert antiviral and antifungal activity.<sup>7–9</sup> Cycloheximide, the best known member of glutarimide antibiotics, is a very strong inhibitor of protein synthesis.<sup>10</sup> It has been reported that the biological activity of cycloheximide is due to a specific binding interaction of the glutarimide moiety with a site on the 60 S ribosomal subunit.<sup>11,12</sup> Of particular interest is the fact that replacement of hydrogen in the CO–NH–CO (imide) group with the methyl or acetyl group results in complete loss of activity of glutarimide drugs.<sup>13</sup> This indicates that the biological activity of these drugs is determined by the specific hydrogen bonding between glutarimide and other molecules in biological systems.

Despite the wealth of published papers on pharmacological and biochemical studies on glutarimide drugs, there is very little

documented work concerning the structure and vibrational spectra of glutarimide. Recently, Xu and Clark<sup>14</sup> have performed experimental and theoretical studies on the electronic spectra of single crystals of glutarimide. The infrared spectra of glutarimide in solutions have been reported by Thompson et al.;<sup>15</sup> however, the proposed band assignment is very ambiguous.

Knowledge of the molecular structure and vibrational spectra of glutarimide is very important for the investigation of intermolecular interactions using vibrational spectroscopic methods. It is expected that these normal modes of glutarimide which involve vibrations of the imide group should reveal substantial changes in frequency due to the formation of hydrogen bonding with other molecules. Therefore, a clear understanding of the vibrational spectrum of a bare glutarimide is indispensable for a correct assignment of the spectra of hydrogen-bonded systems.

In this work we report, for the first time, the infrared spectra of monomeric glutarimide, measured in low-temperature argon and nitrogen matrixes. Extensive calculations of the structure and infrared spectra of glutarimide have been performed at three theory levels: *ab initio* Hartree–Fock, MP2, and density functional (DFT).

To get detailed information on the nature of normal modes, the potential energy distributions (PEDs) were calculated at each theory level. These combined spectroscopic and theoretical investigations allowed us to obtain unequivocal and complete vibrational assignment of the infrared spectrum of glutarimide. It is hoped that these results will be helpful for further studies on interaction of glutarimide drugs with biological molecules and may shed some light on the possible mechanism of the biological activity of glutarimide.

### Experimental Section

Glutarimide (Aldrich) was purified by vacuum sublimation prior to the matrix experiment. The matrix gases, argon (Linde AG) and nitrogen (Technische Gase, Leipzig), were of purity grade 6.0. The method of matrix preparation and the apparatus

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were the same as described elsewhere.<sup>16</sup> The sample was placed in the microoven in the vacuum chamber of a continuous-flow liquid helium cryostat. The vapors of the sample coming out from the heated microoven were mixed with the matrix gas, Ar or N<sub>2</sub> (precooled in a liquid nitrogen trap), and deposited on a CsI window on the coldfinger (10 K) of the cryostat. Glutarimide diluted in the matrix gas was deposited until the absorbance of the most intense carbonyl band was close to 1. The matrix gas/dopant ratio was experimentally adjusted to ensure that only the bands due to the monomer were recorded. The concentration of glutarimide in the matrix was sufficiently low that no bands due to associated species (known from the studies with more concentrated matrixes) were observed. The temperature of the cold window was maintained at 10 K. Such a low temperature prevented any diffusion and association during the matrix formation. No sign of thermal decomposition was observed during the experiment. The infrared spectra were recorded on a Perkin-Elmer Model 580B grating spectrophotometer operating at a resolution 1–3 cm<sup>-1</sup>. Integral intensities of the absorption bands were measured by numerical integration.

### Theoretical Methods and Computational Details

All calculations have been carried out using the Gaussian 94 code<sup>17</sup> running on a Cray J916 (XMP) supercomputer. The geometry optimizations and calculations of the vibrational spectra of glutarimide have been performed with *ab initio* Hartree–Fock (HF) and second-order Møller–Plesset perturbation (MP2) methods (using the frozen core approximation)<sup>18</sup> as well as with density functional theory (DFT).<sup>19</sup> The basis set contained the Huzinaga's (9s5p) primitive set,<sup>20</sup> contracted to the valence double- $\zeta$  collection, according to the scheme proposed by Dunning and Hay.<sup>21</sup> The basis set was augmented with a set of six-component d Gaussian polarization functions for heavy atoms and p polarization functions for hydrogen atoms, proposed by Hariharan and Pople.<sup>22</sup> This basis set, denoted as D95V\*\*, has been successfully used in previous calculations of vibrational spectra for molecules of a similar size.<sup>23,24</sup> The above choice of basis set leads to 155 molecular orbitals for glutarimide.

The DFT studies of glutarimide were performed employing three energy functionals: BP86 (Becke's exchange with Perdew correlation functionals),<sup>25,26</sup> BLYP (Becke–Lee–Yang–Parr),<sup>25,27</sup> and B3LYP (the combination of the Becke's three-parameter hybrid exchange functional<sup>28,29</sup> with the LYP correlation functional). The results obtained applying these functionals are comparable. The B3LYP functional tends to overestimate all frequencies. The BP86 results turned out to be the most accurate and quite similar to those obtained with the MP2 calculations, especially for C=O stretching modes and in the low-frequency region. Thus, the BP86 results have been listed in Table 2.

For each level of theory the geometry optimization of glutarimide was performed assuming C<sub>s</sub> symmetry (with the symmetry plane passing through N<sub>3</sub>, H<sub>15</sub>, C<sub>6</sub>, H<sub>13</sub>, and H<sub>14</sub> atoms). The atom numbering is shown in Scheme 1, and the optimized geometry parameters are compared in Table 1. The harmonic vibrational frequencies and eigenvectors as well as the infrared intensities were subsequently calculated using the analytical second derivatives for *ab initio* methods and numerical differentiation of analytical gradients in the DFT method. To express the normal modes in a molecule-fixed coordinate system, the nonredundant set of internal coordinates was defined as recommended by Fogarasi and Pulay.<sup>30</sup> These coordinates are listed in Table 2. Then the force constant matrices obtained in Cartesian coordinates at each theory level were transformed to internal coordinates, which allowed us to perform normal

**TABLE 1: Geometry (Bond Lengths in Angstroms, Angles in Degrees) and the Dipole Moment of Glutarimide Calculated within Hartree–Fock, DFT (BP86 Functional), and MP2 Approaches. Experimental Data Are from the X-ray Crystal Structure of Glutarimide (Ref 38)**

parameter <sup>a</sup>	HF	BP86	MP2	exptl <sup>b</sup>
C <sub>1</sub> –C <sub>2</sub>	1.513	1.526	1.516	1.508, 1.502
C <sub>1</sub> –C <sub>6</sub>	1.527	1.540	1.530	1.510, 1.486
C <sub>2</sub> –N <sub>3</sub>	1.384	1.408	1.400	1.391, 1.387
C <sub>2</sub> –O <sub>7</sub>	1.195	1.231	1.228	1.224, 1.215
N <sub>3</sub> –H <sub>15</sub>	1.002	1.029	1.017	0.88
C <sub>1</sub> –H <sub>10</sub>	1.082	1.102	1.092	0.98
C <sub>1</sub> –H <sub>12</sub>	1.088	1.108	1.097	1.103
C <sub>6</sub> –H <sub>13</sub>	1.087	1.105	1.095	1.11
C <sub>6</sub> –H <sub>14</sub>	1.084	1.104	1.094	0.95
C <sub>1</sub> –C <sub>2</sub> –N <sub>3</sub>	116.2	115.3	115.3	116.9, 116.1
C <sub>2</sub> –N <sub>3</sub> –C <sub>4</sub>	128.2	129.0	128.7	126.9
C <sub>6</sub> –C <sub>1</sub> –C <sub>2</sub>	112.5	113.0	112.1	113.7, 114.0
C <sub>1</sub> –C <sub>6</sub> –C <sub>5</sub>	110.2	110.6	109.9	111.8
O <sub>7</sub> –C <sub>2</sub> –C <sub>1</sub>	123.4	124.3	124.3	123.8, 124.0
H <sub>15</sub> –N <sub>3</sub> –C <sub>2</sub>	115.9	115.5	115.6	114.1
C <sub>1</sub> –C <sub>2</sub> –C <sub>4</sub> –C <sub>5</sub>	0.0	0.0	0.0	
O <sub>7</sub> –C <sub>2</sub> –C <sub>1</sub> –C <sub>5</sub>	179.0	178.7	178.8	
N <sub>3</sub> –C <sub>2</sub> –C <sub>1</sub> –C <sub>5</sub>	0.2	0.5	0.7	
H <sub>15</sub> –N <sub>3</sub> –C <sub>2</sub> –C <sub>1</sub>	178.9	179.2	179.0	
DH <sup>c</sup>	132.6	133.0	130.3	130.2
$\mu^d$	3.95	3.37	4.21	

<sup>a</sup> The atom numbering is shown in Scheme 1. The calculated structure has C<sub>s</sub> symmetry with the mirror plane passing through N<sub>3</sub>, H<sub>15</sub>, C<sub>6</sub>, H<sub>13</sub>, and H<sub>14</sub> atoms. <sup>b</sup> Due to H-bonding in the crystalline glutarimide, the structure of a molecule is significantly distorted and the corresponding bond lengths and angles on the left and right side of the ring have different experimental values. <sup>c</sup> DH is the dihedral angle between the plane defined by C<sub>1</sub>C<sub>6</sub>C<sub>5</sub> atoms and the plane defined by C<sub>1</sub>C<sub>2</sub>C<sub>4</sub>C<sub>5</sub> atoms. <sup>d</sup> Dipole moment in debyes.

coordinate analysis, as described in ref 31. The potential energy distributions (PEDs)<sup>32</sup> were calculated using our own program, which follows the formulas for PED matrix elements, given in ref 33. Perhaps the most important information obtained from PED calculations is the extent of mixing of various vibrations (expressed in percent contribution of internal coordinates to the normal mode). This allowed us to make a detailed description of the nature of the observed infrared bands.

The calculated frequencies of all normal modes within the HF method were scaled down by 0.90. For MP2 calculations we have used the factor of 0.96 for all normal modes, except for the modes Q<sub>1</sub>–Q<sub>7</sub>, which were scaled by 0.935 for the following reason: it is known that frequencies of the N–H, C–H, and O–H stretching vibrations calculated with both HF and MP2 methods are considerably overestimated when compared with the experiment.<sup>24,34–36</sup> This indicates that the harmonic approximation in *ab initio* calculations is unable to reproduce experimental stretching frequencies for these highly anharmonic vibrations. This problem cannot be resolved by a uniform scaling of all calculated vibrational frequencies since these modes are more anharmonic than the others. Therefore, we have introduced a new scaling factor of 0.935 for the MP2 calculated frequencies of N–H and C–H stretching modes to account for their large anharmonicity. The scaling factor was obtained by a least-squares fit to the infrared and Raman frequencies of N–H and C–H stretches observed for glutarimide and related molecules.<sup>24,34–36</sup>

### Results and Discussion

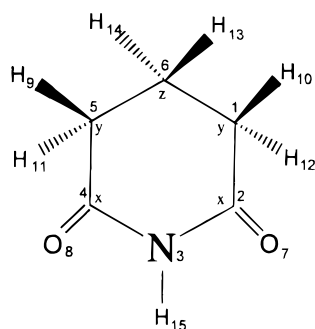
**A. Geometry.** The optimized geometrical parameters and the experimental data from the single-crystal X-ray analysis of glutarimide<sup>38</sup> are listed in Table 1. The calculated data correspond to a molecule of C<sub>s</sub> symmetry in the gas phase.

**TABLE 2: Internal Coordinates Used in the Normal-Mode Analysis of Glutarimide (Atom Numbering as in Scheme 1)**

definition <sup>a</sup>	Description	Symbol
$S_1=r_{1,2}+r_{5,4}$	$\nu(\text{C}_2\text{C}_1) + \nu(\text{C}_4\text{C}_5)$	$\nu^s(\text{C}_x\text{C}_y)$
$S_2=r_{1,2}-r_{5,4}$	$\nu(\text{C}_2\text{C}_1) - \nu(\text{C}_4\text{C}_5)$	$\nu^a(\text{C}_x\text{C}_y)$
$S_3=r_{2,3}+r_{4,3}$	$\nu(\text{C}_2\text{N}) + \nu(\text{C}_4\text{N})$	$\nu^s(\text{C}_x\text{N})$
$S_4=r_{2,3}-r_{4,3}$	$\nu(\text{C}_2\text{N}) - \nu(\text{C}_4\text{N})$	$\nu^a(\text{C}_x\text{N})$
$S_5=r_{1,6}+r_{5,6}$	$\nu(\text{C}_1\text{C}_6) + \nu(\text{C}_5\text{C}_6)$	$\nu^s(\text{C}_y\text{C}_z)$
$S_6=r_{1,6}-r_{5,6}$	$\nu(\text{C}_1\text{C}_6) - \nu(\text{C}_5\text{C}_6)$	$\nu^a(\text{C}_y\text{C}_z)$
$S_7=r_{2,7}+r_{4,8}$	$\nu(\text{C}_2\text{O}_7) + \nu(\text{C}_4\text{O}_8)$	$\nu^s(\text{C}_x\text{O})$
$S_8=r_{2,7}-r_{4,8}$	$\nu(\text{C}_2\text{O}_7) - \nu(\text{C}_4\text{O}_8)$	$\nu^a(\text{C}_x\text{O})$
$S_9=r_{3,15}$	$\nu(\text{N}_3\text{H}_{15})$	$\nu(\text{NH})$
$S_{10}=r_{1,10}+r_{5,9}$	$\nu(\text{C}_1\text{H}_{10}) + \nu(\text{C}_5\text{H}_9)$	$\nu^s \text{CH}_2; (\text{C}_y)$
$S_{11}=r_{1,10}-r_{5,9}$	$\nu(\text{C}_1\text{H}_{10}) - \nu(\text{C}_5\text{H}_9)$	$\nu^a \text{CH}_2; (\text{C}_y)$
$S_{12}=r_{1,12}+r_{5,11}$	$\nu(\text{C}_1\text{H}_{12}) + \nu(\text{C}_5\text{H}_{11})$	$\nu^s \text{CH}_2; (\text{C}_y)$
$S_{13}=r_{1,12}-r_{5,11}$	$\nu(\text{C}_1\text{H}_{12}) - \nu(\text{C}_5\text{H}_{11})$	$\nu^a \text{CH}_2; (\text{C}_y)$
$S_{14}=r_{6,13}+r_{6,14}$	$\nu(\text{C}_6\text{H}_{13}) + \nu(\text{C}_6\text{H}_{14})$	$\nu^s \text{CH}_2; (\text{C}_z)$
$S_{15}=r_{6,13}-r_{6,14}$	$\nu(\text{C}_6\text{H}_{13}) - \nu(\text{C}_6\text{H}_{14})$	$\nu^a \text{CH}_2; (\text{C}_z)$
$S_{16}=\beta_{3,5,4}-\beta_{2,4,3}+\beta_{1,3,2}-\beta_{6,2,1}+\beta_{5,1,6}-\beta_{6,4,5}$	ring def I	$\beta(\text{R}_1)$
$S_{17}=2\beta_{2,4,3}-\beta_{1,3,2}-\beta_{6,2,1}+2\beta_{1,5,6}-\beta_{6,4,5}-\beta_{3,5,4}$	ring def II	$\beta(\text{R}_2)$
$S_{18}=\beta_{1,3,2}-\beta_{6,2,1}+\beta_{6,4,5}-\beta_{3,5,4}$	ring def III	$\beta(\text{R}_3)$
$S_{19}=\beta_{4,15,3}-\beta_{2,15,3}$	$\beta(\text{N}_3\text{H})$	$\beta(\text{NH})$
$S_{20}=\beta_{3,7,2}-\beta_{1,7,2}+\beta_{3,8,4}-\beta_{5,8,4}$	$\beta(\text{C}_2\text{O}_7) + \beta(\text{C}_4\text{O}_8)$	$\beta^s(\text{C}_x\text{O})$
$S_{21}=\beta_{3,7,2}-\beta_{1,7,2}-\beta_{3,8,4}+\beta_{5,8,4}$	$\beta(\text{C}_2\text{O}_7) - \beta(\text{C}_4\text{O}_8)$	$\beta^a(\text{C}_x\text{O})$
$S_{22}=\beta_{13,14,6}$	sciss $\text{C}_6\text{HH}$	sciss $\text{CH}_2; (\text{C}_z)$
$S_{23}=\beta_{10,12,1}+\beta_{9,11,5}$	sciss $\text{C}_y\text{HH}$ (in-phase)	sciss $^s\text{CH}_2; (\text{C}_y)$
$S_{24}=\beta_{10,12,1}-\beta_{9,11,5}$	sciss $\text{C}_y\text{HH}$ (out-of-phase)	sciss $^a\text{CH}_2; (\text{C}_y)$
$S_{25}=\beta_{5,13,6}-\beta_{5,14,6}+\beta_{1,13,6}-\beta_{1,14,6}$	rock $\text{C}_6\text{HH}$	rock $\text{CH}_2; (\text{C}_z)$
$S_{26}=\beta_{5,13,6}+\beta_{5,14,6}-\beta_{1,13,6}-\beta_{1,14,6}$	wag $\text{C}_6\text{HH}$	wag $\text{CH}_2; (\text{C}_z)$
$S_{27}=\beta_{5,13,6}-\beta_{5,14,6}-\beta_{1,13,6}+\beta_{1,14,6}$	twist $\text{C}_6\text{HH}$	twist $\text{CH}_2; (\text{C}_z)$
$S_{28}=A_1-A_2+A_3-A_4+A_5-A_6+A_7-A_8$		rock <sup>s</sup> $\text{CH}_2; (\text{C}_y)$
$S_{29}=A_1-A_2+A_3-A_4-A_5+A_6-A_7+A_8$		rock <sup>a</sup> $\text{CH}_2; (\text{C}_y)$
$S_{30}=A_1+A_2-A_3-A_4+A_5+A_6-A_7-A_8$		wag <sup>s</sup> $\text{CH}_2; (\text{C}_y)$
$S_{31}=A_1+A_2-A_3-A_4-A_5-A_6+A_7+A_8$		wag <sup>a</sup> $\text{CH}_2; (\text{C}_y)$
$S_{32}=A_1-A_2-A_3+A_4+A_5-A_6-A_7+A_8$		twist <sup>s</sup> $\text{CH}_2; (\text{C}_y)$
$S_{33}=A_1-A_2-A_3+A_4-A_5+A_6+A_7-A_8$		twist <sup>a</sup> $\text{CH}_2; (\text{C}_y)$
$S_{34}=\gamma_{15,4,3,2}$	$\gamma(\text{N}_3\text{H})$	(NH)
$S_{35}=\gamma_{8,5,4,3}+\gamma_{7,3,2,1}$	$(\text{C}_2\text{O}_7) + (\text{C}_4\text{O}_8)$	$\gamma^s(\text{C}_x\text{O})$
$S_{36}=\gamma_{8,5,4,3}-\gamma_{7,3,2,1}$	$(\text{C}_2\text{O}_7) - (\text{C}_4\text{O}_8)$	$\gamma^a(\text{C}_x\text{O})$
$S_{37}=\tau_{4,3,2,1}-\tau_{3,2,1,6}+\tau_{2,1,6,5}-\tau_{1,6,5,4}+\tau_{6,5,4,3}-\tau_{5,4,3,2}$		$\tau(\text{R}_1)$
$S_{38}=\tau_{4,3,2,1}-\tau_{2,1,6,5}+\tau_{1,6,5,4}-\tau_{5,4,3,2}$		$\tau(\text{R}_2)$
$S_{39}=2\tau_{3,2,1,6}-\tau_{4,3,2,1}-\tau_{2,1,6,5}+2\tau_{6,5,4,3}-\tau_{1,6,5,4}-\tau_{5,4,3,2}$		$\tau(\text{R}_3)$

<sup>a</sup> The normalizing factors are omitted for clarity.  $r_{ij}$ , the distance between atoms  $i$  and  $j$ ;  $\beta_{ijk}$ , the angle between vectors  $k-i$  and  $k-j$ ;  $\gamma_{ijkl}$ , the angle between the vector  $k-i$  and the plane defined by atoms  $j,k,l$ ;  $\tau_{ijkl}$ , the dihedral angle between the plane defined by atoms  $ij,k$  and the plane defined by atoms  $j,k,l$ ;  $A_1=\beta_{6,10,1}$ ;  $A_2=\beta_{6,12,1}$ ;  $A_3=\beta_{2,10,1}$ ;  $A_4=\beta_{2,12,1}$ ;  $A_5=\beta_{6,9,5}$ ;  $A_6=\beta_{6,11,5}$ ;  $A_7=\beta_{4,9,6}$ ;  $A_8=\beta_{4,11,5}$ . Abbreviations: a, antisymmetric (out-of-phase); s, symmetric (in-phase) with respect to the plane perpendicular to the molecular plane and passing through N<sub>3</sub>, H<sub>15</sub>, C<sub>6</sub>, H<sub>13</sub>, and H<sub>14</sub> atoms.  $\nu$ , stretching;  $\beta$ , in-plane bending;  $\gamma$ , out-of-plane bending; def, deformation; sciss, scissoring; rock, rocking; wag, wagging; twist, twisting;  $\tau$ , torsion.

### SCHEME 1: Atom Numbering in Glutarimide



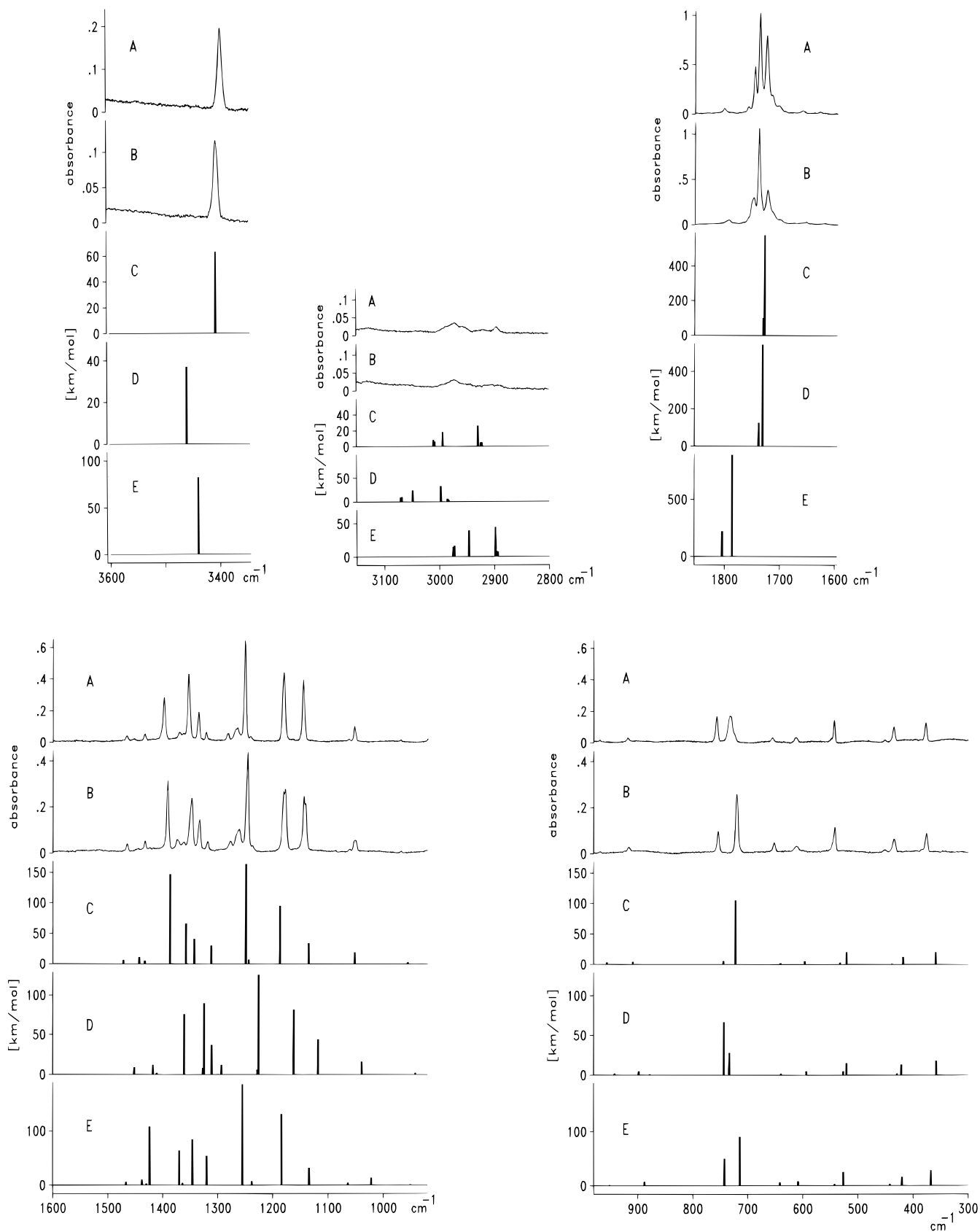
Unfortunately, the experimental gas-phase structure of glutarimide is unknown, whereas the structure of the molecule in the crystal is significantly distorted, due to a strong N-H...O hydrogen bonding, which perturbs the geometry of the ring. Comparison of the calculated and experimental data indicates that the largest discrepancies are encountered for the C-C, C-O, and N-H bond distances. The conformational lability of the ring may affect the experimentally determined C-C bond lengths, as it has been demonstrated in our recent X-ray studies,<sup>39</sup> while the differences in the C-O and N-H bond

lengths are understandable, since these bonds are the most susceptible to intermolecular interactions.

The best agreement with the X-ray data has been obtained at the HF level for C-C, C-N, and N-H bond lengths. However, the calculated C=O bond distance is too short, while it is much better predicted with the MP2 method. The BP86 functional considerably overestimates all experimental bond lengths. It should be noted that the calculated bond angles are well reproduced regardless of the theoretical method applied.

It is apparent that the calculated conformation of the molecule may be described as a half-chair, with one carbon atom (C<sub>6</sub>) slightly out of the essentially coplanar system, formed by five atoms of the glutarimide ring and two oxygen atoms. The calculated dihedral angle between the C<sub>1</sub>C<sub>6</sub>C<sub>5</sub> plane and the molecular framework is in excellent agreement with the experimental value, as is shown in Table 1. It can be concluded that such significant flattening of the glutarimide ring, predicted by theory and also confirmed by experimental data, enables some glutarimide drugs to intercalate between nucleic base pairs in the DNA helix.

**B. Vibrational Spectra.** The infrared spectra of glutarimide measured in low-temperature argon and nitrogen matrices are compared with the theoretical spectra in Figure 1. The



**Figure 1.** Comparison of the experimental IR spectra of glutarimide isolated in nitrogen (A) and argon (B) matrices, with the spectra predicted theoretically at the MP2/D95V\*\* (C), DFT(BP86)/D95V\*\* (D), and HF/D95V\*\* (E) levels.

experimental and calculated (HF, DFT, and MP2) frequencies and infrared intensities are listed in Table 3. Mode description (vibrational assignment) given in this table is based on potential energy distribution obtained at the MP2 level, except for the modes Q<sub>28</sub> and Q<sub>29</sub>, which are better described with PED

calculated at the HF level (as will be shown later). It should be emphasized that these three theoretical methods yielded very similar PED values for all corresponding normal modes but these two. Different PEDs have been indicated in the footnotes of Table 3. The overall agreement between the experimental and



is shifted by  $6\text{ cm}^{-1}$  toward higher wavenumbers. Such a shift is typical for  $\beta(\text{NH})$  bands. From the PED shown in Table 3, it is apparent that the corresponding normal mode ( $Q_{13}$ ) has mainly "NH-bend" character due to the predominant contribution (48%) from the  $\beta(\text{NH})$  internal coordinate, although a significant mixing with the C–N symmetric stretching vibration (32%) is also noted. It is remarkable that the frequency of this mode is so close to that of the corresponding mode, at 1399 and  $1395\text{ cm}^{-1}$ , in the infrared spectra of uracil and thymine, respectively.<sup>36</sup>

As can be seen from Table 3, the  $\beta(\text{NH})$  vibration in glutarimide contributes also to the mode  $Q_{18}$ , assigned to the very strong infrared band at  $1246\text{ cm}^{-1}$  in Ar ( $1251\text{ cm}^{-1}$  in  $\text{N}_2$  matrix). According to the calculated PED, this band can be considered as an analogue of the "amide III band" which occurs around  $1250\text{ cm}^{-1}$  in the spectra of amides and peptides.<sup>40</sup>

The NH out-of-plane bending vibration is particularly sensitive to the environment and susceptible to matrix effects, which may account for the differences in wavenumbers of the corresponding infrared band in low-temperature gas matrices.<sup>35–37</sup> It is also expected that this band will be markedly shifted in the spectra of the hydrogen-bonded systems. Thus, a definite assignment of this vibration is very important for the investigation of intermolecular interactions. In the infrared spectrum of glutarimide in an Ar matrix a strong band is observed at  $720\text{ cm}^{-1}$ . In a nitrogen matrix this band is considerably blue-shifted (by  $12\text{ cm}^{-1}$ ) and broadened, which indicates that it is associated with the NH out-of-plane  $\gamma(\text{NH})$  vibration. The frequency of the corresponding mode  $Q_{29}$  is very well predicted with three theoretical methods, which confirms our assignment. It follows from PED obtained for  $Q_{29}$  that the  $\gamma(\text{NH})$  vibration predominates in this mode; however the calculated contribution from other vibrations is different at each theory level. Furthermore, some discrepancies in PEDs have also been noted for the nearby  $Q_{28}$  mode. The relative infrared intensities of these two bands predicted with theoretical methods differ significantly from each other (see Figure 1 C,D,E). The MP2-calculated infrared intensity of the  $Q_{28}$  mode is seriously underestimated, whereas that of  $Q_{29}$  is overestimated with respect to the experiment.

It is worth noting that infrared intensities calculated at the MP2 level are usually more accurate than those calculated with the HF method using the same basis set.<sup>24,34–35,41–43</sup> However, in our recent MP2/6-31G\*\* calculations we have already encountered a similar problem, since this level of theory failed to predict frequencies and intensities of two normal modes of phenol.<sup>43</sup>

The relative intensities of modes  $Q_{28}$  and  $Q_{29}$  obtained with the DFT method (BP86 energy functional) are also incorrect since they are just opposite those of the experiment, as is shown in Table 3. The results obtained for these modes at the Hartree–Fock level are in much better agreement with the experiment, although the calculated intensity of  $Q_{28}$  is overestimated. It is evident from Table 3 that these discrepancies are related with the different PEDs, calculated at the three theory levels, for  $Q_{28}$  and  $Q_{29}$ . The plausible explanation of these discrepancies is that calculated infrared intensities depend on the forms of normal modes (which are represented by PED values). Different contributions of internal coordinates within these normal coordinates may sum up or nearly cancel the infrared intensity of the corresponding modes. It seems that PEDs for  $Q_{28}$  and  $Q_{29}$  calculated with the Hartree–Fock method best describe the nature of these modes; therefore the HF values have been used for assignment of the corresponding two infrared bands. However, it should be emphasized that the best overall agreement between the observed and calculated frequencies and

infrared intensities of all other modes of glutarimide has been obtained at the MP2/D95V\*\* level.

It should be noted that the corresponding  $\gamma(\text{NH})$  band occurs at much lower frequencies in the infrared spectra of pyrimidine bases in an Ar matrix:  $662$  and  $659\text{ cm}^{-1}$  for uracil<sup>36</sup> and 1-methyluracil,<sup>24</sup> respectively. This is due to the fact that this band arises from almost "pure" NH out-of-plane vibration in these bases.

**C=O Vibrations.** Infrared spectra of monomeric glutarimide exhibit a very complex pattern in the region  $1800\text{--}1600\text{ cm}^{-1}$ , where only two carbonyl stretching modes,  $\nu(\text{C=O})$ , are expected. It is interesting to note that similar complicated structures of the absorption bands arising from C=O stretching vibrations have been observed in the infrared spectra of matrix-isolated uracil, thymine, and its methyl derivatives.<sup>16,24,36–37,44–46</sup> It seems that such a splitting of bands in this frequency region is characteristic of the infrared spectra of all cyclic imides containing two polar C=O groups in the cis position.

It has been concluded that the fundamental transitions corresponding to the stretching motion of these two dipoles enter into strong and multiple Fermi resonances with several overtones and combinations of similar frequencies. The possible combination bands for glutarimide in this frequency range are  $1432 + 376 = 1808$ ;  $1348 + 451 = 1799$ ;  $1181 + 611 = 1792$ ;  $1246 + 541 = 1787$ ;  $1061 + 720 = 1781$ ;  $1392 + 376 = 1768$ ;  $1348 + 376 = 1724$ ;  $1181 + 541 = 1722$ ;  $968 + 754 = 1722$ ;  $1061 + 652 = 1713$ ;  $1334 + 376 = 1710$ ;  $1051 + 652 = 1703$ ;  $916 + 720 = 1636$ ;  $1246 + 376 = 1622\text{ cm}^{-1}$  (in an Ar matrix).

According to our calculations, the symmetric and antisymmetric C=O stretching vibrations of two equivalent carbonyl groups in glutarimide have very similar energy. The frequency of the former is only slightly higher than that of the latter. However, the predicted infrared intensities of these two C=O stretching modes are very different; the antisymmetric vibration generates the infrared band which is 4–5 times more intense than the symmetric counterpart. It should be emphasized that the theoretical results are consistent regardless of the method applied.

The predicted frequencies and the intensity pattern of  $\nu(\text{C=O})$  absorptions have proved to be of great help in assigning the spectra. In both matrices, the band of the highest intensity, located at  $1739\text{ cm}^{-1}$  in Ar and  $1737\text{ cm}^{-1}$  in  $\text{N}_2$ , is undoubtedly associated with the antisymmetric C=O stretching vibration ( $Q_9$  mode). The higher wavenumber band, observed at  $1748\text{ cm}^{-1}$  in an Ar matrix ( $1746\text{ cm}^{-1}$  in  $\text{N}_2$ ), has been attributed to the symmetric C=O stretching vibration ( $Q_8$ ). A third band in this cluster, observed at  $1723\text{ cm}^{-1}$  (Ar) and  $1725\text{ cm}^{-1}$  ( $\text{N}_2$ ), corresponds to some combination tone (as indicated above) which borrows infrared intensity from one of the  $\nu(\text{CO})$  fundamentals through the Fermi resonance mechanism. It should be noted that both the MP2 method and BP86 functional predicted well both the frequencies and infrared intensities of these two modes, whereas the HF method overestimated these values. The corresponding C=O stretching bands in matrix-isolated infrared spectra of pyrimidine bases have been found at very similar wavenumbers.<sup>24,36,37</sup>

The CO in-plane bending vibrations have been assigned to medium-intensity infrared bands at  $536$  and  $391\text{ cm}^{-1}$  in the spectrum of uracil<sup>36,37</sup> and at  $538$  and  $389\text{ cm}^{-1}$  in 1-methyluracil.<sup>24</sup> It is interesting to note that the corresponding antisymmetric and symmetric CO in-plane bending vibrations in glutarimide give rise to the medium-intensity bands at almost identical wavenumbers,  $541\text{ cm}^{-1}$  ( $Q_{33}$ ) and  $376\text{ cm}^{-1}$  ( $Q_{36}$ ), respectively. Although the calculated frequency of the former is consistently underestimated (by about  $15\text{--}20\text{ cm}^{-1}$ ), at all

theory levels, the predicted infrared intensities of these modes are in very good agreement with experimental values.

According to PED, the CO out-of-plane bending vibrations,  $\gamma(\text{CO})$ , in glutarimide are strongly mixed with ring vibrations and  $\text{CH}_2$  deformations; therefore they are not observed as characteristic bands, but contribute to several bands in the range from 860 to about  $500\text{ cm}^{-1}$ . The very weak infrared band at  $611\text{ cm}^{-1}$  corresponds to the mode  $Q_{31}$ , which contains a significant contribution from the antisymmetric  $\gamma(\text{CO})$ . The symmetric  $\gamma(\text{CO})$  vibration contributes to the mode  $Q_{32}$ , but the corresponding band is too weak to be observed in the experimental spectrum. It seems that the extensive coupling of  $\gamma(\text{CO})$  with other vibrations is the likely reason for the anomalously low intensities of the “ $\gamma(\text{CO})$  bands” in the infrared spectrum of glutarimide.

It should be noted that in 1-methyluracil the corresponding  $\gamma(\text{CO})$  modes have significantly higher infrared intensities and also higher frequencies, 802 and  $761\text{ cm}^{-1}$ .<sup>24</sup> This can be attributed to a different coupling of these vibrations in 1-methyluracil.

***CH<sub>2</sub> and Ring Vibrations.*** The C–H stretching region in the infrared spectra of cyclic amides and alkanes is noted for being strongly affected by anharmonic resonances.<sup>40</sup> In glutarimide Fermi resonances may occur between the C–H stretching fundamentals and overtones of  $\text{CH}_2$  deformation modes as well as combinations involving the NH bending or CH bending and C=O stretching vibrations, similarly as in substituted amides. It can be seen from Figure 1 that several weak absorptions appear in the region between  $3000$  and  $2850\text{ cm}^{-1}$ , both in argon and nitrogen matrices. Such a low intensity of the bands due to C–H stretching modes seems to be typical for infrared spectra of monomeric heterocycles isolated in low-temperature matrices.<sup>24,36,37</sup> Therefore, the C–H stretching vibrations of glutarimide have been assigned from the Raman spectrum of crystalline glutarimide,<sup>47</sup> where these modes generate two very intense bands, at  $2973$  and  $2902\text{ cm}^{-1}$ , and a shoulder at  $2925\text{ cm}^{-1}$ . It should be noted that the calculated (and scaled) HF and MP2 frequencies of these modes are quite close to the experimental values, as shown in Table 3. In the assignment of the C–H stretching modes we were guided by the fact that in compounds where the  $\text{CH}_2$  group is attached to a carbonyl group the corresponding symmetric  $\text{CH}_2$  stretching vibration is observed as a strong Raman band around  $2910\text{ cm}^{-1}$ .<sup>40,48</sup> Thus, it has been concluded that a very strong band at  $2902\text{ cm}^{-1}$  in the Raman spectrum of glutarimide corresponds to stretching vibrations in the  $\text{C}_\gamma\text{HH}$  groups, adjacent to  $\text{C}_\alpha=\text{O}$  ( $Q_6$  and  $Q_7$  modes). These two modes have almost identical frequencies, as revealed by calculations; therefore they may give rise to only one band observed in the spectrum. The next strong band at  $2973\text{ cm}^{-1}$  in the Raman spectrum may result from an overlap of three modes,  $Q_2$ ,  $Q_3$ , and  $Q_4$ , since their frequencies, predicted at the MP2 level, are very similar. A shoulder at  $2925\text{ cm}^{-1}$  has been assigned to the stretching vibration ( $Q_5$ ) in the  $\text{C}_\alpha\text{HH}$  group, which is supported by the presence of a similar band in the spectrum of cyclohexane.<sup>49</sup>

The methylene scissoring vibrations (sciss  $\text{CH}_2$ ) in glutarimide have been assigned by comparison with the spectra of related compounds. In cyclohexane, these modes generate weak absorptions in the range  $1475$ – $1450\text{ cm}^{-1}$ .<sup>49</sup> Bellamy<sup>40</sup> has noted for the compounds containing the methylene group adjacent to a carbonyl group that the position of this band is shifted to lower wavenumbers ( $1425$ – $1400\text{ cm}^{-1}$ ) while its infrared intensity increases. A similar effect has also been observed in the matrix-isolated infrared spectrum of glutarimide, and it was confirmed in our calculations. As can be seen from

Table 3, the calculated frequencies of  $Q_{11}$  and  $Q_{12}$  are lower than that of the mode  $Q_{10}$ . The two former correspond to the methylene scissoring vibrations in the  $\text{C}_\gamma\text{HH}$  groups (attached to  $\text{C}_\alpha=\text{O}$ ), whereas the latter arises from the scissoring vibration in the  $\text{C}_\alpha\text{HH}$  group. It seems that these characteristic bands corresponding to stretching and scissoring vibrations of different methylene groups in glutarimide can be used as the diagnostic bands in other spectroscopic studies on larger molecules (e.g. steroids) containing methylene groups in various environments.

The remaining  $\text{CH}_2$  deformation modes, wagging, twisting, and rocking vibrations, appear to be strongly coupled with different ring vibrations in glutarimide. According to PED, the  $\text{CH}_2$  wagging modes have a large contribution to several bands observed in the range  $1374$ – $1050\text{ cm}^{-1}$ , as shown in Table 3. The very strong band at  $1348\text{ cm}^{-1}$  (Ar) has been assigned to the mode  $Q_{15}$ , which corresponds to the “ring breathing vibration” coupled with the  $\text{CH}_2$  wagging vibrations. This is supported by a similar assignment of the strong, characteristic band observed at about  $1360\text{ cm}^{-1}$  in the infrared spectra of antineoplaston A10, the new antitumor drug containing glutarimide.<sup>48,50</sup> We, therefore, suggest that this infrared band can also be used as the “marker band”, for the presence of the glutarimide ring in large molecular systems. The next strong band at  $1246\text{ cm}^{-1}$  in the infrared spectrum of glutarimide in an Ar matrix (assigned to  $Q_{18}$ ) corresponds to the “amide III band”, as discussed earlier.

An assignment of the remaining bands in the range  $1374$ – $1262\text{ cm}^{-1}$  is quite difficult since these bands are weak, broad, and closely spaced. Bands at  $1374$ ,  $1334$ , and  $1318\text{ cm}^{-1}$  have been assigned to the fundamentals  $Q_{14}$ ,  $Q_{16}$ , and  $Q_{17}$ , respectively, since they correspond to bands of similar wavenumbers in the spectra of related compounds.<sup>40,49</sup> A doublet observed at  $1277/1262\text{ cm}^{-1}$  in Ar, which is slightly shifted in a  $\text{N}_2$  matrix, originates, most probably, from combination tones (for example,  $611 + 652 = 1263$  and  $541 + 720 = 1261$  in Ar) which borrow their intensity via anharmonic resonances with the nearby  $Q_{18}$  fundamental mode of the same symmetry. Similarly, a weak band at  $1362\text{ cm}^{-1}$  may arise from Fermi resonance between the combination ( $611 + 754 = 1365\text{ cm}^{-1}$ ) with the fundamental  $Q_{14}$  at  $1374\text{ cm}^{-1}$ . The methylene twisting vibrations ( $Q_{20}$  and  $Q_{21}$ ) have been assigned to the strong infrared bands observed at  $1181$  and  $1141\text{ cm}^{-1}$ . Band splitting observed for these modes in an argon matrix disappears in a nitrogen matrix; therefore, it is ascribed to a matrix effect. As follows from calculations, the third  $\text{CH}_2$  twisting vibration ( $Q_{19}$ ) in glutarimide generates a band of very low infrared intensity. Furthermore, its frequency is very similar to that of the  $Q_{18}$  mode; therefore it is probably hidden under the strong  $Q_{18}$  absorption. Theoretical data indicate that the  $\text{CH}_2$  rocking vibrations are extensively coupled with the ring deformation modes and contribute to extremely weak infrared bands in the range  $1060$ – $850\text{ cm}^{-1}$  and also to several bands in the low-frequency region.

The frequencies of ring deformations  $Q_{34}$ ,  $Q_{35}$ , and  $Q_{36}$  are quite well predicted by theory. As is seen in Table 2, a high level of conformity between the BP86/D95V\*\* and MP2/D95V\*\* results is observed in the range below  $640\text{ cm}^{-1}$ . The remaining three modes,  $Q_{37}$ ,  $Q_{38}$ , and  $Q_{39}$ , corresponding to torsion vibrations, were below the range covered by the matrix experiment. The predicted frequencies of these modes are in the range  $250$ – $90\text{ cm}^{-1}$ . A band observed at  $263\text{ cm}^{-1}$  in the Raman spectrum of solid glutarimide, which was assigned to the ring torsion vibration,<sup>47</sup> corresponds very well to the frequency of  $250\text{ cm}^{-1}$ , calculated at the MP2/D95V\*\* level. It should be mentioned that for uracils<sup>24,36</sup> torsion vibrations of the ring contribute to the normal modes of higher frequencies.

For 2(1*H*)-pyrimidinone<sup>35a</sup> or 4(3*H*)-pyrimidinone<sup>35b</sup> torsional modes give rise to infrared bands near 700 cm<sup>-1</sup>. The very low frequencies of the torsion vibrations of glutarimide correspond to a greater flexibility of the glutarimide ring when compared to the above-mentioned compounds.

## Conclusions

The important conclusions from the present study can be summarized as follows.

1. A good agreement between the calculated (HF, MP2, and DFT/BP86) and experimental infrared intensities and frequencies of glutarimide confirms the reliability of the presented vibrational band assignment.

2. The best overall agreement between the calculated and experimental spectra has been obtained at the MP2/D95V\*\* level, although for some modes, the infrared intensities predicted by HF or DFT methods agreed better with experiment.

3. The HF frequencies (after scaling) are overestimated for most modes, in particular, for C=O stretching vibrations, whereas frequencies calculated at the BP86/D95V\*\* level are closer to MP2 and experimental results.

4. The bond distances predicted with the DFT method (the BP86 functional) are too long, while the frequencies of C=O stretchings and other modes are very well reproduced. This effect can be attributed to some error cancellation.

5. A striking similarity is noted for frequencies of the corresponding CO and NH vibrations in glutarimide and in uracil, thymine, and their methyl derivatives.

6. The significant flattening of the glutarimide ring, predicted by calculations, indicates its structural resemblance to pyrimidine bases.

7. It is suggested that some glutarimide drugs are able to intercalate between nucleic base pairs in the DNA helix, or they may act as an antagonist of uracil and thymine in biological processes. This may interfere with nucleic acid or protein synthesis in cells. These conclusions are in accordance with previous experimental results indicating that some glutarimide drugs may interact with DNA.

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