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Time–resolved resonance FT–IR and raman spectroscopy and density functional theory investigation of vibronic–mode coupling structure in vibrational spectra of nanopolypeptide macromolecule beyond the multi–dimensional franck–condon integrals approximation and density matrix method

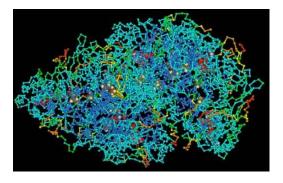
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### **Abstract**

A macromolecule is a very large molecule, such as protein, commonly created by the polymerization of smaller subunits (monomers). They are typically composed of thousands of atoms or more. The most common macromolecules in biochemistry are biopolymers (nucleic acids, proteins, carbohydrates and lipids) and large non–polymeric molecules (such as lipids and macrocycles). Synthetic macromolecules include common plastics and synthetic fibers as well as experimental materials such as carbon nanotubes. Parameters such as FT–IR and Raman vibrational wavelengths and intensities for single crystal Nano polypeptide macromolecule are calculated using density functional theory and were compared with empirical results. The investigation about vibrational spectrum of cycle dimers in crystal with carboxyl groups from each molecule of acid was shown that it leads to create Hydrogen bonds for adjacent molecules. The current study aimed to investigate the possibility of simulating the empirical values. Analysis of vibrational spectrum of Nano polypeptide macromolecule is performed based on theoretical simulation and FT–IR empirical spectrum and Raman empirical spectrum using density functional theory in levels of HF/6–31G\*, HF/6–31++G\*\*, MP2/6–31G, MP2/6–31++G\*\*\*, BLYP/6–31G, BLYP/6–31G and B3LYP6–31–HEG\*\*\*. Vibration modes of methylene, carboxyl acid and phenyl cycle are separately investigated. The obtained values confirm high accuracy and validity of results obtained from calculations.



Molecular structure of a Nano polypeptide macromolecule [1-42].

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Key words: Vibronic Structure, Vibrational Spectra, Multi-Dimensional Franck-Condon Integrals Approximation, Density Functional Theory (DFT), Nano Polypeptide Macromolecule, Non-Focal Functions of Becke, Correlation Functions of Lee-Yang-Parr, Time-Resolved Resonance, FT-IR and Raman Spectroscopy, Density Matrix Method (DMM)

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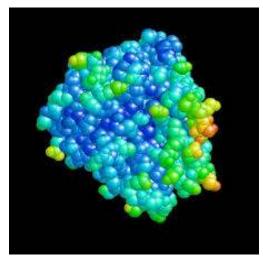
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### Introduction

A macromolecule is a very large molecule, such as protein, commonly created by the polymerization of smaller subunits (monomers). They are typically composed of thousands of atoms or more. The most common macromolecules in biochemistry are biopolymers (nucleic acids, proteins, carbohydrates and lipids) and large non-polymeric molecules (such as lipids and macrocycles). Synthetic macromolecules include common plastics and synthetic fibers as well as experimental materials such as carbon nanotubes. Density Functional Theory (DFT) is one of the most powerful calculation methods for electronic structures [5-7]. Numerous results have been previously studied and indicate successful use of these methods [8-10]. The theory is one of the most appropriate methods for simulating the vibrational wavenumbers, molecular structure as well as total energy. It may be useful to initially consider the calculated results by density functional theory using HF/6-31G\*, HF/6-31++G\*\*, MP2/6-31G, MP2/6-31++G\*\*, BLYP/6-31G, BLYP/6-31++G\*\*, B3LYP/6-31G and B3LYP6-31-HEG\*\* approach [11-16]. It should be noted that calculations are performed by considering one degree of quantum interference as well as polarization effects of 2d orbitals in interaction [17-364].

## **Details of calculations**

All calculations of molecular orbital in the base of ab are performed by Gaussian 09. In calculation process, the structure of Nano polypeptide macromolecule molecule (Figure 1) is optimized and FT-IR and Raman wavenumbers are calculated using HF/6-31G\*, HF/6-31++G\*\*, MP2/6-31G, MP2/6-31++G\*\*, BLYP/6-31G, BLYP/6-31++G\*\*, B3LYP/6-31G and B3LYP6-31-HEG\*\* base. All optimized structures are adjusted with minimum energy. Harmonic vibrational wavenumbers are calculated using second degree of derivation to adjust convergence on potential surface as good as possible and to evaluate vibrational energies at zero point. In optimized structures considered in the current study, virtual frequency modes are not observed which indicates that the minimum potential energy surface is correctly chosen. The optimized geometry is calculated by minimizing the energy relative to all geometrical quantities without forcing any constraint on molecular symmetry. Calculations were performed by Gaussian 09. The current calculation is aimed to maximize structural optimization using density functional theory. The calculations of density functional theory are performed by HF/6-31G\*, HF/6-31++G\*\*, MP2/6-31G,



**Figure 1.** Different sections of the extensive two–dimensional hydrogen bond network in a Nano polypeptide macromolecule [43-93]

MP2/6-31++G\*\*, BLYP/6-31G, BLYP/6-31++G\*\*, B3LYP/6-31G and B3LYP6-31-HEG\*\* function in which non-focal functions of Becke and correlation functions of Lee-Yang-Parr beyond the Franck-Condon approximation are used. After completion of optimization process, the second order derivation of energy is calculated as a function of core coordination and is investigated to evaluate whether the structure is accurately minimized. Vibrational frequencies used to simulate spectrums presented in the current study are derived from these second order derivatives. All calculations are performed for room temperature of 518 (K).

# Vibration analysis

Analysis of vibrational spectrum of Nano polypeptide macromolecule is performed based on theoretical simulation and FT–IR empirical spectrum and Raman empirical spectrum using density functional theory in levels of HF/6–31G\*, HF/6–31++G\*\*, MP2/6–31G, MP2/6–31++G\*\*, BLYP/6–31G, BLYP/6–31++G\*\*, B3LYP/6–31G and B3LYP6–31–HEG\*\*. Vibration modes of methylene, carboxyl acid and phenyl cycle are separately investigated.

C–H stretching vibrations in single replacement of benzene cycles are usually seen in band range of 3000–4000 cm<sup>-1</sup>. Weak Raman bands are at 3193 cm<sup>-1</sup> and 3207 cm<sup>-1</sup>. C–C stretching mode is a strong Raman mode at 1211cm<sup>-1</sup>. Raman weak band is seen at 1667 cm<sup>-1</sup>, too. Bending mode of C–H is emerged as a weak mode at 1388 cm<sup>-1</sup> and 1187 cm<sup>-1</sup> and a strong band at 1291 cm<sup>-1</sup> in Raman spectrum. Raman is considerably active in the range of 1000–2000 cm<sup>-1</sup> which 1189 cm<sup>-1</sup> indicates this issue.

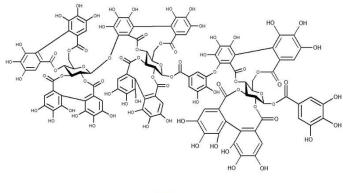
C–H skew–symmetric stretching mode of methylene group is expected at 3199  $\rm cm^{-1}$  and its symmetric mode is expected at 3001  $\rm cm^{-1}$ . Skew–symmetric stretching mode of CH $_2$  in Nano polypeptide macromolecule has a mode in mid–range of Raman spectrum at 3000–3530  $\rm cm^{-1}$ . When this mode is symmetric, it is at 3098  $\rm cm^{-1}$  and is sharp. The calculated wavenumbers of higher modes are at 3064  $\rm cm^{-1}$  and 3098  $\rm cm^{-1}$  for symmetric and skew–symmetric stretching mode of methylene, respectively.

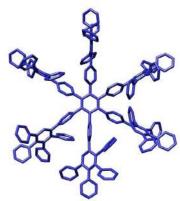
Scissoring vibrations of  $\mathrm{CH}_2$  are usually seen at the range of 1528–1589 cm<sup>-1</sup> which often includes mid–range bands. Weak bands at 1555 cm<sup>-1</sup> are scissoring modes of  $\mathrm{CH}_2$  in Raman spectrum. Moving vibrations of methylene are usually seen at 1473 cm<sup>-1</sup>. For the investigated chemical in the current study, these vibrations are at 1341 cm<sup>-1</sup> were calculated using density functional theory. Twisting and rocking vibrations of  $\mathrm{CH}_2$  are seen in Raman spectrum at 900 cm<sup>-1</sup> and 1200 cm<sup>-1</sup>, respectively, which are in good accordance with the results at 915 cm<sup>-1</sup> and 1185 cm<sup>-1</sup>, respectively.

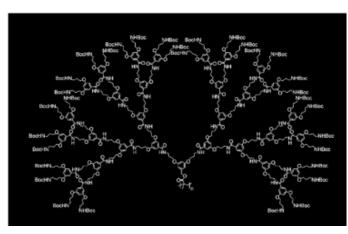
In a non-ionized carboxyl group (COOH), stretching vibrations of carbonyl [C=O] are mainly observed at the range of 1880–11890 cm<sup>-1</sup>. If dimer is considered as an intact constituent, two stretching vibrations of carbonyl for symmetric stretching are at 1765–1795 cm<sup>-1</sup> in Raman spectrum. In the current paper, stretching vibration of carbonyl mode is at 1803 cm<sup>-1</sup> which is a mid-range value.

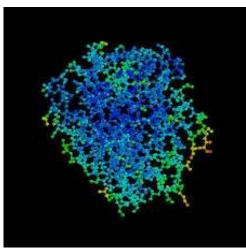
Stretching and bending bands of hydroxyl can be identified by width and band intensity which in turn is dependent on bond length of Hydrogen. In dimer form of Hydrogen bond, stretching band of O–H is of a strong Raman peak at 1387 cm<sup>-1</sup> which is due to in–plain metamorphosis mode. Out–of–plain mode of O–H group is a very strong mode of peak at 1056 cm<sup>-1</sup> of Raman spectrum. The stretching mode of C–O (H) emerges as a mid–band of Raman spectrum at 1267 cm<sup>-1</sup>.

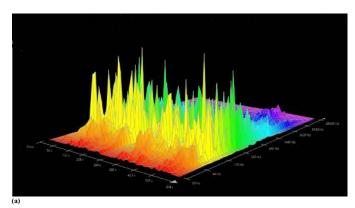
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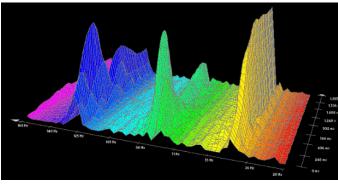


Figure 2. 3D Simulation of (a) FT-IR spectrum and (b) Raman spectrum of Nano polypeptide macromolecule

Lattice vibrations are usually seen at the range of 0–800 cm<sup>-1</sup>. These modes are induced by rotary and transferring vibrations of molecules and vibrations and are including Hydrogen bond. Bands with low wavenumbers of Hydrogen bond vibrations in FT–IR and Raman spectrum (Figure 2) are frequently weak, width and unsymmetrical. Rotary lattice vibrations are frequently stronger than transferring ones. Intra–molecular vibrations with low wavenumbers involving two–bands O–H ...O dimer at 89 cm<sup>-1</sup>, 196 cm<sup>-1</sup> and 259 cm<sup>-1</sup> are attributed to a rotary moving of two molecules involving in–plain rotation of molecules against each other.

# Conclusion and summary

Calculations of density functional theory using HF/6–31G\*, HF/6–31++G\*\*, MP2/6–31G, MP2/6–31++G\*\*, BLYP/6–31G, BLYP/6–31++G\*\*, B3LYP/6–31G and B3LYP6–31-HEG\*\* levels were used to obtain vibrational wavenumbers and intensities in single crystal of Nano polypeptide macromolecule. Investigation and consideration of vibrational spectrum confirm the formation of dimer cycles in the investigated crystal with carboxyl groups from each Hydrogen molecule of acid protected from adjacent molecules. The calculated vibrational spectrum which obtains from calculations of density functional theory is in good accordance with recorded empirical values which indicates successful simulation of the problem. The obtained results indicate that the results obtained from theoretical calculations are valid through comparing with empirical recorded results.

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