

COMMUNICATION

Intermolecular Interaction in 2-Aminopyridine: A Density Functional Study

Manoj Majumder, Tamal Goswami, Anirban Misra^{1*}, Soumik Bardhan, Swapan

K. Saha^{2*}

Department of Chemistry, University of North Bengal, Siliguri 734013, West Bengal, India

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Abstract: The absorption and fluorescence property of 2-amino pyridine (2-AP) and its dimer (2AP)₂ have been investigated using dispersion corrected density functional theory (DFT-D) method using B3LYP-D functional along with triple- ζ TZ2P basis in ADF suite of program. For the equilibrium geometries of the dimer, the ETS-NOCV calculations are performed. The formation of H-bond has been confirmed by the calculation of synergy (ΔE_{syn}) as well as from the spectral shift. A close agreement of the calculated spectra with that of experimental results has been found suggesting the dimers to be the preferable states of 2-AP in water medium.

AMS subject classifications: 74E40, 78M50

Key words: Density functional theory, 2-amino pyridine, absorption, fluorescence, H-bond

The excited state relaxation is of paramount importance in the photochemical process. Recently it has been noticed that the excited state relaxation can be induced through intermolecular Hydrogen bonding [1-4]. Photo induced proton-coupled electron transfers, where the proton donor and acceptor are held together by hydrogen bond (H-bond), have attracted considerable interest in recent years [5]. A number of studies by Han and co-workers [6-12] have shown that the gradual strengthening of the H-bond brings about the stabilization of the electronically excited state. H-bonded systems are ubiquitous within the

¹ Corresponding author. Email address: anirbanmisra@yahoo.com

² Corresponding author. Email address: ssahanbu@hotmail.com

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biological macromolecules and hence the H-bond induced relaxation of excited state is deemed to be important in biological systems as well (13-15, 16). Pyridine and its derivatives are currently finding immense interest in the scientific community due to their luminescence property. Some of amino pyridine derivatives show anaesthetic properties and are used as drugs for certain brain disease [17-19]. 2-amino pyridine (2-AP) is used in the preparation of cytidine which is used as drug [20]. In spite of this biomedical application the photochemistry of this molecule has not been thoroughly investigated.

In this study we report a systematic density functional investigation of the absorption and fluorescence property of 2-amino pyridine (2-AP) and its dimer (2AP)₂. As the dimerization of 2-AP occurs through hydrogen bonding (H-bonding), it becomes essential to unveil the effect of intermolecular H-bonding interactions in the photochemical behaviour of 2-AP. It has been reported earlier that H-bonding plays a pivotal role in the photochemical description of a molecule. Moreover, the simple electronic structure of this compound offers simplicity and encourages studying the effect of H-bonding interaction in the photochemistry of this compound as a benchmark.

For the computation of weak interactions like H-bond, the correlated *ab initio* techniques such as second-order Møller-Plesset perturbation theory (MP2) are in wide use [21]. In recent years *ab initio* and density functional theory (DFT) based methods have become popular tool in the investigations of structure and electronic properties of molecules. DFT offers a cost effective way for the quantum chemical investigation of the molecular properties and has been proved to be efficient in producing accurate results [22]. Although the density functional methods, like the popular B3LYP functional, are reported to be incompetent for the accurate description of H-bonding [23], the inclusion of correction due to dispersion interactions in the scenario has been proved to be effective. In the present work, we opted for the dispersion corrected DFT (DFT-D) method for the computation of absorption and fluorescence spectra. All the DFT and time dependent DFT (TDDFT) calculations are performed using ADF suit of program [24]. The B3LYP functional has been used, augmented with dispersion correction developed by Grimme (B3LYP-D) [25], along with triple- ζ TZ2P basis. The solvent effects have been estimated using the conductor-like screening model (COSMO) implemented in ADF [26].

Structures of the monomer and the dimer are optimized at the B3LYP-D/TZ2P level. The optimized ground state structures of the monomer and the dimer are represented in Figures 1 and 2. The dimer is found to prefer a non-planar geometry at ground and excited states (Figure 2).

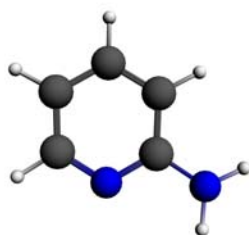


Figure 1: B3LYP-D/TZ2P optimized geometry of 2-Aminopyridine at its ground state.

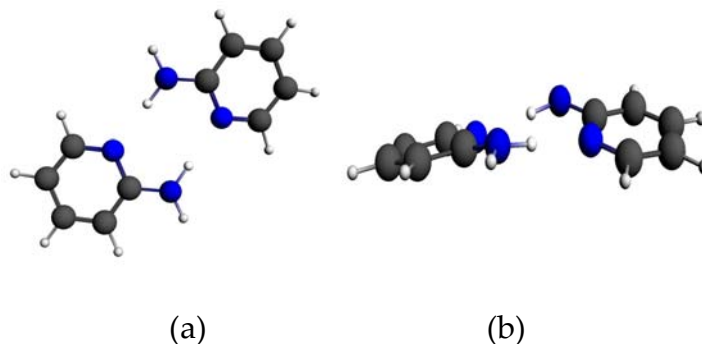


Figure 2: The H-bonded dimer (a) front view and (b) side view

To reveal the cooperativity due to H-bonding, an energy decomposition analysis (EDA) in the framework of the Kohn-Sham molecular orbital (MO) model is performed. In this approach, the interaction energy (ΔE_{int}) is partitioned into electrostatic interaction (ΔV_{elstat}), Pauli repulsive orbital interaction (ΔE_{pauli}), attractive orbital interaction (ΔE_{oi}) and dispersion interactions (ΔE_{disp}) as shown in the following,

$$\Delta E_{int} = \Delta V_{elstat} + \Delta E_{pauli} + \Delta E_{oi} + \Delta E_{disp} . \quad (1)$$

This method combines the extended transition state (ETS) energy decomposition approach [27] with the natural orbitals for chemical valence (NOCV) density analysis method [28]. The ETS-NOCV calculations are performed for the equilibrium geometries of the ground and excited states of the dimer. The formation of H-bond has been confirmed by the calculation of synergy (ΔE_{syn}) from the difference of interaction energy (ΔE_{int}) with and without H-bond interactions

$$\Delta E_{syn} = \Delta E_{int-eq} - \Delta E_{int-noneq} . \quad (2)$$

Here, ΔE_{int-eq} corresponds to the interaction energy of the dimer in its equilibrium geometry. Next, the equilibrium geometry is distorted by moving the monomer units far from each other, so that no H-bond formation is possible, and the interaction energy (ΔE_{int-

noneq) is estimated. A negative value of synergy indicates stabilization through the formation of H-bonding. ΔE_{int} is the energy change in systems due to formation of the association from individual units [29]. This is interpreted as the amount of energy required to promote the separated fragments to form the dimeric structure [30]. The synergy due to H-bonding is estimated through eq 2 and the results are given in Table 1.

Table 1: Bonding energy (kcal/mol) decomposition of ground and excited states of the dimer

Dimer	ΔV_{elstat}	ΔE_{pauli}	ΔE_{oi}	ΔE_{disp}	ΔE_{int}	ΔE_{syn}
Distorted Geometry	-0.30	-0.12	-0.03	-0.07	-0.52	--
Excited State	-25.93	29.92	-14.64	-4.30	-14.94	-14.42
Ground State	-26.24	30.31	-14.85	-4.26	-15.05	-14.53

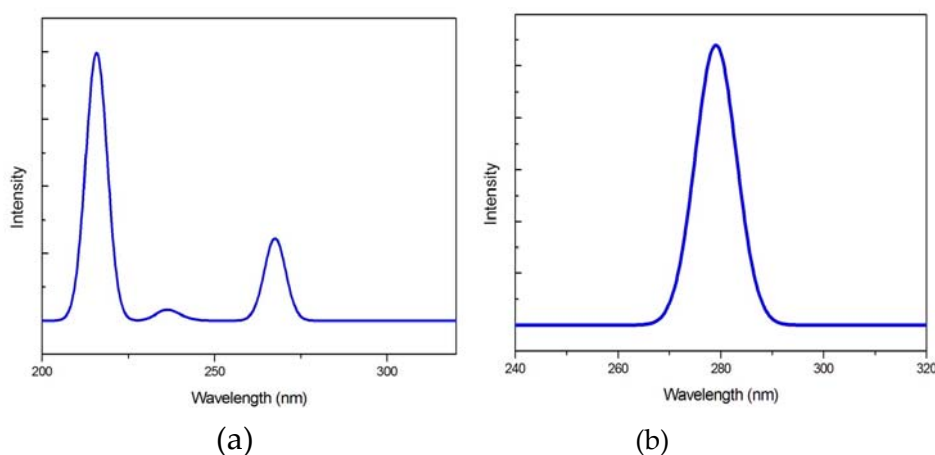


Figure 3: Absorption Spectra of (a) monomer unit and (b) hydrogen bonded dimer.

From Table 1, it is clear that the hydrogen-bond formation is possible at both the ground and excited states. From the comparison of ΔE_{syn} values in Table 1, one can infer a greater stabilization of the H-bonded dimer at its ground state than the corresponding excited state. However, from the components of ΔE_{int} it is observed that the dispersion energy change (ΔE_{disp}), which corresponds to the H-bonding interaction, is greater in the excited state than the ground state. Hence, it can be concluded that the H-bonding interaction is strengthened at the excited state.

The absorption and fluorescence spectra are shown in Figures 3 and 4. It becomes prominent from the figures that there occurs a red-shift of the absorption and fluorescence peaks in the dimeric state compared to the monomer (Table 2). This red-shift can be attributed to the lowering in the HOMO-LUMO gap (ΔE_{HL}) due to H-bond formation. Table

3 accumulates the HOMOs and the LUMOs of the species in its different states and a lowering of the ΔE_{HL} is clearly visible. From the previous studies on H-bonded systems it becomes evident that polarization effects promote H-bonding interaction [31]. It has also been seen that there occurs a mixing of the empty/occupied orbital on one fragment in the presence of another fragment due to polarization effect. From the nature of the HOMO and LUMO orbitals in Table 3, the mixing of two fragments is critically exposed. Hence, the diminished HOMO-LUMO gap can be explained as an outcome of the H-bonding interaction [32-33].

Table 2: Comparison of the calculated value λ_{max} of absorption and fluorescence with experimentally reported values.

System	λ_{max} of absorption in nm		λ_{max} of fluorescence in nm	
	Calculated	Experimental	Calculated	Experimental
2-AMP monomer	216	292 ^(a)	342	363 ^(a)
2-AMP dimer	279		359	

^(a) Experimentally reported (in reference [37]) λ_{max} of absorption and fluorescence in water medium

The red-shift in the fluorescence spectra can be explained from the stabilization of the S_1 electronic state of the dimer due to the formation of the H-bonding. The decrease in the energy gap for the $S_0 \leftarrow S_1$ leads to a red-shift of the fluorescence λ_{max} value.

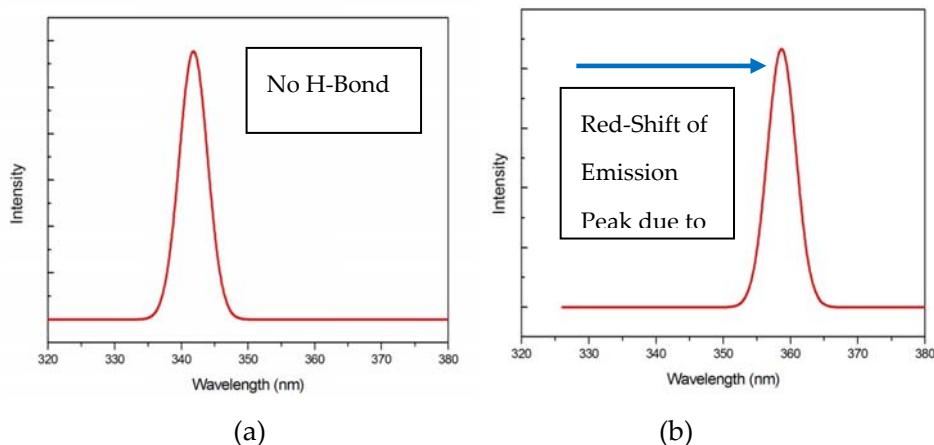
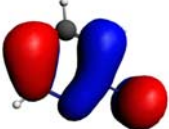
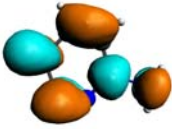
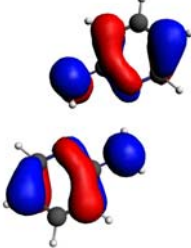
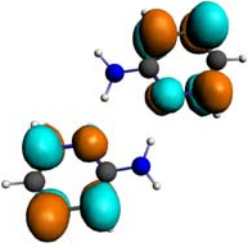


Figure 4: Fluorescence Spectra of (a) monomer unit and (b) hydrogen bonded dimer.

The calculated values of λ_{max} of both the absorption and fluorescence spectra of the 2-AP monomer and dimer are given in Table 2. The close agreement of the calculated values with

the experimental results suggests that the dimers are the most preferable states of 2-AP in water solution. This can also be inferred from the Table 1 since the high negative synergy of the dimer in both the ground and excited states indicates the formation of H-bonding. The formation or the change in the strength of the intermolecular hydrogen bonds following the excitation, may simply lead to new vibronic dissipative modes that couple the excited and the ground states [34-36]. Hence, the study of relaxation induced red shift in the emission spectra due to strengthening of H-bond seems to be useful tool to unveil many biological systems [38].

Table 3: HOMOs and LUMOs of the monomer and the dimer with corresponding energy values and HOMO-LUMO gap (ΔE_{HL}).

System	Orbitals	Energy in eV	ΔE_{HL} in eV
2-AP monomer	 HOMO	-5.8743	6.0044
	 LUMO	0.1257	
2-AP dimer	 HOMO	-5.6443	4.9446
	 LUMO	-0.6997	

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