# A density functional theory study on the porphyrin isomers: effect of meso-bridge length, relative stabilities, *cis-trans* isomerism

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Dedicated to Dr. A. V. Rama Rao on his 70th birthday

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#### **Abstract**

B3LYP/6-311+G\*\* calculations are employed to study the structures, stabilities of 1,2 (syn) and the 1,3 (anti) tautomeric forms of porphyrin isomers with varying meso-bridge lengths. A total of 46 structures obtained by considering the geometrical isomers of 1,2 (syn) and the 1,3 (anti) tautomeric forms, 44 were characterized as distinct minima on potential energy surface. Among the anti and syn tautomeric form the porphycene isomer 4a and 4a' is computed to be the most stable with a strong intramolecular hydrogen bonding. A  $-(CH)_2-$  linker in Z form with significant  $\pi$ -delocalization and the E form isomers are somewhat distorted into bowl-like structure in order to avoid the steric repulsion between the inner protons. In the isomers with trimethine interpyrrole linker, the E forms 7b and 7b' are having lower energy than the Z form due to the unstrained bond angles in these isomers and the Z form is destabilized by severe bond angle. In isomer [4.0.0.0], which has connectivity with the maximum possible bipyrrolic, linkage Z isomer is having high energy this is due to possibly strained bonds that are directly connected to the pyrrole rings.

Keywords: Porphyrin isomers, density functional theory, tautomerism, meso-bridges, minima

### Introduction

The study of porphyrins and related compounds has been developed into a subject of interdisciplinary interest due to their importance and relevance spanning a range of fields such as chemistry, biology and medicine. Porphyrins with their extended  $\pi$ -electron networks and exhibit high stability and have displayed applications in advanced materials as components in organic metals and in photodynamic therapy in the treatment of cancer and dermotological

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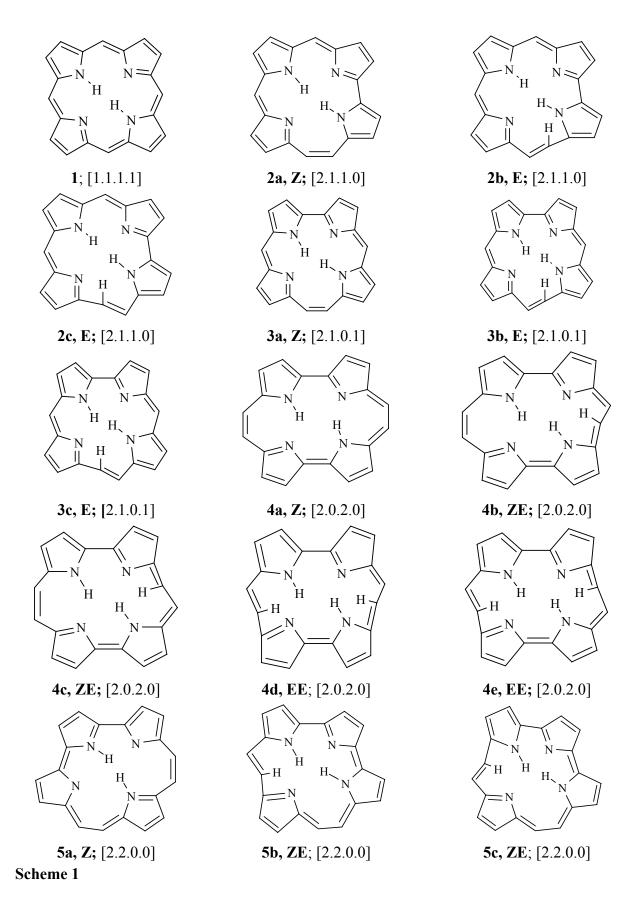
diseases.<sup>3</sup> The biological significance and the potential technological applications of porphyrins and related isomers are traced to their unique structural features.<sup>4,5,6</sup>

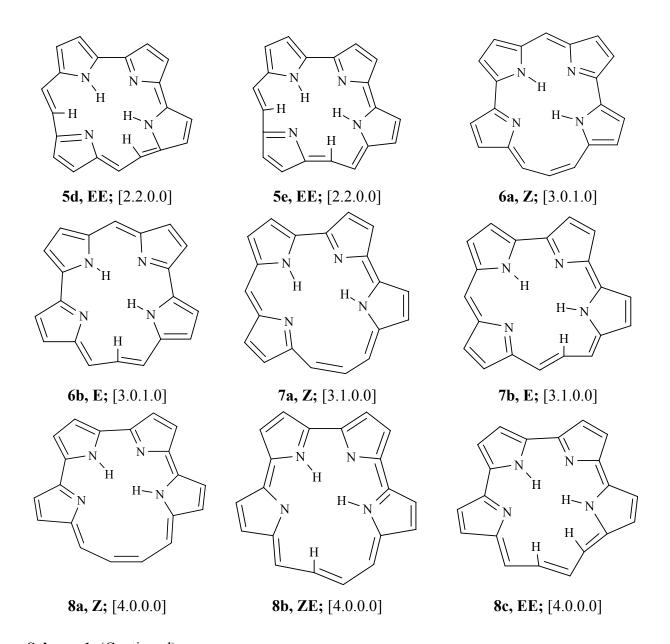
Recent successes in the syntheses of porphyrin isomers have opened up a new direction in porphyrin chemistry. The research groups of Vogel, Sessler, Furuta and others have synthesized a wide variety of porphyrin analogues. Many of the porphyrins are of interest as novel ligands for transition metals, and some have potential practical applications. Porphycenes, the first porphyrin isomers to be synthesized show promise as drugs in photodynamic cancer therapy. Structural variants such as ring or bridge extended, reshuffled, inverted invert

Wu et al. carried out density functional calculations using both the BLYP/3-21G and BLYP/6-31G\*\* methods on free-base porphyrin and its possible isomers with an N<sub>4</sub> – metal coordination core.<sup>20</sup> Ghosh et al. examined whether tetrapyrrolic isomers can accommodate trans double bonds.<sup>18</sup> while these studies addressed some important issues on the stability ordering of the porphyrin isomers, the structural and stabilities of various other possible porphyrin isomers were not considered. Considering the importance of the understanding the relative stability of the porphyrin isomers, we would like to explore the causative reasons such as angle strain, internal hydrogen bonding, inner proton repulsion, meso bridege length, cis-trans isomerism etc. <sup>20,21,25</sup>

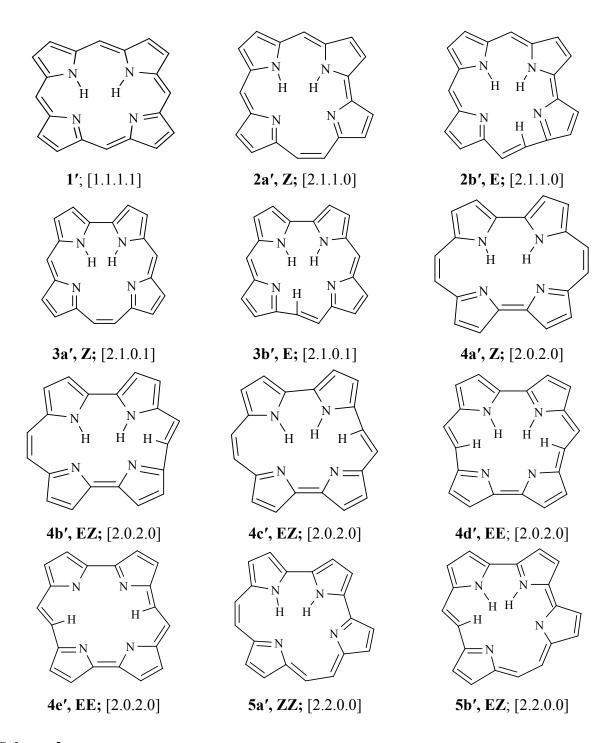
In this present study we concentrated particularly on tetrapyrrolic isomers of porphyrin isomers with N<sub>4</sub> cores. Two types of tautomers (a) *anti* (the imino protons on the I and III ring, Scheme 1); (b) *syn* (imino protons at II and I ring, Scheme 2) were taken into consideration. In this study, eight classes, depending on the nature of bridging, were considered, viz. freebase porphyrin [1.1.1.1], hemiporphycene [2.1.1.0], corrphycene [2.1.0.1], porphycene [2.0.2.0], isoporphycene [3.0.1.0], [2.2.0.0], [3.1.0.0] and [4.0.0.0]. Schemes 1 and 2 depicts the structures considered along with the general nomenclature used in the study for these isomers. Except for the freebase 1, all other classes involve geometrical isomerism Z (cis) and E (trans), so overall 46 porphyrin isomers of *syn* and *anti* tautomeric forms were studied.

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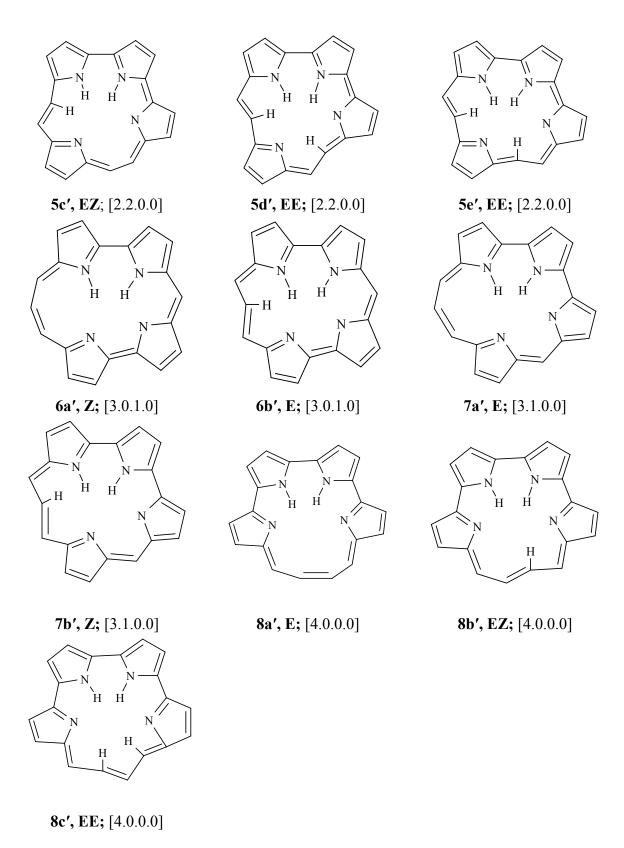




Scheme 1. (Continued).



Scheme 2



## Scheme 2. Continued

#### **Computational details**

B3LYP/6-31G calculations on all the structures considered in the study were optimized with planar geometries and the resultant stationary points were subjected to frequency calculations. Frequency calculations reveal that most of the planar structures do not correspond to minima. In such cases, following the normal modes of the imaginary frequencies, the minimum energy structures were obtained which display significant out-of-plane distortions. Many of these nonplanar minimum energy structures are highly puckered and in some cases the pyrrolic rings got inverted. This show that the structures have very high strain in the planar geometries, so we intended to calculate the energy difference between the planar structure and the puckered nonplanar structures and reported as distortion energies. The cavity size also large for the puckered structures. The EE isomers are more puckered compared to the EZ isomers. All the minima are characterized by frequencies calculations, which shows all real frequencies. Single point calculations were done using 6-311+G\*\* basis set on B3LYP/6-31G optimized geometries. 6-311+G\*\* basis set is a triple ζ basis set with polarization function on the heavy atoms and hydrogens as well as diffuse functions on the heavy atoms. The choice of using a B3LYP/6-31G geometries is entirely due to the economic reasons. As the previous studies 6,18-20 indicate that a better quality basis set is required to get better estimates, single point calculations were done at higher level of theory. All the calculations were performed using Gaussian 98 suite of programs.<sup>26</sup>

#### **Results and Discussion**

The different types of isomers are possible by varying methine bridges in addition to the parent porphyrin moiety. Considering all the possible geometric isomers, a total of 46 structures of 1,3 (anti) and 1,2 (syn) tautomers were considered in the present study, are given in Schemes 1 and 2. The relative energies were calculated from the energy difference between porphyrin isomer 1 in anti isomers and porphyrin isomer 1' in syn isomers.

## Anti and syn tautomeric forms

The relative energies, enthalpies, Highest Occupied Molecular Orbital (HOMO) and Lowest Unoccupied Molecular Orbital (LUMO) energies, distortion energies of *anti* tautomers of porphyrin isomers obtained at B3LYP level using the 6-31G and 6-311+G\*\* basis sets are given in Table 1. The B3LYP/6-31G optimized geometries of *anti* tautomeric structures and the relative energies with enthalpy correction are given in Figure 1. The variation in relative energies of *anti* (1,3) and *syn* (1,2) tautomeric porphyrin isomers obtained at B3LYP/6-31G and 6-311+G\*\* levels of theory are given in Figure 3. Considering all the possible cis-trans isomers of 1,3 (anti) and 1,2 (syn) tautomers totally 46 porphyrin isomers are resulted. Among 46 isomers 8c collapsed to the EZ isomer 8a and Z isomer 6a' collapsed to the anti tautomeric structure 6a due to closely presented imido protons. Thus, the total number of isomers reduced to 44. Porphyrin isomer 4a is the most stable isomeric form. The porphyrin isomer 4a is -0.4 kcal/mol more stable than the porphyrin isomer 1.

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**Table 1.** Relative energies ( $\Delta E$ , in kcal/mol), Enthalpies ( $\Delta H$ , in kcal/mol), HOMO ( $E_{HOMO}$ , in eV) and LUMO ( $E_{LUMO}$ , in eV) energies and the number of imaginary frequencies (NIMG) of *anti* tautomer of porphyrin isomers obtained at B3LYP level using the 6-31G and 6-311+G\*\* basis sets

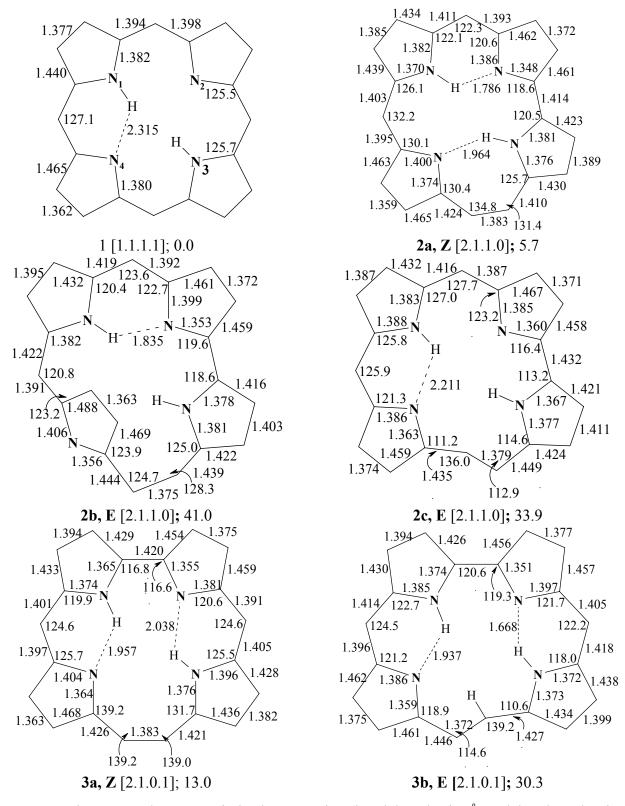
Struct.			B3LYP/6-311+G** <sup>a</sup>						
	ΔΕ	NIMG	$\Delta E^{b}$	ΔΗ	$E_{\text{HOMO}}$	$E_{LUMO}$	ΔΕ	$\Delta E^{b}$	$\Delta E^{c}$
1a	0.0	0	0.0	0.0	-5.16	-2.30	0.0	0.0	0.0
2a	4.4	0	0.0	0.0	-5.30	-2.60	5.7	0.0	5.7
<b>2</b> b	42.7	1	15.5	-0.1	-5.23	-2.77	41.1	15.6	41.0
<b>2</b> c	36.2	2	15.2	-0.7	-5.34	-2.77	34.6	15.7	33.9
3a	12.2	0	0.0	0.2	-5.35	-2.39	12.8	0.0	13.0
<b>3</b> b	29.9	1	1.9	-0.6	-5.32	-2.49	30.9	2.6	30.3
<b>3c</b>	29.7	1	2.1	-0.5	-5.33	-2.51	30.2	3.4	29.7
4a	-3.0	0	0.0	-0.7	-5.34	-2.88	0.3	0.0	-0.4
<b>4b</b>	56.3	1	14.0	-0.5	-5.12	-2.99	53.7	15.0	53.2
<b>4c</b>	47.2	2	28.2	-1.2	-5.30	-3.18	45.1	27.7	43.9
<b>4d</b>	82.6	3	30.2	-1.0	-5.09	-2.92	76.8	33.2	75.8
<b>4e</b>	82.9	2	26.7	-1.0	-5.09	-2.92	76.7	30.0	75.7
5a	33.5	1	0.1	0.3	-5.31	-2.96	32.7	0.4	33.0
<b>5</b> b	50.1	2	7.7	-0.4	-5.24	-3.06	48.8	8.8	48.4
5c	62.1	1	16.9	-0.6	-5.20	-2.95	57.7	19.3	57.1
5d	83.1	3	25.5	-0.9	-5.12	-2.96	77.0	29.1	76.1
5e	86.4	3	32.5	-1.2	-5.19	-3.11	79.2	35.7	78.0
6a	19.2	0	0.0	-0.1	-5.36	-2.48	21.2	0.0	21.1
<b>6b</b>	27.5	1	1.0	0.1	-5.26	-2.54	27.9	1.2	28.0
7a	42.2	0	0.0	0.2	-5.30	-2.78	41.4	0.0	41.6
<b>7b</b>	33.2	2	0.5	-0.2	-5.23	-2.79	33.0	1.1	32.8
8a	75.4	1	1.6	-1.2	-4.89	-2.64	70.6	4.2	69.4
8b	42.2	1	0.1	-0.3	-5.23	-3.04	41.0	1.6	40.7

<sup>&</sup>lt;sup>a</sup> Single point calculations were done on B3LYP/6-31G optimized geometries.

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<sup>&</sup>lt;sup>b</sup> The energy computed by taking the difference between the planar structure and the corresponding minimum energy structure. All these values are in kcal/mol.

<sup>&</sup>lt;sup>c</sup> The relative energies with enthalpy correction at B3LYP/6-311+G\*\* level.



**Figure 1.** The B3LYP/6-31G optimized geometries (bond lengths in Å and bond angles in degrees) of *anti* porphyrin isomers. The relative energies obtained at B3LYP/6-311+G\*\* with enthalpy correction are given in kcal/mol.

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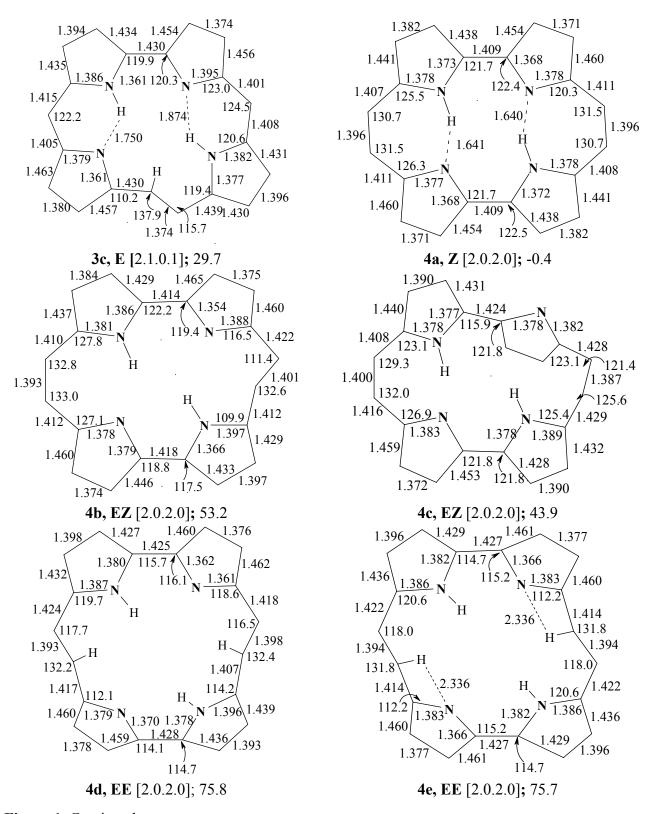


Figure 1. Continued

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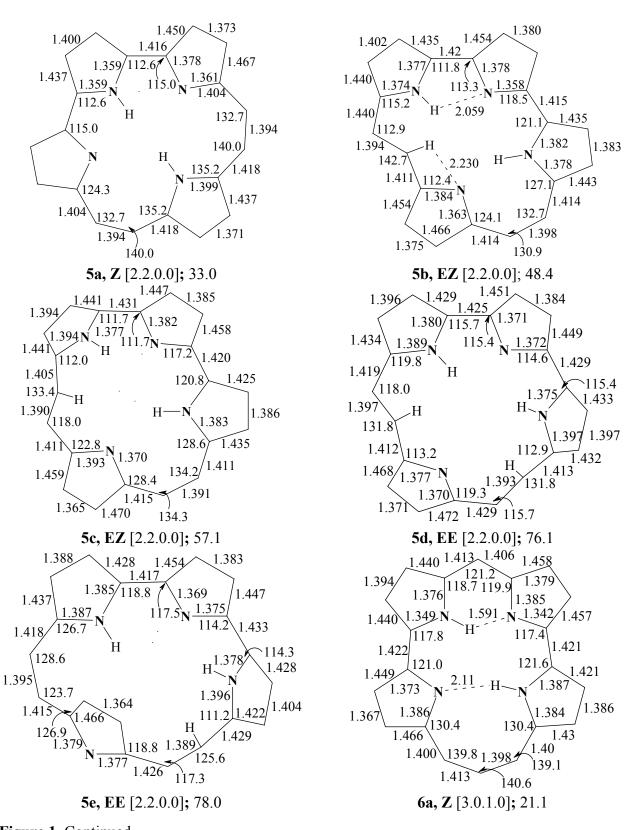


Figure 1. Continued

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1.453

1.417

1.373

1.443

1.449

1.409

1.379

117.0

1.428

1.386

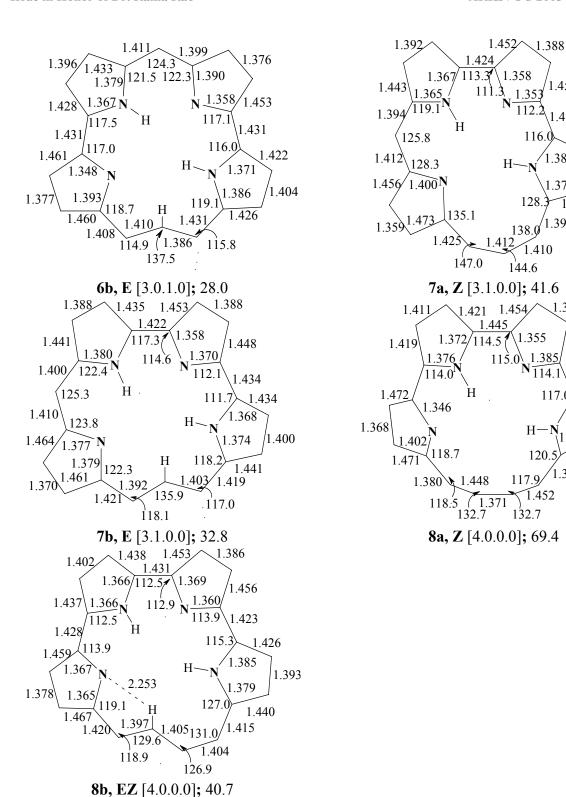
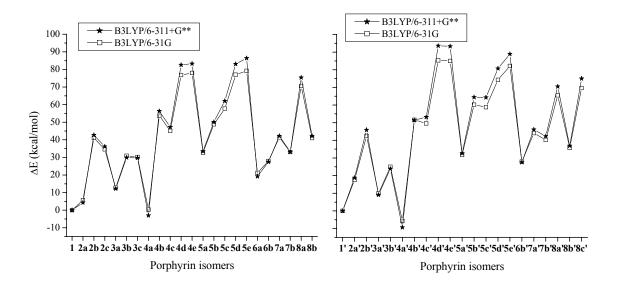


Figure 1. Continued

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Isomers 5a, 6a, 7a and 8a are much less stable than porphyrin isomer 4a and 1. In hemiporphycenes [2.1.1.0], the Z isomer 2a is more stable than the E isomers 2b and 2c. Isomer 2c is more stable than 2b by 7 kcla/mol and the isomer 2b undergoes ring inversion due to a highly repulsive interaction between tautomeric carbons and inner proton repulsion. In corrphycenes [2.1.0.1], the Z isomer 3a is more stable than the E isomers 3b and 3c. Isomers 3b and 3c are having energy difference around 0.6 kcal/mol. In porphycenes [2.0.2.0], three different isomeric forms are possible: Z form (4a), EZ form (4b and 4c) and EE form (4d and 4e). Among porphycene isomers, 4a is most stable due to strong hydrogen bonding. In EZ form and EE forms, the EZ form isomers 4b and 4c more stable than the EE form isomers 4d and 4e by 20-30 kcal/mol. The destabilizations of these isomers are traced due to the inner proton repulsion between the imino proton and proton present in the methine bridge. In isomer [2.2.0.0], like porphycene isomers, three different types of isomers are possible for these isomers. In EZ form and EE forms, the EZ form isomers 5b and 5c more stable than the EE form isomers 5d and 5e this is due to the presence of inner CH group. In isomers [3.0.1.0], the Z isomer is strongly favored energetically over the E isomer. The E forms 7b and 7b' are having lower energy than the Z form due to the unstrained bond angles in these isomers. In isomers [4.0.0.0] which has connectivity with the maximum possible bipyrrolic linkages and all the methine bridging groups are placed together, the EZ isomer 8b is having lower energy than the Z form 8a. Attempts to locate the 8c was futile, the structure is collapsing with the isomer 8a.

The relative energies, enthalpies, HOMO and LUMO energies, distortion energies of *syn* tautomers of porphyrin isomers obtained at B3LYP level using the 6-31G and 6-311+G\*\* basis sets are given in Table 2. The B3LYP/6-31G optimized geometries of *anti* tautomeric structures and the relative energies with enthalpy correction are given in Figure 2. The variation in relative energies of *anti* (1,3) and *syn* (1,2) tautomeric porphyrin isomers obtained at B3LYP/6-31G and 6-311+G\*\* levels of theory are given in Figure 2.



**Figure 2.** Relative energies of *anti* tautomers (a) and *syn* tautomers (b) of porphyrin isomers obtained at B3LYP/6-31G and 6-311+G\*\* levels of theory.

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In the *syn* tautomeric structures of porphyrin isomer, **4a'** is the most stable isomeric form and is about 6 kcal/mol more stable than porphyrin due to its exceptional strong hydrogen bonding. Isomers **5a'**, **7a'** and **8a'** are much less stable than porphyrin isomer **1'**. Similarly like *anti* porphyrin isomers, hemiporphycenes [2.1.1.0] and correphycenes [2.1.0.1], the Z isomer **2a** and **3a** are more stable than the E isomers **2b** and **3b** this is due to the inner proton repulsion.

**Table 2.** Relative energies ( $\Delta E$ , in kcal/mol), Enthalpies ( $\Delta H$ , in kcal/mol), HOMO ( $E_{HOMO}$ , in eV) and LUMO ( $E_{LUMO}$ , in eV) energies and the number of imaginary frequencies (NIMG) of *syn* tautomer of porphyrin isomers obtained at B3LYP level using the 6-31G and 6-311+G\*\* basis sets

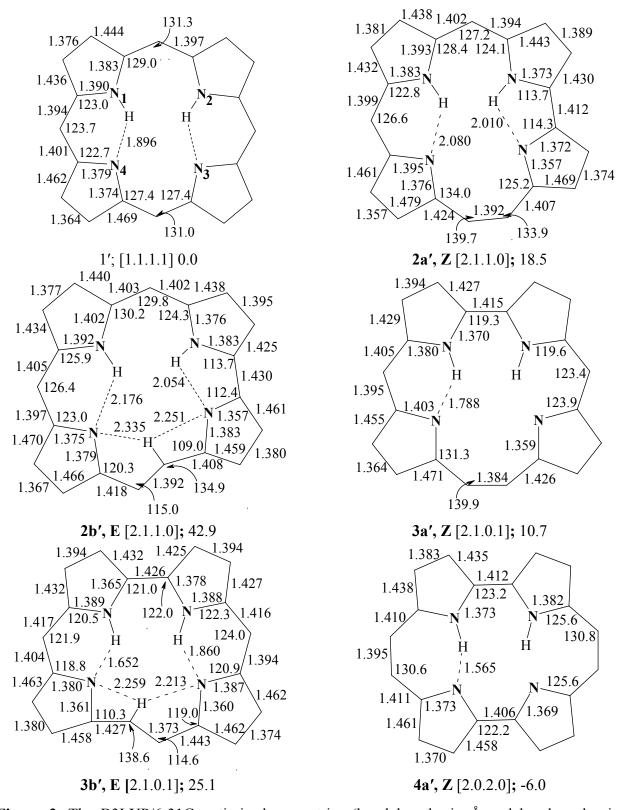
			B3LYP	B3LYP/6-311+G** <sup>a</sup>					
Struct.	ΔΕ	NIMAG	ΔE <sup>b</sup>	ΔΗ	Еномо	$E_{LUMO}$	ΔΕ	$\Delta E^{b}$	$\Delta E^{c}$
1a'	0.0	0	0.0	0.0	-5.19	-2.34	0.0	0.0	0.0
2a'	18.8	0	0.0	1.0	-5.24	-2.67	17.5	0.0	18.5
2b'	45.9	1	10.4	0.4	-5.11	-2.75	42.5	12.3	42.9
3a'	9.1	0	0.0	0.8	-5.36	-2.43	9.9	0.0	10.7
3b'	23.9	1	3.5	0.1	-5.31	-2.46	25.0	2.5	25.1
4a'	-9.3	0	0.0	-0.3	-5.33	-2.87	-5.7	0.0	-6.0
4b'	51.4	2	24.6	0.1	-5.20	-3.09	49.6	26.4	49.7
4c'	53.2	1	18.0	0.1	-5.25	-3.03	51.6	18.6	51.7
4d'	93.6	4	45.2	0.3	-5.14	-2.95	85.0	44.7	85.3
4e'	93.3	3	34.3	-0.3	-5.19	-2.95	85.3	31.9	85.0
5a'	32.7	0	0.0	1.2	-5.33	-2.98	31.8	0.0	33.0
5b'	64.4	3	13.4	0.7	-5.26	-3.09	60.2	13.3	60.9
5c'	64.3	3	25.6	0.5	-5.18	-2.99	58.8	24.6	59.3
5d'	80.7	3	32.9	0.3	-5.14	-2.95	74.3	29.9	74.6
5e'	88.9	3	29.7	0.0	-4.87	-2.95	82.1	27.1	82.1
6b'	27.6	3	2.5	0.4	-5.21	-2.53	27.9	2.2	28.3
7a′	46.1	1	0.5	1.0	-5.31	-2.82	44.1	0.1	45.1
7b′	42.3	1	3.2	0.9	-5.25	-2.84	40.2	2.9	41.1
8a'	70.5	2	10.4	0.1	-4.94	-2.53	65.5	9.2	65.6
8b'	36.9	2	3.4	1.0	-5.29	-3.04	35.7	2.3	36.7
8c'	75.0	1	29.2	0.2	-5.22	-3.09	69.6	26.2	69.8

<sup>&</sup>lt;sup>a</sup> Single point calculations were done on B3LYP/6-31G optimized geometries.

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<sup>&</sup>lt;sup>b</sup> The energy computed by taking the difference between the planar structure and the corresponding minimum energy structure. All these values are in kcal/mol.

<sup>&</sup>lt;sup>c</sup> The relative energies with enthalpy correction at B3LYP/6-311+G\*\* level.



**Figure 3.** The B3LYP/6-31G optimized geometries (bond lengths in Å and bond angles in degrees) of *syn* porphyrin isomers. The relative energies obtained at B3LYP/6-311+G\*\* with enthalpy correction are given in kcal/mol.

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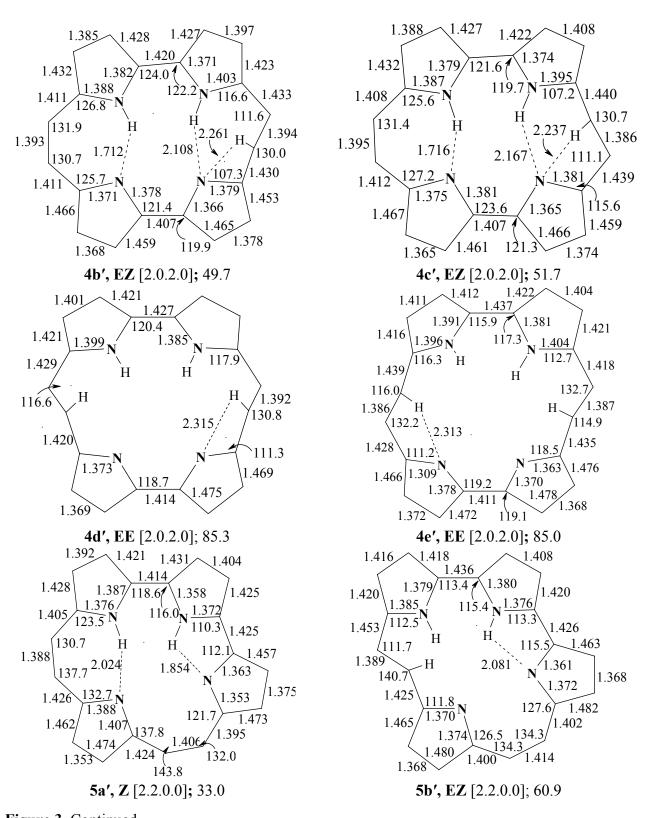


Figure 3. Continued

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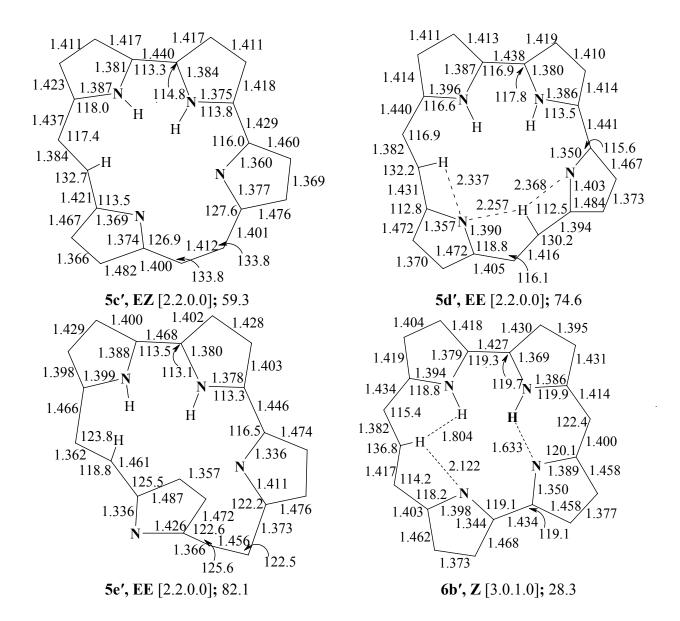


Figure 3. Continued

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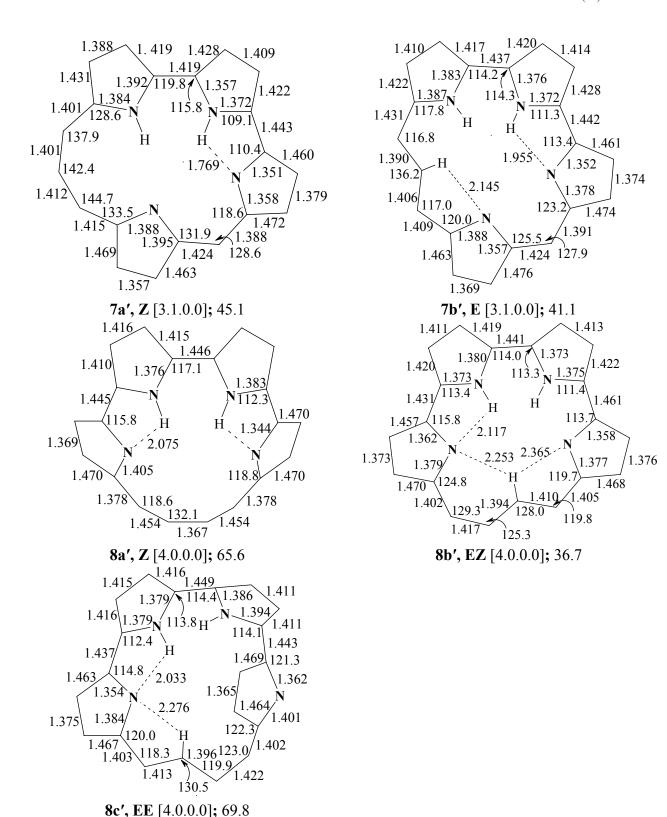


Figure 3. Continued

In porphycenes [2.0.2.0] and [2.2.0.0], like *anti* porphyrin isomers there are three different isomeric forms. Isomer **4a'** is most stable than all other porphycene isomers. This is due to strong hydrogen bonding. In EZ form and EE forms, the EZ form isomers **4b'** and **4c'** more stable than the EE form isomers **4d'** and **4e'**. The Z isomer **5a** is more stable than the EZ isomers **5b** and **5c** and **EE** isomer **5d** and **5e**. In EZ form and EE forms, the EZ form isomers **5b** and **5c** more stable than the EE form isomers **5d** and **5e** this is due to the presence of inner CH group.

In isomers [3.0.1.0], attempts to locate the planar form of **6a'** was futile, the structure is collapsing with the *anti* tautomeric isomer **6a**. In [3.1.0.0], the E form is more stable than the Z forms. The E form **7b'** are having lower energy than the Z form due to less steric strain in these isomers. In isomers [4.0.0.0], which have connectivity with the maximum possible bipyrrolic linkages the EZ, isomer **8b** is having lower energy than the Z and EE form **8a** and **8c** due to less angle strain. Comparison of Table 1 and 2 indicates that the trends in the relative energy ordering of the 1,3 (anti) and 1,2 (syn) tautomeric forms are very similar.

#### Geometrical isomerism

Except for the porphyrin isomers 1 and 1' geometrical isomerism is possible for all the isomeric forms considered for the *syn* and *anti* isomers. The relative stabilities of the geometrical isomers have attracted the theoreticians and experimentalist. The E and Z form especially affects the metal-binding properties of ligand. The important bond lengths and bond angles for all the *anti* and *syn* porphyrin isomers obtained at B3LYP/6-31G optimization are given in Figure 3.

In hemiporphyrin [2.1.1.0] and corrporphycene [2.1.0.1], three isomers are considered in *anti* tautomeric forms (one Z form and two E forms). In *syn* tautomeric forms one Z form and one E form is considered. In porphycene [2.0.2.0] and [2.2.0.0] five isomers one Z form, two EZ form and two EE form in both *syn* and *anti* tautomeric forms. The isomers [2.1.1.0], [2.1.0.1], [2.0.2.0], [2.2.0.0] having two –(CH)– linkages, prefer Z isomers 2a, 2a', 3a, 3a', 4a, 4a', 5a and 5a' this preference was due to the less steric interactions involving the inner CH group comparing to its E, EE and EZ isomers.

In isomers isoporphycene [3.0.1.0] and [3.1.0.0], one E form and one Z form was considered in both *syn* and *ant*i tautomers. In isomers [4.0.0.0], which have connectivity with the maximum possible bipyrrolic linkages one Z form, one EZ form and one EE form was considered. In [3.1.0.0] the E forms are more stable than Z forms. The E forms **7b**, **7b'** are more stable than the Z forms **7a** and **7a'** due to the less angle strain in these isomers. Isomers [4.0.0.0] **8a** and **8a'** are computed to be least stable compare to all other Z isomers and EZ isomer **8b** and **8b'** favored than Z isomers (**8a** and **8a'**) and EE isomers (**8c'**), this may be due to less angle strain in the EZ form. In isomers porphycenes [2.0.2.0], [2.2.0.0] the EZ isomers are more stable than the EE isomers in both the *anti* and *syn* tautomeric forms.

#### Effect of meso-bridge length on the relative stabilities

The relative energy differences seem to critically depend on the length and position of the meso bridges. In hemiporphycenes [2.1.1.0], corrporphycenes [2.1.0.1], porphycenes [2.0.2.0],

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[2.2.0.0] having two –(CH)– linkages, prefer Z isomers 2a, 2a', 3a, 3a', 4a, 4a', 5a and 5a' this preference was due to the less steric interactions involving the inner CH group comparing to its E, EE and EZ isomers.

When a  $-(CH)_2$ - linker is in the Z form, isomers are nearly planar with significant  $\pi$ -delocalization. In contrast, the corresponding E isomers have distorted structures with bowl-like geometries, which are traced due to steric repulsive interactions involving the inner protons. But, [3.1.0.0] the E forms are more stable than the Z forms, indicating higher angle strain in the latter. In isomers [4.0.0.0], which have connectivity with the maximum possible bipyrrolic linkages, Z isomer is computed to be relatively unstable. The variation in number of methine groups drastically alters the stability in these class of compounds.

#### Angle strain

Next to the hydrogen-bonding interactions, the strained bond angles dictate the relative stability ordering of the porphyrin isomers. All skeletal bond angle for the *anti* and *syn* structures are given in Figure 1 and Figure 3 respectively. The lower stability of isomers **5a**, **5a'**, **6a**, **7a**, **7a'**, **8a** and **8a'** in both syn and anti forms may directly traced to the severe angle strain in these isomers. The isomers of types [2.2.0.0], isoporphycenes [3.0.1.0], [3.1.0.0], [4.0.0.0] are destabilized due to the absence of bridging methine groups, which results in the angle strain for the tetrapyrroles and possibly due to the strained bonds that are directly connect to pyrrole rings. Bond angle strain is perhaps the most important factor contributing to the high energies in Z form in all [3.0.1.0], [4.0.0.0] isomers. The E isomers **7b** and **7b'** are having lower energy than the Z form, due to the unstrained bond angles in these isomers and the Z form is destabilized by severe bond angle strain in the trimethine interpyrrole linker. Isomers **8a** and **8c'** are computed to be the least stable among the [4.0.0.0.] isomers. The EZ isomers (**8b** and **8b'**) appear to be much less strained and importantly one of the hydrogens of the methine chain involve in hydrogen bond with the nitrogens.

#### **Distortion energies**

The  $\pi$ -delocalization and planarity has been treated as important factors to impart stability to porphyrins. However, as the planar form of a large number of isomers are not minima on the potential energy hypersurface, the puckering is often related to the strain in the planar form. We termed the energy difference between the planar structure and the corresponding minimum energy structure as the distortion energy. The distortion energies, which quantifies the amount strain in the planar geometry of the isomers, of the *anti* and *syn* porphyrin isomers are given in Table 1 and Table 2. In *anti* porphyrin isomers all the Z form isomers except **5a** and **8a** are planar. All other isomers have positive distortion energies, indicating that the nonplanar structures are more stable. In *syn* porphyrin isomers all the Z form isomers except **7a'** and **8a'** are planar. Isomers having positive distortion energies, signifying that nonplanar structures are more stable. The distortion energies are more for the EE form isomers than EZ forms. This is due to the inner proton repulsions in case of EE isomers.

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### **Hydrogen bonding**

Porphyrin isomers show some strong hydrogen bonds in most of the structures due to the close proximity of the four nitrogen atoms. In *anti* porphyrin isomer 1, the N---H--N distance is 2.315 Å and in *syn* porphyrin isomer 1' the distance is 1.896 Å (Figure 1 and 3). In *anti* and *syn* porphycene isomers 4a and 4a' features strong hydrogen bonding with H---N distances 1.640 Å and 1.565 Å respectively. Strong hydrogen bonding exists in isomers 6a and 3b. Isomers 3a-3c, 2a', 3a', 3b', 4b', 4c', 5a' and 7b' shows moderate hydrogen bonding. The difference between the stabilities of corrphycenes (3a-3c and 3a'-3b') and hemiporphycenes (2a-2c and 2a'-2b') can also to be traced by ring strain, as hemiporphycene is better hydrogen bonded than corrphycene. In general the *syn* tautomers have strong hydrogen bonds compare to the corresponding *anti* tautomers. In stability of all the low energy porphyrin isomers is clearly a manifestation of strong NH---N hydrogen bond resulting in short hydrogen bonding distances. Extremely long and unequal H-N-C angles leads to a high energy isomer.

#### **Central cavity**

The four adjacent N-N distances of all the *syn* and *anti* porphyrin isomers are given in Tables 3 and 4. The sum of the four N-N distances of the individual isomers may be taken as a measure of the effective cavity sizes of the porphyrin isomers. These measures give a straightforward indication of the cavity size of the porphyrin isomers. The inner protons were not considering when estimating the cavity sizes as normally complexation is invariably associated with displacement of the protons.

The porphyrin isomer 1 is having large cavity size with a square geometry (2.937Å). In corrphycenes, isomers 3b and 3c have large N<sub>3</sub>-N<sub>4</sub> distances 4.123 and 4.151 Å respectively. In isomers 4d, 4e, 5b and 5c the imino proton distances are very large due to more puckering of these isomers. In porphycenes, the isomers with EE form (4d and 4e) have large N<sub>2</sub>-N<sub>3</sub> and N<sub>4</sub>-N<sub>1</sub> distances and isomer 4e has similar N<sub>1</sub>-N<sub>2</sub> and N<sub>3</sub>-N<sub>4</sub> distances. Isomers 5b, 5c and 5d (N<sub>3</sub>-N<sub>4</sub> distance also large) have large N<sub>4</sub>-N<sub>1</sub> distances range from 4.207 to 4.246 Å this is due to the distortion into bowl-like structure in order to avoid the steric repulsion between the inner CH group. Isomers with [3.0.1.0], [3.1.0.0] and [4.0.0.0] connectivity the N<sub>3</sub>-N<sub>4</sub> various from 3.098 to 4.589 Å. Like *anti* porphyrin isomers the *syn* porphyrin isomers 4d, 4e, 5b, 5c and 5d isomers also have large N<sub>4</sub>-N<sub>1</sub> distances. Isomer 4d has similar N<sub>4</sub>-N<sub>1</sub> and N<sub>2</sub>-N<sub>3</sub> distances. In isomers [4.0.0.0], which have connectivity with the maximum possible bipyrrolic linkage, Z isomer and EZ isomer have large N<sub>4</sub>-N<sub>1</sub> distance. Isomers with [3.1.0.0] and [4.0.0.0] connectivity the imino proton distance is less compare to other *syn* porphyrin isomers.

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**Table 3.** The adjacent N---N distances and imino proton distances (Å) of *anti* tautomer of porphyrin isomers obtained at B3LYP level using the 6-31G basis set

Struct.	N <sub>1</sub> -N <sub>2</sub>	N <sub>2</sub> -N <sub>3</sub>	N <sub>3</sub> -N <sub>4</sub>	N <sub>4</sub> -N <sub>1</sub>	Н-Н
1a	2.937	2.937	2.937	2.937	2.208
2a	2.375	2.760	2.897	3.236	2.354
<b>2</b> b	2.646	2.756	-a-	-a-	2.470
<b>2</b> c	2.951	2.574	3.693	2.840	2.448
3a	2.642	2.735	3.555	2.727	2.268
<b>3</b> b	2.793	2.534	4.123	2.691	2.569
3c	2.796	2.684	4.151	2.549	2.744
4a	2.864	2.627	2.864	2.627	2.375
<b>4b</b>	2.823	3.326	2.729	2.887	2.391
4c	-a-	-a-	2.865	2.649	2.576
4d	2.624	4.178	2.565	4.153	3.305
<b>4e</b>	2.590	4.173	2.590	4.173	3.334
5a	2.522	3.257	3.257	2.522	2.010
5b	2.497	2.780	2.695	4.246	3.182
5c	2.472	2.766	3.052	4.376	3.109
<b>5</b> d	2.610	2.601	4.107	4.207	2.755
5e	2.725	2.597	-a-	-a-	3.216
6a	2.451	2.766	3.098	2.758	2.242
6 <b>b</b>	2.680	2.650	3.732	2.674	2.254
7a	2.458	2.534	3.660	2.817	2.352
<b>7</b> b	2.620	2.457	4.291	2.772	2.605
8a	2.586	2.606	4.589	2.613	2.635
8b	2.489	2.567	4.267	2.506	2.491

<sup>-</sup>a- the imidazole ring is inverted.

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**Table 4.** The adjacent N---N distances and imino proton distances (Å) of *syn* tautomer of porphyrin isomers obtained at B3LYP level using the 6-31G basis set

Struct.         N <sub>1</sub> -N <sub>2</sub> N <sub>2</sub> -N <sub>3</sub> N <sub>3</sub> -N <sub>4</sub> N <sub>4</sub> -N <sub>1</sub> H-H           1a'         3.241         2.692         3.158         2.692         2.125           2a'         2.978         2.528         3.284         2.866         1.746           2b'         3.168         2.518         4.012         2.883         2.158           3a'         2.756         2.633         3.623         2.633         2.083           3b'         2.864         2.513         4.104         2.658         2.335           4a'         2.916         2.582         2.867         2.582         2.340           4b'         2.933         3.070         2.821         2.690         2.306           4c'         2.834         3.001         2.885         2.704         2.147           4d'         2.827         3.889         2.732         3.888         2.008           4e'         2.707         3.875         2.750         4.065         2.076           5a'         2.672         2.414         3.396         2.995         1.892           5b'         2.595         2.588         4.115         4.063         2.044						
2a'         2.978         2.528         3.284         2.866         1.746           2b'         3.168         2.518         4.012         2.883         2.158           3a'         2.756         2.633         3.623         2.633         2.083           3b'         2.864         2.513         4.104         2.658         2.335           4a'         2.916         2.582         2.867         2.582         2.340           4b'         2.933         3.070         2.821         2.690         2.306           4c'         2.834         3.001         2.885         2.704         2.147           4d'         2.827         3.889         2.732         3.888         2.008           4e'         2.707         3.875         2.750         4.065         2.076           5a'         2.672         2.414         3.396         2.995         1.892           5b'         2.595         2.558         3.047         3.927         1.949           5c'         2.612         2.580         3.113         4.142         2.033           5d'         2.726         2.588         4.115         4.063         2.044           5e'	Struct.	$N_1$ - $N_2$	$N_2$ - $N_3$	N <sub>3</sub> -N <sub>4</sub>	$N_4$ - $N_1$	Н-Н
2b'       3.168       2.518       4.012       2.883       2.158         3a'       2.756       2.633       3.623       2.633       2.083         3b'       2.864       2.513       4.104       2.658       2.335         4a'       2.916       2.582       2.867       2.582       2.340         4b'       2.933       3.070       2.821       2.690       2.306         4c'       2.834       3.001       2.885       2.704       2.147         4d'       2.827       3.889       2.732       3.888       2.008         4e'       2.707       3.875       2.750       4.065       2.076         5a'       2.672       2.414       3.396       2.995       1.892         5b'       2.595       2.558       3.047       3.927       1.949         5c'       2.612       2.580       3.113       4.142       2.033         5d'       2.726       2.588       4.115       4.063       2.044         5e'       2.664       2.589       -a-       -a-       -a-       2.206         6b'       2.781       2.525       2.745       3.544       2.059 <t< th=""><th>1a'</th><th>3.241</th><th>2.692</th><th>3.158</th><th>2.692</th><th>2.125</th></t<>	1a'	3.241	2.692	3.158	2.692	2.125
3a'       2.756       2.633       3.623       2.633       2.083         3b'       2.864       2.513       4.104       2.658       2.335         4a'       2.916       2.582       2.867       2.582       2.340         4b'       2.933       3.070       2.821       2.690       2.306         4c'       2.834       3.001       2.885       2.704       2.147         4d'       2.827       3.889       2.732       3.888       2.008         4e'       2.707       3.875       2.750       4.065       2.076         5a'       2.672       2.414       3.396       2.995       1.892         5b'       2.595       2.558       3.047       3.927       1.949         5c'       2.612       2.580       3.113       4.142       2.033         5d'       2.726       2.588       4.115       4.063       2.044         5e'       2.664       2.589       -a-       -a-       -a-       2.206         6b'       2.781       2.525       2.745       3.544       2.059         7a'       2.705       2.363       3.003       3.399       1.948 <t< th=""><th>2a'</th><th>2.978</th><th>2.528</th><th>3.284</th><th>2.866</th><th>1.746</th></t<>	2a'	2.978	2.528	3.284	2.866	1.746
3b'       2.864       2.513       4.104       2.658       2.335         4a'       2.916       2.582       2.867       2.582       2.340         4b'       2.933       3.070       2.821       2.690       2.306         4c'       2.834       3.001       2.885       2.704       2.147         4d'       2.827       3.889       2.732       3.888       2.008         4e'       2.707       3.875       2.750       4.065       2.076         5a'       2.672       2.414       3.396       2.995       1.892         5b'       2.595       2.558       3.047       3.927       1.949         5c'       2.612       2.580       3.113       4.142       2.033         5d'       2.726       2.588       4.115       4.063       2.044         5e'       2.664       2.589       -a-       -a-       2.206         6b'       2.781       2.525       2.745       3.544       2.059         7a'       2.705       2.363       3.003       3.399       1.948         7b'       2.596       2.480       2.923       4.181       1.981         8a' <t< th=""><th>2b'</th><th>3.168</th><th>2.518</th><th>4.012</th><th>2.883</th><th>2.158</th></t<>	2b'	3.168	2.518	4.012	2.883	2.158
4a'       2.916       2.582       2.867       2.582       2.340         4b'       2.933       3.070       2.821       2.690       2.306         4c'       2.834       3.001       2.885       2.704       2.147         4d'       2.827       3.889       2.732       3.888       2.008         4e'       2.707       3.875       2.750       4.065       2.076         5a'       2.672       2.414       3.396       2.995       1.892         5b'       2.595       2.558       3.047       3.927       1.949         5c'       2.612       2.580       3.113       4.142       2.033         5d'       2.726       2.588       4.115       4.063       2.044         5e'       2.664       2.589       -a-       -a-       2.206         6b'       2.781       2.525       2.745       3.544       2.059         7a'       2.705       2.363       3.003       3.399       1.948         7b'       2.596       2.480       2.923       4.181       1.981         8a'       2.699       2.567       4.662       2.568       1.923         8b' <t< th=""><th>3a'</th><th>2.756</th><th>2.633</th><th>3.623</th><th>2.633</th><th>2.083</th></t<>	3a'	2.756	2.633	3.623	2.633	2.083
4b'       2.933       3.070       2.821       2.690       2.306         4c'       2.834       3.001       2.885       2.704       2.147         4d'       2.827       3.889       2.732       3.888       2.008         4e'       2.707       3.875       2.750       4.065       2.076         5a'       2.672       2.414       3.396       2.995       1.892         5b'       2.595       2.558       3.047       3.927       1.949         5c'       2.612       2.580       3.113       4.142       2.033         5d'       2.726       2.588       4.115       4.063       2.044         5e'       2.664       2.589       -a-       -a-       -a-       2.206         6b'       2.781       2.525       2.745       3.544       2.059         7a'       2.705       2.363       3.003       3.399       1.948         7b'       2.596       2.480       2.923       4.181       1.981         8a'       2.699       2.567       4.662       2.568       1.923         8b'       2.579       2.492       4.198       2.571       1.900 <th>3b'</th> <th>2.864</th> <th>2.513</th> <th>4.104</th> <th>2.658</th> <th>2.335</th>	3b'	2.864	2.513	4.104	2.658	2.335
4c'       2.834       3.001       2.885       2.704       2.147         4d'       2.827       3.889       2.732       3.888       2.008         4e'       2.707       3.875       2.750       4.065       2.076         5a'       2.672       2.414       3.396       2.995       1.892         5b'       2.595       2.558       3.047       3.927       1.949         5c'       2.612       2.580       3.113       4.142       2.033         5d'       2.726       2.588       4.115       4.063       2.044         5e'       2.664       2.589       -a-       -a-       2.206         6b'       2.781       2.525       2.745       3.544       2.059         7a'       2.705       2.363       3.003       3.399       1.948         7b'       2.596       2.480       2.923       4.181       1.981         8a'       2.699       2.567       4.662       2.568       1.923         8b'       2.579       2.492       4.198       2.571       1.900	4a'	2.916	2.582	2.867	2.582	2.340
4d'       2.827       3.889       2.732       3.888       2.008         4e'       2.707       3.875       2.750       4.065       2.076         5a'       2.672       2.414       3.396       2.995       1.892         5b'       2.595       2.558       3.047       3.927       1.949         5c'       2.612       2.580       3.113       4.142       2.033         5d'       2.726       2.588       4.115       4.063       2.044         5e'       2.664       2.589       -a-       -a-       2.206         6b'       2.781       2.525       2.745       3.544       2.059         7a'       2.705       2.363       3.003       3.399       1.948         7b'       2.596       2.480       2.923       4.181       1.981         8a'       2.699       2.567       4.662       2.568       1.923         8b'       2.579       2.492       4.198       2.571       1.900	4b'	2.933	3.070	2.821	2.690	2.306
4e'       2.707       3.875       2.750       4.065       2.076         5a'       2.672       2.414       3.396       2.995       1.892         5b'       2.595       2.558       3.047       3.927       1.949         5c'       2.612       2.580       3.113       4.142       2.033         5d'       2.726       2.588       4.115       4.063       2.044         5e'       2.664       2.589       -a-       -a-       2.206         6b'       2.781       2.525       2.745       3.544       2.059         7a'       2.705       2.363       3.003       3.399       1.948         7b'       2.596       2.480       2.923       4.181       1.981         8a'       2.699       2.567       4.662       2.568       1.923         8b'       2.579       2.492       4.198       2.571       1.900	4c'	2.834	3.001	2.885	2.704	2.147
5a'       2.672       2.414       3.396       2.995       1.892         5b'       2.595       2.558       3.047       3.927       1.949         5c'       2.612       2.580       3.113       4.142       2.033         5d'       2.726       2.588       4.115       4.063       2.044         5e'       2.664       2.589       -a-       -a-       -a-       2.206         6b'       2.781       2.525       2.745       3.544       2.059         7a'       2.705       2.363       3.003       3.399       1.948         7b'       2.596       2.480       2.923       4.181       1.981         8a'       2.699       2.567       4.662       2.568       1.923         8b'       2.579       2.492       4.198       2.571       1.900	4d'	2.827	3.889	2.732	3.888	2.008
5b'       2.595       2.558       3.047       3.927       1.949         5c'       2.612       2.580       3.113       4.142       2.033         5d'       2.726       2.588       4.115       4.063       2.044         5e'       2.664       2.589       -a-       -a-       -a-       2.206         6b'       2.781       2.525       2.745       3.544       2.059         7a'       2.705       2.363       3.003       3.399       1.948         7b'       2.596       2.480       2.923       4.181       1.981         8a'       2.699       2.567       4.662       2.568       1.923         8b'       2.579       2.492       4.198       2.571       1.900	4e'	2.707	3.875	2.750	4.065	2.076
5c'       2.612       2.580       3.113       4.142       2.033         5d'       2.726       2.588       4.115       4.063       2.044         5e'       2.664       2.589       -a-       -a-       -a-       2.206         6b'       2.781       2.525       2.745       3.544       2.059         7a'       2.705       2.363       3.003       3.399       1.948         7b'       2.596       2.480       2.923       4.181       1.981         8a'       2.699       2.567       4.662       2.568       1.923         8b'       2.579       2.492       4.198       2.571       1.900	5a'	2.672	2.414	3.396	2.995	1.892
5d'       2.726       2.588       4.115       4.063       2.044         5e'       2.664       2.589       -a-       -a-       -a-       2.206         6b'       2.781       2.525       2.745       3.544       2.059         7a'       2.705       2.363       3.003       3.399       1.948         7b'       2.596       2.480       2.923       4.181       1.981         8a'       2.699       2.567       4.662       2.568       1.923         8b'       2.579       2.492       4.198       2.571       1.900	5b'	2.595	2.558	3.047	3.927	1.949
5e'       2.664       2.589       -a-       -a-       2.206         6b'       2.781       2.525       2.745       3.544       2.059         7a'       2.705       2.363       3.003       3.399       1.948         7b'       2.596       2.480       2.923       4.181       1.981         8a'       2.699       2.567       4.662       2.568       1.923         8b'       2.579       2.492       4.198       2.571       1.900	5c'	2.612	2.580	3.113	4.142	2.033
6b'       2.781       2.525       2.745       3.544       2.059         7a'       2.705       2.363       3.003       3.399       1.948         7b'       2.596       2.480       2.923       4.181       1.981         8a'       2.699       2.567       4.662       2.568       1.923         8b'       2.579       2.492       4.198       2.571       1.900	5d'	2.726	2.588	4.115	4.063	2.044
7a'       2.705       2.363       3.003       3.399       1.948         7b'       2.596       2.480       2.923       4.181       1.981         8a'       2.699       2.567       4.662       2.568       1.923         8b'       2.579       2.492       4.198       2.571       1.900	5e'	2.664	2.589	-a-	-a-	2.206
7b'       2.596       2.480       2.923       4.181       1.981         8a'       2.699       2.567       4.662       2.568       1.923         8b'       2.579       2.492       4.198       2.571       1.900	6b'	2.781	2.525	2.745	3.544	2.059
8a'       2.699       2.567       4.662       2.568       1.923         8b'       2.579       2.492       4.198       2.571       1.900	7a′	2.705	2.363	3.003	3.399	1.948
<b>8b'</b> 2.579 2.492 4.198 2.571 1.900	7b'	2.596	2.480	2.923	4.181	1.981
	8a'	2.699	2.567	4.662	2.568	1.923
<b>8c'</b> 2.632 -aa- 2.533 2.107	8b'	2.579	2.492	4.198	2.571	1.900
oc 2.052 u u 2.555 2.107	8c'	2.632	-a-	-a-	2.533	2.107

<sup>-</sup>a- the imidazole ring is inverted.

# **Conclusions**

The present study reports density functional calculations on a series of *syn* and *anti* tautomeric forms of porphyrin isomers. The relative energy differences seem to critically depend on the length and position of the meso bridges. The isomers [2.1.1.0], [2.1.0.1], [2.0.2.0], [2.2.0.0]

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having two -(CH)- linkages, prefer Z isomer than E isomers. Isomers [2.2.0.0], [3.0.1.0], [3.1.0.0], [4.0.0.0] are destabilized by the presence of severe angle strain in the  $-(CH)_n$ -linkages. The present computational study touches upon a few important aspects of porphyrin chemistry, such as the relative stabilities of tautomers, geometrical isomers. This study provides a good insight about the geometry, energetics and reactivity of various porphyrin isomers and the perturbations that are caused to these isomers upon shuffling hydrogen positions. The variation in number of methine groups drastically alters the stability in these classes of compounds

Several factors, such as angle strain, inner proton repulsion, hydrogen bonding, length of the bridge and the type of geometrical isomerism- E or Z, play important roles in deciding the relative stability ordering of the porphyrin isomers. In  $-(CH)_2$ - linkers the Z forms are more stable than E form due to significant  $\pi$ -delocalization and hydrogen bonding, even in the presence of significant angle strain in it. The E form isomers are distorted into bowl-like structure in order to avoid the steric repulsion between the inner CH group. In trimethine interpyrrole linker E forms are having lower energy than the Z form due to the unstrained bond angles in these isomers. In isomers [4.0.0.0], which have connectivity with the maximum possible bipyrrolic linkages, Z isomer is having high energy this is due to possibly strained bonds that are directly connect to pyrrole rings.

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