

Hydrogen Bonding Influence of 1,10-Phenanthroline on Five-Coordinate High-Spin Imidazole-Ligated Iron(II) Porphyrinates

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The influence of a hydrogen bond to the coordinated imidazole on the geometric and electronic structure of iron has been further studied in new complexes of five-coordinate high-spin imidazole-ligated iron(II) porphyrinates. With 1,10-phenanthroline (1,10-phen) as the hydrogen-bond acceptor, several new octaethylporphyrin dianion (OEP) and *meso*-tetraphenylporphyrin dianion (TPP) derivatives have been synthesized and characterized by X-ray crystallography and Mössbauer spectroscopy. In all three new structures, the porphyrin molecules and 1,10-phenanthroline molecules have been found with a ratio of 1:1. All the porphyrin derivatives are five-coordinate 2-methylimidazole-ligated iron(II) species. 1,10-Phenanthroline is hydrogen bonded to the coordinated imidazole to form two unequal hydrogen bonds. The Fe–N_p and Fe–N_{im} bond lengths and displacement of the iron atom out of the porphyrin plane are similar to those in imidazole-ligated species. Mössbauer measurements showed remarkable temperature dependence; the analysis of the data obtained in applied magnetic field for [Fe(OEP)(2-MeHIm)] · (1,10-phen) gave a negative quadrupole splitting value and large asymmetry parameters. All the structural and Mössbauer properties suggest that these new hydrogen-bonded species have the same electronic configuration as imidazole-ligated species.

Introduction

Many biologically important proteins have heme (iron porphyrin) as the active site. These heme proteins have a variety of functions, including oxygen carriers in mammals, electron carriers in photosynthesis and respiration, and catalysts for a variety of biochemical reactions involving O₂, H₂O₂, etc. Their functionalities have been influenced by many factors. Hydrogen bonding is one such factor believed to play an important role. One important hydrogen bond in heme proteins is between the coordinated imidazole of a histidine residue and an adjacent amino acid residue. Crystal structures of heme proteins show that such hydrogen bonds are very widespread.^{1–3} The strength of these hydrogen bonds is thought to vary from very weak proton donation to

complete proton donation to form the imidazolate ligand.^{4–8} There has been substantial speculation that hydrogen bonding in heme proteins could control the relative stability of the oxidation state of iron and thus control reactivity of the heme protein.⁹

For example, a particularly striking pair of heme protein families are involved in such hydrogen bonds, the globins and the peroxidases. The heme in globins is the same *b*-type heme in the peroxidase heme, and they both have a coordinated imidazole from a histidyl residue of the protein chain as the proximal ligand. The proximal imidazole in the globins forms a hydrogen bond to a backbone carbonyl oxygen atom. In peroxidases, such as horseradish peroxidase,^{10,11} the proximal histidine (His170) is ligated to the

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heme center and also forms a hydrogen bond with a highly conserved aspartate group Asp247, a stronger hydrogen bond acceptor than that of backbone carbonyl oxygen. But they have much different chemistry. The biological function of globins is the reversible transport and storage of dioxygen, whereas peroxidases can catalyze the conversion of hydrogen peroxide to water and/or the oxidation of substrates. It has been postulated that the hydrogen bond stabilizes higher oxidation states of iron and distinctly alters the chemical behavior of the peroxidases relative to the globins. The importance of such hydrogen bonds has been commented on and calculated on but, to our knowledge, has not been experimentally investigated in a systematic manner.

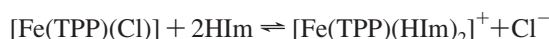
We have been studying the nature of various five-coordinate iron(II) porphyrinates.^{12–17} Much of this work has focused on the nature of high-spin imidazole-ligated species.^{13,14,16} As part of that work, we reported¹⁵ the effects of deprotonating the imidazole ligand in five-coordinate iron(II) porphyrinate derivatives of the type [Fe(Por)(2-MeHIm)].¹⁸ Both imidazole- and imidazolate-ligated iron(II) porphyrinates exhibit an $S = 2$ (quintet) state, but the structural parameters of the coordination groups are distinct with both axial and equatorial bond distance differences and large differences in the displacement of iron from the porphyrin plane. Distinctive features in the Mössbauer spectra obtained in applied magnetic fields show that the doubly occupied d orbital is different in imidazole- vs imidazolate-ligated iron(II) porphyrinates. The positive sign of the quadrupole splitting in the imidazolate derivative shows that the doubly occupied orbital must be the d_{xy} orbital whereas the negative sign in the imidazole derivative is consistent only with a low-symmetry orbital comprised of a hybrid of d_{xz} , d_{yz} , and d_{xy} . This change in the d-electron configuration is clearly consonant with all observed differing features of the two classes.

These differences are similar to those of the globins and the peroxidases. Reduced horse radish peroxidase (HRP) has been studied by Mössbauer spectroscopy and compared with

deoxymyoglobin.^{19,20} These Mössbauer studies, in a strong magnetic field, showed remarkable differences between reduced HRP and deoxymyoglobin (deoxyMb) even though both are five-coordinate hemes with histidine as the axial ligand. Reduced HRP has a positive quadrupole splitting constant ($V_{zz} > 0$) and a rather small asymmetry constant,²⁰ whereas deoxyMb has a large asymmetry constant and a negative value of the quadrupole splitting constant ($V_{zz} < 0$). This strongly argues for different properties of the axial ligand in the five-coordinate iron(II) states of reduced HRP and deoxyMb.

On the basis of these studies, we have begun further investigations with iron(II) porphyrinates of hydrogen-bonded species to fill the gap between imidazole- and imidazolate-ligated porphyrinates. They will also help us understand how hydrogen bonds influence the molecular and electronic structure of the high-spin species. Recently, we presented the first of our investigations of hydrogen bond formation with a coordinated imidazole in a high-spin iron(II) porphyrinate system, [Fe(TPP)(2-MeHIm)]₂·2-MeHIm.²¹ This species has two independent five-coordinate, high-spin iron(II) porphyrinate sites: one with a strong hydrogen bond to the imidazole and the second with a “neutral” imidazole ligand. This complex has been studied by X-ray and neutron diffraction as well as Mössbauer spectroscopy to assess the effects of hydrogen-bonded imidazole as a ligand. These studies show that there is a clear difference in the iron(II) electronic structure between the two iron sites, distinctions that can be attributed to the presence of a hydrogen bond to the coordinated imidazole in the one site. But in this case, the two different porphyrin molecules in the asymmetric unit make it difficult to unambiguously assign their electronic configuration. We have attempted to obtain additional hydrogen-bonded imidazole-ligated iron(II) porphyrinates for characterization. Success requires finding suitable hydrogen-bond acceptors that are either weak ligands or nonligands to avoid coordination competition with the imidazole (2-methylimidazole).

Balch and co-workers^{22,23} have reported that 1,10-phenanthroline (1,10-phen) forms hydrogen bonds with coordinated imidazole in six-coordinate iron(III) porphyrinates.



They showed that the equilibrium in the reaction above was significantly shifted to the right by the addition of 1,10-phenanthroline. Balch et al. investigated this system by visible spectroscopy and NMR.²² In the NMR experiments, diamagnetic cobalt(III) derivatives were substituted for the

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paramagnetic iron(III) center. Observed proton shifts are consistent with the formation of a hydrogen bond between imidazole and 1,10-phenanthroline. No complexes were isolated.

In this paper, we report new, isolated, imidazole-ligated, hydrogen-bonded iron(II) species with 1,10-phenanthroline as the hydrogen-bond acceptor. The species are [Fe(TPP)(2-MeHIm)]·(1,10-phen) (two crystalline forms) and [Fe(OEP)(2-MeHIm)]·(1,10-phen). The effects of the hydrogen bond have been investigated by crystallographic characterization and Mössbauer studies. The consequences of the relatively weak hydrogen bonds formed between coordinated imidazole and 1,10-phenanthroline are found to have a relatively small or no effect on the electronic structure of the high-spin iron(II) species.

Experimental Section

General Information. All reactions and manipulations for the preparation of the iron(II) porphyrin derivatives (see below) were carried out under argon using a double-manifold vacuum line, Schlenkware, and cannula techniques. Chlorobenzene was washed with concentrated sulfuric acid, then with water until the aqueous layer was neutral, dried with MgSO₄, and distilled twice over P₂O₅ under argon. Hexanes were distilled over sodium benzophenone. 2-Methylimidazole (Aldrich) was recrystallized from toluene/methanol and dried under vacuum. All other chemicals were used as received from Aldrich or Fisher. The free-base porphyrin *meso*-tetraphenylporphyrin (H₂TPP) was prepared according to the work of Adler et al.²⁴ Octaethylporphyrin (H₂OEP) was prepared from formaldehyde and 3,4-diethylpyrrole.²⁵ The metalation of the free-base porphyrin to give [Fe(Por)Cl] (Por = TPP, OEP) was done as previously described.²⁶ [Fe(Por)]₂O was prepared according to a modified Fleischer preparation.²⁷

Mössbauer measurements were performed on a constant acceleration spectrometer from 4.2 to 300 K with optional small field and in a 9 T superconducting magnet system (Knox College). Samples for Mössbauer spectroscopy were prepared by immobilization of the crystalline material in Apiezon M grease.

IR spectra were recorded on a Nicolet Nexus 670 FT-IR spectrometer as KBr pellets. Both [Fe(Por)(2-MeHIm)] and [Fe(Por)(2-MeHIm)]·(1,10-phen) crystalline samples were measured.

Synthesis of [Fe(TPP)(2-MeHIm)]·(1,10-phen) (Form A). [Fe(TPP)]₂O (32 mg, 0.024 mmol) was mixed with ethanethiol (1 mL) in chlorobenzene (9 mL) and stirred for three days at room temperature. The resulting solution of the four-coordinate [Fe(II)-(TPP)] was transferred into a Schlenk flask containing 2-methylimidazole (17 mg, 0.21 mmol) and 1,10-phenanthroline (50 mg, 0.28 mmol). The mixture was stirred for 15 min. X-ray quality crystals were obtained in 8 mm × 250 mm sealed glass tubes by liquid diffusion using hexanes as nonsolvent.

Synthesis of [Fe(TPP)(2-MeHIm)]·(1,10-phen) (Form B). A similar procedure as the above was used. [Fe(TPP)]₂O (105 mg,

0.078 mmol), ethanethiol (2.5 mL), chlorobenzene (25 mL), 2-methylimidazole (44 mg, 0.54 mmol), and 1,10-phenanthroline (154 mg, 0.86 mmol) were used.

Synthesis of [Fe(OEP)(2-MeHIm)]·(1,10-phen). A similar procedure as the above was used. [Fe(OEP)]₂O (28 mg, 0.023 mmol), ethanethiol (1 mL), chlorobenzene (7 mL), 2-methylimidazole (18 mg, 0.22 mmol), and 1,10-phenanthroline (46 mg, 0.26 mmol) were used.

UV–Vis Spectroscopy. UV–vis spectra were recorded on a Perkin-Elmer Lambda 19 UV/vis/near-IR spectrometer and were obtained in a specially designed combined 1- and 10-mm inert atmosphere cell. A solution of four-coordinate [Fe(II)(TPP)] (0.5×10^{-3} mol/L) was prepared as the above. UV–vis spectra of iron porphyrinates at different concentration of 2-methylimidazole and 1,10-phenanthroline were measured. The concentrations of 2-methylimidazole for UV–vis measurements range from 0.11 to 1.07 mol/L, the concentrations of 1,10-phenanthroline range from 4.8×10^{-3} to 8.6×10^{-1} mol/L.

X-ray Structure Determinations. Single-crystal experiments were carried out on a Bruker Apex system with graphite monochromated Mo–K α radiation ($\lambda = 0.71073$ Å). The structures were solved by direct methods and refined against F^2 using SHELXTL,^{28,29} subsequent difference Fourier syntheses led to the location of the remaining non-hydrogen atoms. All non-hydrogen atoms were refined anisotropically if not remarked otherwise below. Hydrogen atoms were added with the standard SHELXL-97 idealization methods. The program SADABS³⁰ was applied for the absorption correction. Brief crystal data and intensity collection parameters for the crystalline complexes are shown in Table 1. Complete crystallographic details, atomic coordinates, anisotropic thermal parameters, and fixed hydrogen atom coordinates are given in the Supporting Information.

[Fe(TPP)(2-MeHIm)]·(1,10-phen) (Form A). A red crystal with the dimensions $0.49 \times 0.43 \times 0.14$ mm³ was used for the structure determination. Crystal data were collected at room temperature as the crystals cracked on cooling. The structure was refined in space group $P2_1/c$. The asymmetric unit contains one porphyrinate molecule, one 1,10-phenanthroline, and one chlorobenzene solvate. The C₆H₅Cl was badly disordered, showing rotation around an axis perpendicular to the molecular plane and passing through the ring center. No models could satisfy this disorder. SQUEEZE³¹ was used to model this disordered chlorobenzene. A 1030 Å^3 void volume and electron density equivalent to 233.1 e^- are consistent with four C₆H₅Cl in the unit cell, resulting in a 1:1 ratio of solvent to porphyrin.

[Fe(TPP)(2-MeHIm)]·(1,10-phen) (Form B). A red crystal with the dimensions $0.34 \times 0.21 \times 0.19$ mm³ was used for the structure determination. Crystal data were collected at 100 K. The structure was refined in space group $P2_1/c$. The asymmetric unit contains two porphyrinate molecules, two 1,10-phenanthrolines, and two chlorobenzene solvates. For both porphyrin molecules, two phenyl rings are disordered, each over two positions with 50% occupancy.

An attempted refinement in a small unit cell containing just one porphyrin required addition disorder in the porphyrin and resulted in a less satisfactory refinement. To verify the correct choice of

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Table 1. Brief Crystallographic Data and Data Collection Parameters

	[Fe(TPP)(2-MeHIm)]·(1,10-phen) (Form A)	[Fe(TPP)(2-MeHIm)]·(1,10-phen) (Form B)	[Fe(OEP)(2-MeHIm)]·(1,10-phen)
formula	C ₄₈ H ₃₄ FeN ₆ ·C ₁₂ H ₈ N ₂ ·C ₆ H ₅ Cl	2C ₄₈ H ₃₄ FeN ₆ ·2C ₁₂ H ₈ N ₂ ·2C ₆ H ₅ Cl	2C ₄₀ H ₅₀ FeN ₆ ·2C ₁₂ H ₈ N ₂
FW	1043.42	2086.83	1701.83
<i>a</i> , Å	11.8059(10)	24.6264(9)	25.319(2)
<i>b</i> , Å	21.8824(16)	18.7689(7)	19.1907(14)
<i>c</i> , Å	21.3200(17)	23.2011(8)	18.3944(13)
β, deg	100.143(4)	106.726(2)	
<i>V</i> , Å ³	5421.8(7)	10270.1(6)	8937.6(12)
<i>Z</i>	4	4	4
space group	<i>P</i> 2 ₁ / <i>c</i>	<i>P</i> 2 ₁ / <i>c</i>	<i>Pca</i> 2 ₁
<i>D_c</i> , g/cm ³	1.278	1.350	1.265
<i>F</i> (000)	2168	4336	3616
μ, mm ⁻¹	0.377	0.398	0.383
crystal dimens, mm	0.49 × 0.43 × 0.14	0.34 × 0.21 × 0.19	0.44 × 0.28 × 0.20
absorption correction		SADABS	
radiation, MoKα, λ		0.71073 Å	
<i>T</i> , K	293(2)	100(2)	100(2)
total data collected	59973	130581	181135
unique data	14107 (<i>R</i> _{int} = 0.030)	23997 (<i>R</i> _{int} = 0.032)	24415 (<i>R</i> _{int} = 0.046)
unique obsd data [<i>I</i> > 2σ(<i>I</i>)]	8398	20261	21117
refinement method		on <i>F</i> ² (SHELXL)	
final <i>R</i> indices [<i>I</i> > 2σ(<i>I</i>)]	<i>R</i> ₁ = 0.0428, <i>wR</i> ₂ = 0.1101	<i>R</i> ₁ = 0.0678, <i>wR</i> ₂ = 0.1521	<i>R</i> ₁ = 0.0499, <i>wR</i> ₂ = 0.1228
final <i>R</i> indices [for all data]	<i>R</i> ₁ = 0.0765, <i>wR</i> ₂ = 0.1195	<i>R</i> ₁ = 0.0806, <i>wR</i> ₂ = 0.1587	<i>R</i> ₁ = 0.0605, <i>wR</i> ₂ = 0.1292

cell, the two porphyrins were overlaid, one on the top of the other. This indicated differences in the conformations of the phenyl ring and β carbons, confirming the cell choice.

[Fe(OEP)(2-MeHIm)]·(1,10-phen). A red crystal with the dimensions 0.44 × 0.28 × 0.20 mm³ was used for the structure determination. Crystal data were collected at 100 K. The structure was refined in space group *Pca*2₁. The asymmetric unit contains two porphyrinate molecules and two 1,10-phenanthroline molecules. The methyl carbon atom of one ethyl group in the second porphyrin molecule was found to be disordered over two positions, a major and a minor position (C(22a) and C(22b)). Their distances to C(221) were constrained to be 1.52 Å, and their thermal displacement parameters were constrained to be equal to each other. After the final refinement, the occupancy of the major orientation was found to be 75%.

PLATON³¹ suggested a possible higher symmetry space group, *Pbcn*, but several peripheral ethyl groups do not fit in this symmetry. The refinement as disordered parts under *Pbcn* gave unsatisfactorily high *R*₁/*wR*₂ values. Thus, the space group *Pca*2₁ was used in the final refinement, which gave *R*₁ = 0.0499, *wR*₂ = 0.1228.

Results

New five-coordinate iron(II) porphyrin complexes with a hydrogen-bonded 2-methylimidazole as the fifth ligand have been synthesized by reaction with 2-methylimidazole in the presence of excess 1,10-phenanthroline. Three crystalline derivatives of two porphyrins have been obtained. Two of them are polymorphs of [Fe(TPP)(2-MeHIm)]·(1,10-phen) and are called forms **A** and **B**; the third is an OEP derivative, [Fe(OEP)(2-MeHIm)]·(1,10-phen). Two of these three crystalline species have two independent porphyrinate molecules in the asymmetric unit, called molecules **1** and **2**. In the figures and tables, the following atom naming convention has been used: *Q*(*n*), *Q*(*ny*), and *Q*(*nyy*), where *Q* is the atom type, *n* refers to molecule 1 or 2, and *y* or *yy* are further numbers and letters needed to completely specify the atom.

Thus similar atoms in the two molecules except phenyl rings in [Fe(TPP)(2-MeHIm)]·(1,10-phen) (form **B**) have the same name except for the digit *n*.

In all three crystalline derivatives, there are both porphyrin molecules and 1,10-phenanthroline molecules. The porphyrin molecules are all five-coordinate imidazole-ligated iron(II) species and a 1,10-phenanthroline molecule is hydrogen bonded to the coordinated 2-methylimidazole. As an example, the ORTEP diagram of [Fe(TPP)(2-MeHIm)]·(1,10-phen) (form **B**, molecule **1**) is illustrated in Figure 1. ORTEP diagrams of the four additional structures and a complete listing of bond distances and angles are given in the Supporting Information.

The average value of the Fe–N_p bond distances are 2.084(5) Å for [Fe(TPP)(2-MeHIm)]·(1,10-phen) (**A**), 2.080(9), 2.081(6) Å for [Fe(TPP)(2-MeHIm)]·(1,10-phen) (**B**), and 2.088(14), 2.087(14) Å for [Fe(OEP)(2-MeHIm)]·(1,10-phen). The Fe–N_{im} bond length is 2.1289(13) Å for [Fe(TPP)(2-MeHIm)]·(1,10-phen) (**A**), 2.125(3), 2.120(3) Å for [Fe(TPP)(2-MeHIm)]·(1,10-phen) (**B**), and 2.135(2), 2.131(2) Å for [Fe(OEP)(2-MeHIm)]·(1,10-phen), respectively.

Two crystalline species of TPP and OEP derivatives were studied with variable-temperature Mössbauer spectroscopy. For the TPP derivative, the Mössbauer sample was made by the method for [Fe(TPP)(2-MeHIm)]·(1,10-phen) (form **A**). These Mössbauer parameters at various temperatures are given in Table 2.

Discussion

Our previous studies have shown that strong hydrogen bonding²¹ to or deprotonation¹⁵ of the coordinated imidazole in five-coordinate iron(II) porphyrinate species significantly influences both the structural and electronic properties of the resulting species. The structural consequences of these effects are schematically illustrated in Figure 2. Limiting behavior

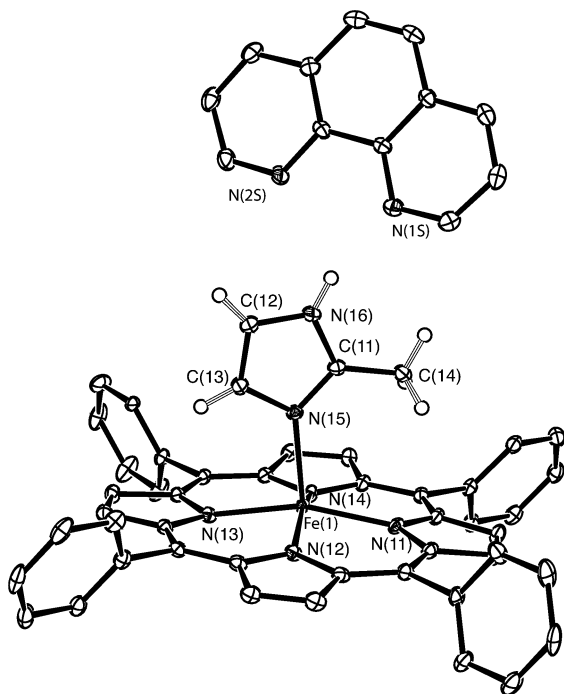


Figure 1. ORTEP diagram of $[\text{Fe}(\text{TPP})(2\text{-MeHIm})]\cdot(1,10\text{-phen})$ (form B molecule 1). For clarity, one orientation of the disordered phenyl rings and the hydrogen atoms of the porphyrin ligand have been omitted. 50% probability ellipsoids are depicted.

Table 2. Variable-Temperature Mössbauer Parameters (mm/s) for $[\text{Fe}(\text{TPP})(2\text{-MeHIm})]\cdot(1,10\text{-phen})$ and $[\text{Fe}(\text{OEP})(2\text{-MeHIm})]\cdot(1,10\text{-phen})$

[Fe(TPP)(2-MeHIm)]·(1,10-phen)					
T (K)	298	200	100	20	
ΔE_Q	1.46	1.65	2.06	2.12	
δ_{Fe}	0.80	0.86	0.88	0.90	
[Fe(OEP)(2-MeHIm)]·(1,10-phen)					
T (K)	295	200	100	25	4.2
ΔE_Q	1.73	1.80	1.92	2.01	1.96
δ_{Fe}	0.82	0.89	0.89	0.90	0.94

is shown by the averaged values displayed by a number of imidazole complexes depicted at the left side of the figure. The other limiting case, deprotonation of the N–H of the coordinated imidazole to yield coordinated imidazolate, is shown at the right. Are these differences meaningful? We believe that the differences in geometry around the iron atom in the two cases reflect a difference in the electronic structure of iron. Experimental study of the electronic structure of iron in these systems is best investigated with Mössbauer spectroscopy. In addition to significant differences in the magnitude of the quadrupole splitting constants, the sign of the quadrupole splitting is an important distinction between imidazole- and imidazolate-ligated iron(II) species. The imidazolate derivatives have both larger values of the quadrupole splitting and a positive value for the sign. This distinction in sign of the quadrupole splitting arises from differing symmetry of the doubly occupied orbital, which in the imidazolate species has essentially pure d_{xy} character, i.e., the doubly occupied orbital is in the plane of the heme. This appears to be the usual electronic state for high-spin iron(II), whereas the imidazole-ligated species, including deoxyhemoglobin and deoxymyoglobin, have a very low symmetry orbital comprised of a hybrid d_{xy} , d_{xz} , and d_{yz} set

in which the major direction of the electron density is oblique to the heme plane and all Fe–N bond directions.

The coordination geometry of the hydrogen-bonded species shown in the center of Figure 2 suggests that the hydrogen bond has a real effect on the structure. The structural parameters suggest that the hydrogen bond leads to an “imidazole-like” coordinated ligand. The Mössbauer analysis of this system is unfortunately complicated by the presence of a second iron center in the solid-state sample. The assignment of a positive sign for the quadrupole doublet, similar to that of the imidazolate derivatives was judged most likely, but an absolute, unambiguous assessment of the sign was not possible.

In order to further investigate this question, we sought to prepare additional imidazole-ligated iron(II) porphyrinates in which the imidazole was hydrogen bonded to an external acceptor. Suitable hydrogen-bond acceptors in such a system must either be weak ligands or a nonligand so as to avoid coordination competition with 2-methylimidazole. Attempts using species including 2-methylpyridine, 2-chloropyridine, proton sponge, and DBU failed because they failed to give crystalline materials,³² failed to yield a hydrogen-bonded species with the coordinated imidazole or, most interestingly, yielded the already known species with a 2-methylimidazole solvate hydrogen bonded to the coordinated imidazole.²¹ One hydrogen-bonding acceptor that did yield interesting new materials was 1,10-phenanthroline.

1,10-Phenanthroline has been shown as a hydrogen-bond acceptor to coordinated imidazole.²² By addition of excess 1,10-phenanthroline, our experiments gave new species, $[\text{Fe}(\text{TPP})(2\text{-MeHIm})]\cdot(1,10\text{-phen})$ and $[\text{Fe}(\text{OEP})(2\text{-MeHIm})]\cdot(1,10\text{-phen})$, in which 1,10-phenanthroline is H bonded to the coordinated imidazole. These species have been characterized by X-ray crystallography, Mössbauer spectroscopy, and IR and UV–vis spectroscopic measurements.

IR Spectroscopy. IR has been used to demonstrate the formation of a hydrogen bond to coordinated imidazole in porphyrins³³ or other species.³⁴ In our case, in order to verify the formation of a hydrogen-bonded species, IR spectra of $[\text{Fe}(\text{Por})(2\text{-MeHIm})]$ and $[\text{Fe}(\text{Por})(2\text{-MeHIm})]\cdot(1,10\text{-phen})$ have been measured.

The spectra of $[\text{Fe}(\text{OEP})(2\text{-MeHIm})]$ and $[\text{Fe}(\text{OEP})(2\text{-MeHIm})]\cdot(1,10\text{-phen})$ are shown in Figure 3. The sharp band at 3360 cm^{-1} for $[\text{Fe}(\text{OEP})(2\text{-MeHIm})]$ was assigned as the N–H stretch. But in $[\text{Fe}(\text{OEP})(2\text{-MeHIm})]\cdot(1,10\text{-phen})$, it has shifted, and two new broad bands arise at lower frequencies. One is at 3050 cm^{-1} which is the same position as the C–H stretch in 1,10-phenanthroline. The other band is at 3100 cm^{-1} which can reasonably be assigned the hydrogen-bonded N–H stretch. The spectra of $[\text{Fe}(\text{TPP})(2\text{-MeHIm})]$ and $[\text{Fe}(\text{TPP})(2\text{-MeHIm})]\cdot(1,10\text{-phen})$ showed similar behavior. Such a change indicates the formation of

(32) Many of these failed crystallization experiments yielded oils suggesting the presence of several species.

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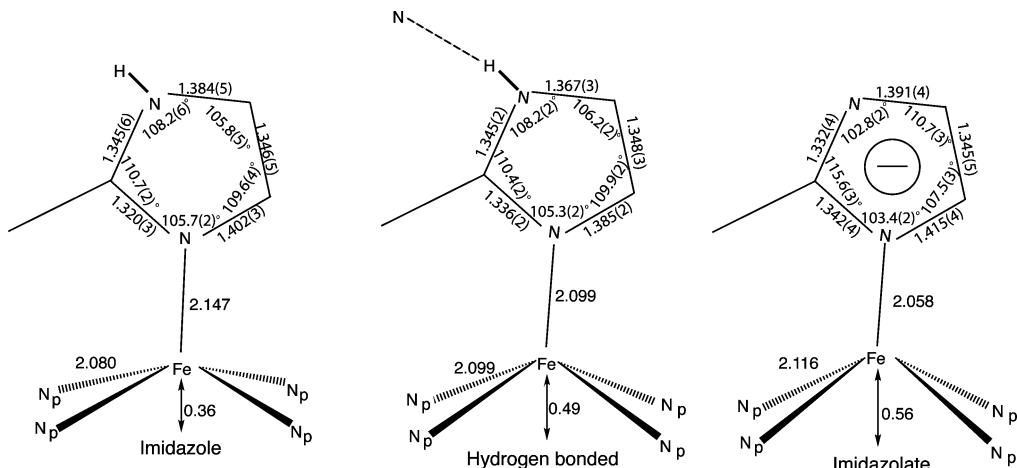


Figure 2. Schematic diagrams of the coordination group geometries found for a series of imidazole-ligated high-spin iron(II) porphyrinates (left), an analogous species with a strong hydrogen bond to the coordinated imidazole (center), and imidazolate-ligated high-spin iron(II) porphyrinates (right).

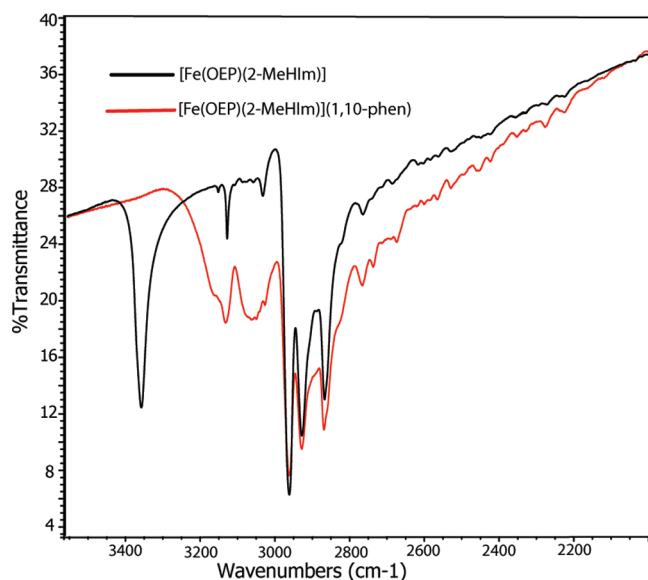


Figure 3. IR spectra of $[\text{Fe}(\text{OEP})(2\text{-MeHIm})] \cdot (1,10\text{-phen})$ and $[\text{Fe}(\text{OEP})(2\text{-MeHIm})]$ between 3500 and 2000 cm^{-1} .

a hydrogen bond in new species, which is similar to that for an imidazole-coordinated manganese complex.³⁴

Structural Studies. All new porphyrin molecules are five-coordinate 2-methylimidazole-ligated species. 1,10-Phenanthroline does not coordinate to the iron center in the solid state, presumably as the result of the steric configuration between 1,10-phenanthroline and the porphyrin ring. But, as expected, 1,10-phenanthroline has formed hydrogen bonds with the coordinated imidazole in a 1:1 ratio of 2-methylimidazole and 1,10-phenanthroline.

Two views of the hydrogen-bonding pattern are shown in Figure 4. The hydrogen bonds between 2-methylimidazole and 1,10-phenanthroline are three-center hydrogen bonds: the hydrogen of the imidazole nitrogen is hydrogen bonded to both nitrogen atoms of 1,10-phenanthroline; two hydrogen bonds are formed. Individual distances and angles for the five independent molecules are listed in Table 3. In all structures, the two hydrogen bonds are not equal: one is stronger with a shorter $\text{N} \cdots \text{N}$ distance (average 2.97(2) versus 3.08(3) Å), shorter intermolecular $\text{H} \cdots \text{N}$ distance (average 2.13(4) versus 2.52(5) Å), and larger $\text{N}-\text{H} \cdots \text{N}$

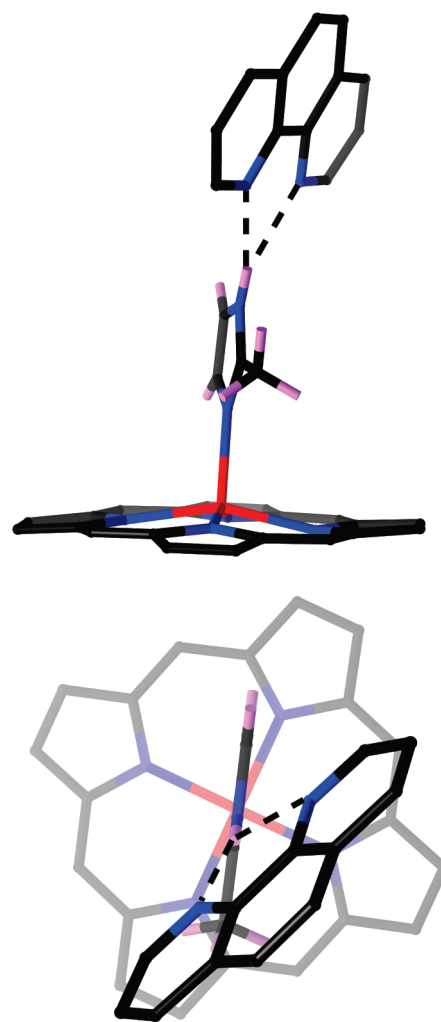


Figure 4. Two views of the hydrogen-bonding interaction between the coordinated imidazole and 1,10-phenanthroline. For clarity, only the 24 atoms of the porphyrin core are shown in the figure.

angle (average 162(6) versus 122(2)°). It should be noted that all values involving the hydrogen atom of the imidazole are based on an idealized position of this hydrogen atom. The dihedral angle between the 2-methylimidazole plane and the corresponding 1,10-phenanthroline plane ranges from 37.6 to 46.5°.

Table 3. Hydrogen Bond Parameters Between Imidazole and 1,10-Phenanthroline

complex	N _{Im} –H distance ^a	N···H distance ^a	N _{Im} ···N _{phen} distance ^a	N _{Im} –H···N _{phen} angle ^b	dihedral angle ^{b,c}
[Fe(TPP)(2-MeHIm)]·(1,10-phen) (form A)	0.86	2.14	2.980(2)	165.0	43.7
[Fe(TPP)(2-MeHIm)]·(1,10-phen) (form B, mol 1)	0.86	2.48	3.032(2)	123.0	
	0.88	2.09	2.949(4)	164.1	40.8
	0.88	2.52	3.089(4)	122.7	
[Fe(TPP)(2-MeHIm)]·(1,10-phen) (form B, mol 2)	0.88	2.20	3.006(4)	151.8	37.6
	0.88	2.48	3.059(4)	123.9	
[Fe(OEP)(2-MeHIm)]·(1,10-phen) (mol 1)	0.88	2.10	2.954(3)	163.7	46.5
	0.88	2.57	3.104(3)	119.6	
[Fe(OEP)(2-MeHIm)]·(1,10-phen) (mol 2)	0.88	2.12	2.979(3)	165.2	42.4
	0.88	2.57	3.113(3)	121.1	

^a In angstroms. ^b Value in degrees. ^c Dihedral angle between the imidazole plane and the corresponding phenanthroline plane.

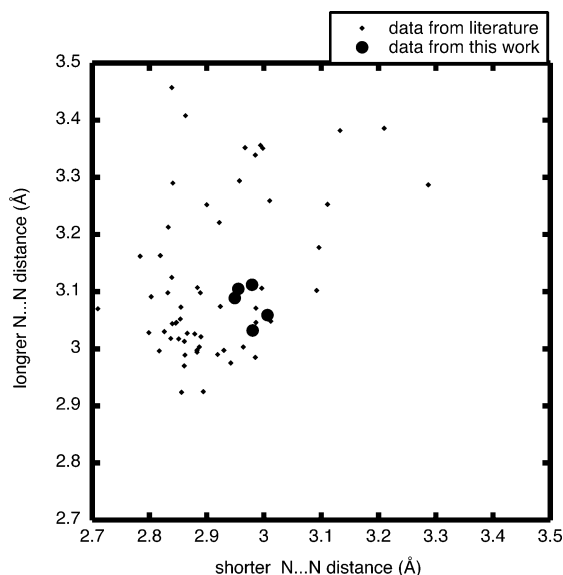


Figure 5. Results from a Cambridge Structural Database survey on the hydrogen bonds formed by 1,10-phenanthroline to a nitrogen atom hydrogen-bond donor in a range of differing crystal structures, showing the correlations between the two intermolecular N···N distances associated with the hydrogen bonds. The shorter distance is given on the x-axis, and the longer one, on the y-axis. The large black circles are the results for our current samples.

Is there anything unusual about the hydrogen bonds between the imidazole and the 1,10-phenanthroline? How to evaluate the strength of these hydrogen bonds? One useful comparison is to utilize the Cambridge Structural Database³⁵ to assess the hydrogen bonds formed by 1,10-phenanthroline in a range of crystal structures. We found a total of 60 instances of three-center hydrogen bonds formed by 1,10-phenanthroline to a nitrogen atom hydrogen-bond donor in a solid state structure. Since all of these are X-ray diffraction studies, the hydrogen atom positions are relatively uncertain. Thus, the best distances to compare are the intermolecular N···N distances. The correlations between these two N···N distances associated with the hydrogen bonds are shown in Figure 5; the shorter N···N distance is shown on the x-axis, and the longer one is the y-axis. The big black circles are the results for our current samples. The figure shows that the N···N distances in these new species are well within the range of hydrogen bonds associated with 1,10-phenan-

throline in other species and the observed asymmetry is typical.

Are there structural changes caused by these hydrogen bonds? The overall structural features are those expected for a high-spin iron(II) complex.^{14,36} These include large equatorial Fe–N_p bond distances, a significant out-of-plane displacement of the iron(II) atom, and a radial expansion of the core (an increase in the size of the central hole).³⁷ The average Fe–N_p bond distances range from 2.080(9) to 2.088(14) Å and the Fe–N_{Im} bond lengths range from 2.120(3) to 2.135(2) Å; both are similar to the values for other imidazole-ligated porphyrinates as shown in Table 4. As shown in Table 4, the out-of-plane displacement of the iron atom out of the mean 24-atom porphyrin core (Δ) and the plane defined by the four pyrrole nitrogen atoms (ΔN_4) also fit the range for those imidazole-ligated porphyrinates. The displacements range from 0.38 to 0.49 Å out of the 24-atom plane and from 0.36 to 0.39 Å out of the four pyrrole nitrogen atom plane. The radii of the porphyrin cores, given by Ct···N in Table 4, are nearly identical at 2.049 and 2.051 Å.

The steric bulk of the imidazole 2-methyl group leads to, in all derivatives examined to date, an off-axis tilt of the axial Fe–N_{Im} bond and a rotation of the imidazole ligand that leads to unequal Fe–N_{Im}–C_{Im} angles. The tilt angles range from 2.1° for [Fe(OEP)(2-MeHIm)]·(1,10-phen) (mol 2) to 8.8° for [Fe(TPP)(2-MeHIm)]·(1,10-phen) (A). This tilting is the partial result of minimizing the interaction between the bulky imidazole methyl group and the porphyrin core. Two important angles associated with the imidazole ligands (Fe–N_{Im}–C_{Im}) are given in Table 4. They range from 130.8(2) to 133.7(2)° on the methyl side and 121.2(16) to 123.8(2)° on the other side. The off-axis tilt and imidazole rotation are correlated so as to maximize the distance between the 2-methyl group and porphyrin core atoms, the values are similar to those for other imidazole-ligated species.

The above structural data suggest that these new species are essentially identical to those of other imidazole-ligated species and indicates that the imidazole···1,10-phen hydrogen bonds do not have an obvious influence on the structure at iron. This is quite distinct from the strong hydrogen bond in [Fe(TPP)(2-MeHIm)]₂·2-MeHIm,²¹ which has remarkable

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Table 4. Selected Bond Distances (Å) and Angles (deg) for New Structures and Related Species^a

complex ^b	Fe–N _p ^{c,d}	Fe–N _{im} ^d	ΔN ₄ ^{d,e}	Δ ^{d,f}	Ct···N ^d	Fe–N–C ^{g,h}	Fe–N–C ^{g,i}	θ ^{g,j}	φ ^{g,k}	ref
[Fe(TPP)(2-MeHIm)]·(1,10-phen) (A)	2.084(5)	2.1289(13)	0.38	0.47	2.049	132.03(12)	123.09(11)	8.8	18.5	tw
[Fe(TPP)(2-MeHIm)]·(1,10-phen) (B)	2.080(9)	2.125(3)	0.36	0.39	2.049	130.8(2)	123.8(2)	6.6	17.7	tw
	2.081(6)	2.120(3)	0.36	0.38	2.051	133.7(2)	121.5(2)	5.4	7.8	tw
[Fe(OEP)(2-MeHIm)]·(1,10-phen) (A)	2.088(14)	2.135(2)	0.39	0.49	2.051	132.75(17)	121.20(16)	2.7	8.1	tw
	2.087(14)	2.131(2)	0.39	0.47	2.051	133.43(17)	121.66(17)	2.1	13.8	tw
[Fe(OEP)(1,2-Me ₂ Im)]	2.080(6)	2.171(3)	0.37	0.45	2.047	132.7(3)	121.4(2)	3.8	10.5	16
[Fe(OEP)(2-MeHIm)]	2.077(7)	2.135(3)	0.34	0.46	2.049	131.3(3)	122.4(3)	6.9	19.5	16
[Fe(TPP)(1,2-Me ₂ Im)]	2.079(8)	2.158(2) ^l	0.36	0.42	2.048	129.3(2)	124.9(2)	11.4	20.9	14
[Fe(TTP)(2-MeHIm)]	2.076(3)	2.144(1)	0.32	0.39	2.050	132.8(1)	121.4(1)	6.6	35.8	14
[Fe(Tp–OCH ₃ PP)(2-MeHIm)]	2.087(7)	2.155(2) ^l	0.39	0.51	2.049	130.4(2)	123.4(2)	8.6	44.5	14
[Fe(Tp–OCH ₃ PP)(1,2-Me ₂ Im)]	2.077(6)	2.137(4)	0.35	0.38	2.046	131.9(3)	122.7(3)	6.1	20.7	14
[Fe(TPP)(2-MeHIm)](2-fold)	2.086(8)	2.161(5)	0.42	0.55	2.044	131.4(4)	122.6(4)	10.3	6.5	38
[Fe(TPP)(2-MeHIm)]·1.5C ₆ H ₅ Cl	2.073(9)	2.127(3) ^l	0.32	0.38	2.049	131.1(2)	122.9(2)	8.3	24.0	13
average of the eight	2.080(5)	2.147(16)	0.36(3)	0.44(6)	2.048(2)	131.4(12)	122.7(11)	7.8(24)	22.9	
[Fe(TPP)(2-MeHIm)] (mol 1)	2.080(8)	2.120(2)	0.36	0.41	2.050	131.6(1)	122.4(1)	9.2	16.0	21
[Fe(TPP)(2-MeHIm)] (mol 2)	2.099(7)	2.099(2)	0.49	0.55	2.040	129.0(1)	125.7(1)	7.6	22.9	21
[K(222)][Fe(OEP)(2-MeIm [−])]	2.113(4)	2.060(2)	0.56	0.65	2.036	136.6(2)	120.0(2)	3.6	37.4	15
[K(222)][Fe(OEP)(2-MeIm [−])]	2.118(13)	1.999(5)	0.56	0.66	2.044	129.6(3)	126.7(3)	9.8	23.4	15
		2.114(5)				133.6(4)	121.9(4)	6.5	21.6	
[Fe(TpivPP)(2-MeIm [−])] [−]	2.11(2)	2.002(15)	0.52	0.65	2.045	NR ^m	NR	5.1	14.7	39
average of the three	2.114(4)	2.044(54)	0.55(2)	0.65(1)	2.042(5)					

^a Estimated standard deviations are given in parentheses. ^b All complexes are high spin. ^c Averaged value. ^d In angstroms. ^e Displacement of iron from the mean plane of the four pyrrole nitrogen atoms. ^f Displacement of iron from the 24-atom mean plane of the porphyrin core. ^g Value in degrees. ^h 2-Carbon, methyl substituted. ⁱ Imidazole 4-carbon. ^j Off-axis tilt (deg) of the Fe–N_{im} bond from the normal to the porphyrin plane. ^k Dihedral angle between the plane defined by the closest N_p–Fe–N_{im} and the imidazole plane in degrees. ^l Major imidazole orientation. ^m Not reported.

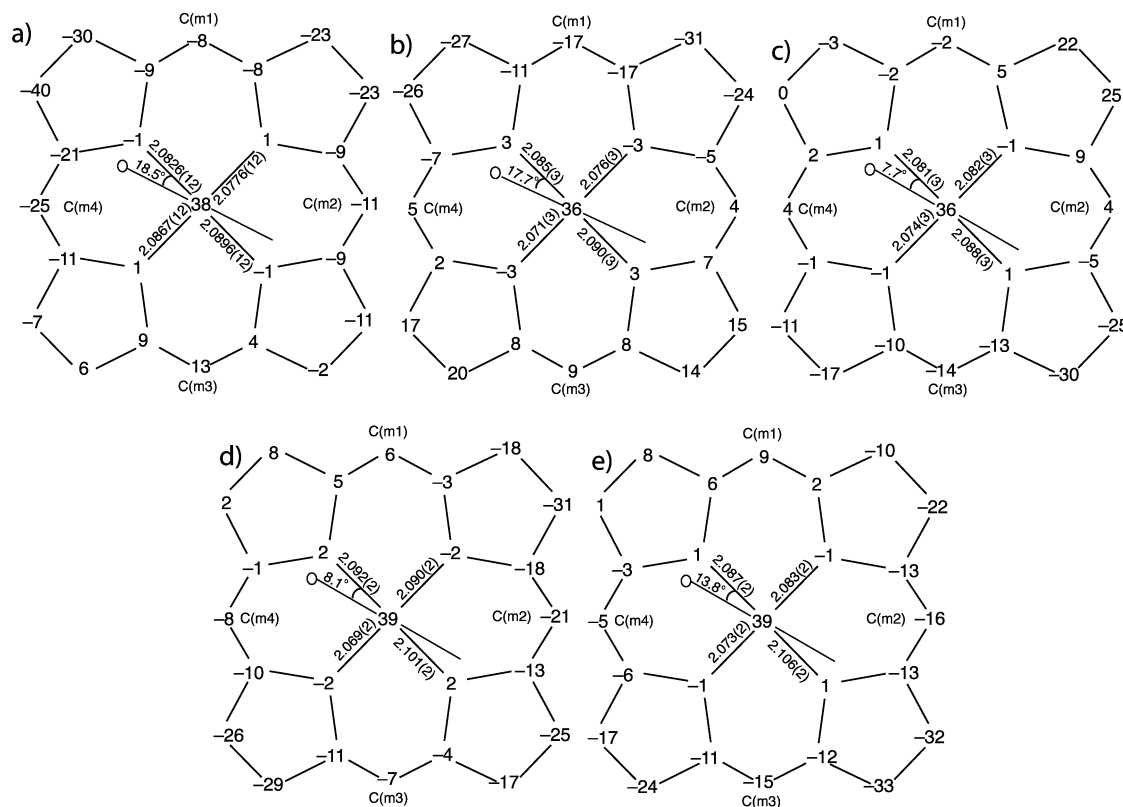


Figure 6. Formal diagrams of the porphyrinato cores of (a) [Fe(TPP)(2-MeHIm)]·(1,10-phen) (A); (b) [Fe(TPP)(2-MeHIm)]·(1,10-phen) (B, molecule 1); (c) [Fe(TPP)(2-MeHIm)]·(1,10-phen) (B, molecule 2); (d) [Fe(OEP)(2-MeHIm)]·(1,10-phen) molecule 1; (e) [Fe(OEP)(2-MeHIm)]·(1,10-phen) molecule 2. Illustrated are the displacements of each atom from the mean plane of the four pyrrole nitrogen in units of 0.01 Å. Positive values of displacement are toward the imidazole ligand. The diagrams also show the orientation of the imidazole ligand with respect to the atoms of the porphyrin core. The location of the methyl group at the 2-carbon position is represented by the circle.

effects on the structure at iron, including much larger Fe–N_p distances (difference of ~0.02 Å) and iron displacement (difference of ~0.10 Å). These data allow us to tentatively conclude that there is a gradation of effects from hydrogen bonding on the structure of the high-spin, imidazole-ligated iron porphyrinates.

Figure 6 gives formal diagrams of the porphyrin cores of the new iron(II) structures. Given are the displacements of each atom from the mean plane of the four pyrrole nitrogen in units of 0.01 Å. An analogous diagram showing atomic displacements from the mean plane of the 24-atom core is given in the Supporting Information (Figure S5). The

orientation of the imidazole ligand with respect to the core atoms are shown by the line with the circle representing the methyl group bound at the 2-carbon atom position. The dihedral angles (ϕ) between the imidazole ligand and the plane defined by N(n1), Fe(n), and N(n5) are all less than 20° . Also included on the diagrams of Figure 6 are the individual Fe–N_p bond lengths. The pair of Fe–N_p distances closer to imidazole plane are longer than the other pair (average 2.090(8) versus 2.078(7) Å), which could be caused by a repulsive interaction between imidazole group and the close N(1) and N(3) atoms when the dihedral angle is small.

As can be seen in Figure 6, the porphyrin cores have somewhat different conformations even within the same crystal. This is consistent with our previous observation that there is not a *single* preferred core conformation for imidazole-ligated high-spin iron(II) complexes.^{14,16} Core conformations have also been analyzed by the normal structural decomposition (NSD) method provided by Sheltnutt et al.⁴⁰ Table S1 of the Supporting Information lists the NSD out-of-plane displacements of known high-spin iron(II) porphyrinates ligated with a 2-methylimidazole or 1,2-dimethylimidazole axial ligand. This further confirms that there was no common pattern for the core conformations.

In addition to the hydrogen bonds, there are remarkable π – π stacking interactions associated with 1,10-phenanthroline. For [Fe(TPP)(2-MeHIm)]·(1,10-phen) (A), as shown in Figure 7a, one aromatic ring of a symmetry related phenanthroline (symmetry operator #1: $x, 1.5 - y, -0.5 + z$) is overlapped with a six-membered chelating ring in porphyrin (Fe(1), N(1), N(2), C(a2), C(m1), C(a3)) and the dihedral angle between them is 15.9° . The shortest intermolecular distance is Fe(1)···C(9S#1) = 3.261 Å. For [Fe(OEP)(2-MeHIm)]·(1,10-phen), phenanthroline hydrogen bonded to molecule 1 stacks over another phenanthroline hydrogen bonded to molecule 2 from a symmetry related unit (symmetry operator #2: $1.5 - x, y, -0.5 + z$). They are almost parallel to each other with the small dihedral angle of 0.7° . The closest intermolecular distance is C(7S)···C(23S#2) = 3.479 Å.

For [Fe(TPP)(2-MeHIm)]·(1,10-phen) (B), there are two kinds of π – π interactions. One is like that for [Fe(OEP)(2-MeHIm)]·(1,10-phen), a phenanthroline hydrogen bonded to molecule 1 stacks over another phenanthroline hydrogen bonded to molecule 2 from a symmetry related unit (symmetry operator #3: $x, 1.5 - y, 0.5 + z$). They are almost parallel to each other with a small dihedral angle of 2.6° . The closest intermolecular distance is C(1S)···C(17S#3) = 3.516 Å. Another π – π interaction is between phenanthroline and one pyrrole ring of porphyrin as shown in Figure 7c. One pyrrole ring in molecule 1 (consisting of N(11), C(1a1), C(1a2), C(1b1), C(1b2)) is overlapped with a phenyl

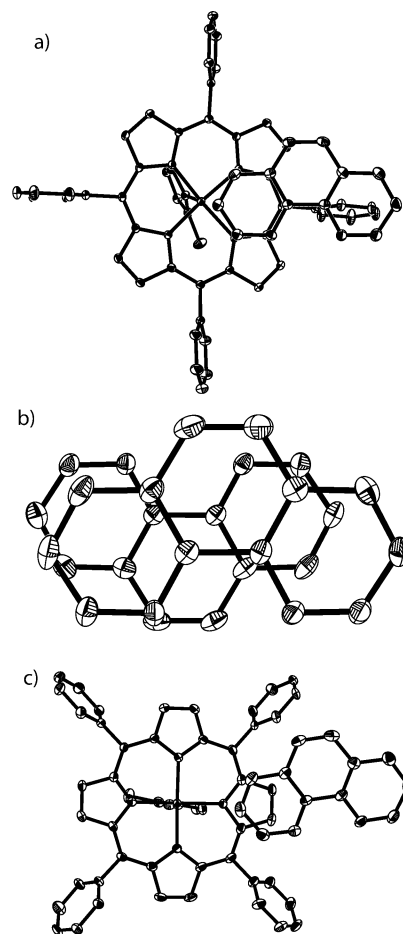


Figure 7. Stacking diagrams showing the π – π interactions. (a) In [Fe(TPP)(2-MeHIm)]·(1,10-phen) (form A), all atoms are contoured at the 10% probability level for clarity; (b) in [Fe(OEP)(2-MeHIm)]·(1,10-phen), all atoms are contoured at the 50% probability level; (c) in [Fe(TPP)(2-MeHIm)]·(1,10-phen) (form B), all atoms are contoured at the 50% probability level.

ring (consisting of N(3S), C(19S), C(20S), C(21S), C(22S), C(23S)) in a symmetry related phenanthroline (symmetry operator #4: $x, y - 1, z$). The dihedral angle between them is 18.1° . The closest intermolecular distance is C(1a1)···C(22S#4) = 3.311 Å. Another pyrrole ring in molecule 2 (consisting of N(23), C(2a5), C(2a6), C(2b5), C(2b6)) is overlapped with a phenyl ring of phenanthroline (consisting of N(1S), C(1S), C(2S), C(3S), C(4S), C(12S)). The corresponding dihedral angle is 10.3° , and the closest intermolecular distance is C(2b5)···C(1S) = 3.322 Å. Obviously, these π – π interactions are important factors to stabilize the solid-state structures.

Electronic Structure. The X-ray structures show that the new phenanthroline species are strongly similar to the imidazole-ligated species. Our previous studies suggest that imidazole- and imidazolate-ligated five-coordinate iron(II) porphyrinates form two distinct groups with different electronic configurations.^{14,15} Mössbauer spectra of the imidazolate-ligated species show a large positive value of the quadrupole splitting with a small asymmetry parameter (η), consistent with a ground-state doubly occupied d_{xy} orbital, $(d_{yz})^2(d_{xz})^1(d_{yz})^1(d_{z^2})^1(d_{x^2-y^2})^1$, i.e., an orbital in the heme plane. On the other hand, Mössbauer spectra of the imidazole-

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- (40) Sun, L.; Sheltnutt, J. A. Program NSD is available via the internet at <http://jasheln.unm.edu> (accessed April 18, 2008).

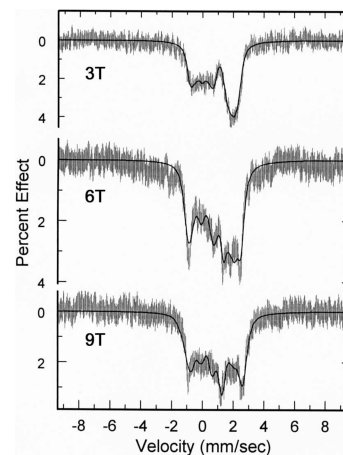
Table 5. Solid-State Mössbauer Parameters for Five-Coordinate, High-Spin, Imidazole-Ligated Iron(II) Porphyrinates and Related Species

complex	ΔE_q^a	δ_{Fe}^a	η^b	Γ^c	T, K	ref
[Fe(OEP)(2-MeHIm)]•(1,10-phen)	-1.93	0.94	0.76	0.43	4.2	tw
[Fe(TPP)(2-MeHIm)]•(1,10-phen)	-2.12 ^d	0.90		0.30	20	tw
[Fe(OEP)(1,2-Me ₂ Im)]	-2.19	0.92	0.50	0.37	4.2	16
[Fe(OEP)(2-MeHIm)]	-1.94	0.90	0.48	0.41	4.2	16
[Fe(Tp-OCH ₃ PP)(1,2-Me ₂ Im)]	-2.44	0.95	0.68	0.46	4.2	14
[Fe(Tp-OCH ₃ PP)(2-MeHIm)]	-2.18	0.94	0.58	0.58	4.2	14
[Fe(TPP)(1,2-Me ₂ Im)]	-1.93	0.92	0.53	0.44	4.2	14
[Fe(TTP)(2-MeHIm)]	-1.95	0.85	0.63	0.42	4.2	14
[Fe(TTP)(1,2-Me ₂ Im)]	-2.06	0.86	0.58	0.43	4.2	14
[Fe(TPP)(2-MeHIm)]	-2.40	0.92	0.8	0.50	4.2	13
[Fe(TPP)(2-MeHIm)(2-fold)]	-2.28	0.93	0.8	0.31	4.2	41
[Fe(TPP)(1,2-Me ₂ Im)]	-2.16	0.92	0.7	0.25	4.2	41
[Fe((Piv) ₂ C ₈ P)(1-MeIm)]	-2.3 ^d	0.88	0.40	0.40	4.2	50
deoxyHb	-2.40	0.92	0.7	0.30	4.2	41
deoxyMb	-2.22	0.92	0.7	0.34	4.2	41
[Fe(TPP)(2-MeHIm)] (mol 1)	-2.40 ^e	0.92	0.90	0.37	4.2	21
[Fe(TPP)(2-MeHIm)] (mol 2)	+2.94 ^e	0.97	0.71	0.58	4.2	21
[K(222)][Fe(OEP)(2-MeIm ⁻)]	+3.71	1.00	0.22	0.31	4.2	15
[K(222)][Fe(TPP)(2-MeIm ⁻)]	+3.60	1.00	0.02	0.32	4.2	15
[Fe(Tp _{piv} P)(2-MeIm)] ⁻	+3.51 ^f	0.97			77	39
[Fe(OC ₆ H ₅)(TPP)] ⁻	+4.01	1.03	0.25	0.38	4.2	51
[Fe(O ₂ CCH ₃)(TP _{piv} P)] ⁻	+4.25	1.05	0.30	0.30	4.2	52
[Fe(OCH ₃)(TP _{piv} P)] ⁻	+3.67 ^f	1.03		0.40	4.2	53
[Fe(OC ₆ H ₅)(TP _{piv} P)] ⁻	+3.90 ^f	1.06		0.38	4.2	53
[NaC ₁₂ H ₂₄ O ₆][Fe(Tp _{piv} P)(SC ₆ HF ₄)]	+2.38 ^f	0.84		0.28	4.2	54
[Na(222)][Fe(T _{piv} PP)(SC ₆ HF ₄)]	+2.38 ^f	0.83		0.32	4.2	54
[Fe(T _{piv} PP)(SC ₂ H ₅)] ⁻	+2.18	0.83	0.80	0.30	4.2	54
[Fe(T _{piv} PP)Cl] ⁻	+4.36 ^f	1.01		0.31	77	55

^a millimeters per second. ^b Asymmetry parameter. ^c Line width, fwhm.^d Sign not determined experimentally, presumed negative. ^e Sign is based on a best fit. Because of the complexity of this two-site model, the set of parameters that result in this fit are not necessarily unique. ^f Sign not determined experimentally, presumed positive.

ligated species, including deoxymyoglobin and deoxyhemoglobin, show a negative value of the quadrupole splitting with large asymmetry parameters. The best interpretation is that these species have an unusual ground-state in which the doubly occupied d orbital can be best described as a hybrid orbital comprised of the d_{xz} , d_{yz} , and d_{xy} orbitals, i.e., a very low symmetry orbital. Our recent study shows that a hydrogen bond to imidazole in [Fe(TPP)(2-MeHIm)]₂•2-MeHIm²¹ can have significant effects on Mössbauer properties as well as on structure, yielding a species with properties very much like that of an imidazolate.

Accordingly, the new hydrogen bonded species, [Fe(TPP)(2-MeHIm)]•(1,10-phen) and [Fe(OEP)(2-MeHIm)]•(1,10-phen), were characterized by Mössbauer spectroscopy. For the TPP derivative, the value of quadrupole splitting (ΔE_q) observed at 20 K is 2.12 mm/s, which is similar to the previously reported high-spin iron(II) species as shown in Table 5;^{14,41} the large value of the isomer shift ($\delta \sim 0.90$ mm/s) is also consistent with high-spin iron(II).⁴² There is a substantial temperature variation of the quadrupole splitting values as shown in Table 2. The value of ΔE_q decreases significantly as the temperature is increased. The observed temperature dependence of ΔE_q is similar to the variation seen for imidazole-ligated samples.^{14,15,41,43–49} As discussed previously,¹⁴ the explanation for this temperature variation

**Figure 8.** Data and fits obtained for [Fe(OEP)(2-MeHIm)]•(1,10-phen) at (a) 3, (b) 6, and (c) 9 T applied magnetic field.

is that there are close-lying excited states. The excited states could have the same or differing spin multiplicity relative to the ground state. These data clearly suggest that the new hydrogen bonded species has the same electronic structure as the imidazole-ligated species. The quadrupole splitting values of the OEP derivative, as shown in Table 2, had similar values, but a smaller temperature variation than many other imidazole-ligated species. In order to further study its electronic structure, the Mössbauer spectra were measured in applied magnetic fields.

The application of applied magnetic field Mössbauer spectroscopy provides more detailed information concerning the electronic ground state. The Mössbauer data in strong magnetic fields shown in Figure 8 were fit with the spin Hamiltonian model used by Kent et al.⁴¹

$$H = D[S_z^2 - \frac{1}{3}S(S+1)] + E(S_x^2 - S_y^2) + \vec{H} \cdot \vec{g} \cdot \vec{S} + H^Q - g_N \beta_N \vec{H} \cdot \vec{I} + \vec{S} \cdot \vec{A} \cdot \vec{I}$$

where D and E are the axial and rhombic zero-field splitting parameters that describe the fine structure of the $S = 2$ multiplet, $\vec{A}$ is the magnetic hyperfine tensor, and H^Q gives the nuclear quadrupole interaction:

$$H^Q = \frac{eQV_{zz}}{12} [3I_z^2 - I(I+1) + \eta(I_x^2 - I_y^2)]$$

Q is the quadrupole moment of the ⁵⁷Fe nucleus and $\eta = (V_{xx} - V_{yy})/V_{zz}$, where V_{ii} are components of the electric field gradient. The quadrupole splitting and isomer shift were constrained to the values determined from the zero-field data.

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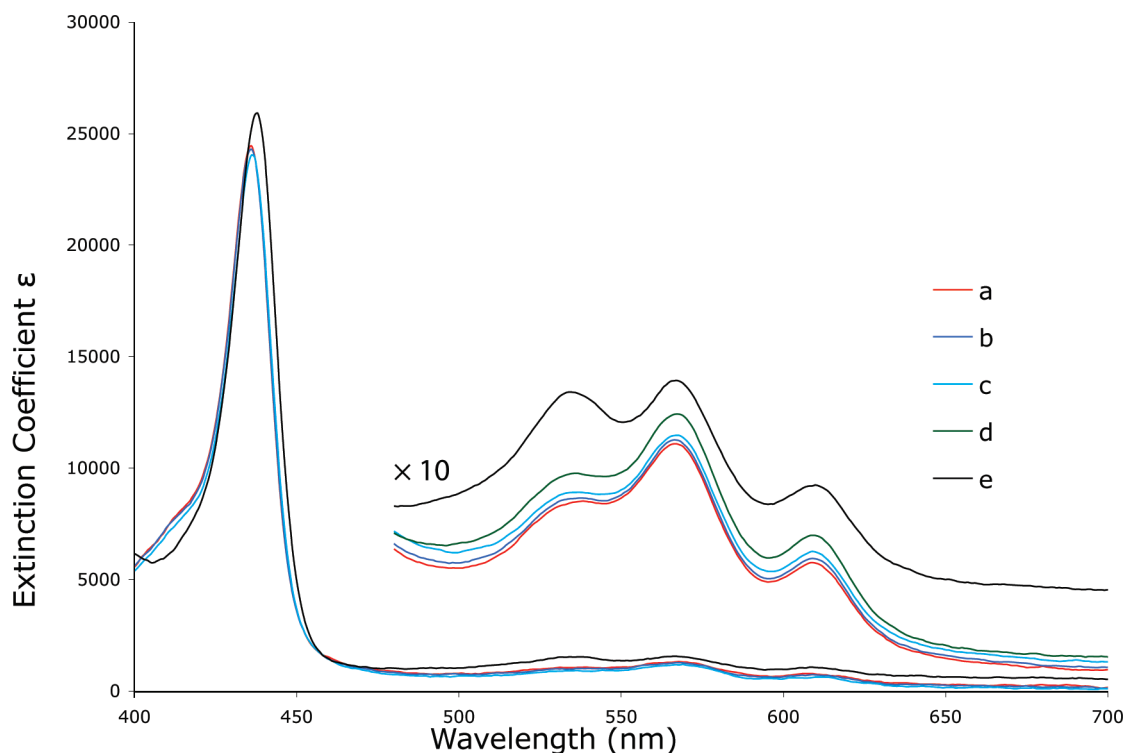


Figure 9. UV-vis spectra taken under argon in CH_2Cl_2 solution. The concentration of $[\text{Fe}(\text{TPP})]$ is 0.5×10^{-3} mol/L; concentrations of 2-methylimidazole are 0.11 mol/L for a, b, c, and d and 1.07 mol/L for e; concentrations of 1,10-phenanthroline are (a) 0, (b) 4.8×10^{-3} , (c) 0.08, (d) 0.16, (e) 0.86 mol/L. The enlarged spectra from 480 to 700 nm are measured in the 1-mm UV cell.

Complete fitting data are given in Table S2 of the Supporting Information.

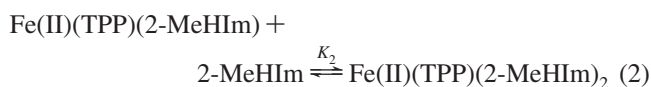
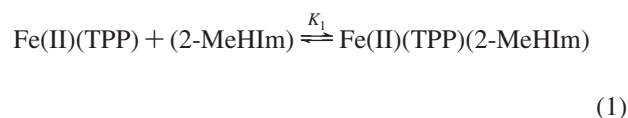
An analysis of the spectra shows the largest component of the electric field gradient, V_{zz} , has a negative value and hence the sign of the quadrupole splitting value is also negative. The asymmetry parameters η for $[\text{Fe}(\text{OEP})(2\text{-MeHIm})] \cdot (1,10\text{-phen})$ (0.76) is relatively large, which is consistent with those for imidazole-ligated species.^{14,41,44–49} The large asymmetry parameter indicates the low symmetry of the electric field gradient (EFG), which is also reflected in the solid-state structure. The results are consistent with our earlier studies of imidazole-ligated species, and we can conclude that the moderate hydrogen bonding in these phenanthroline derivatives is inadequate to lead to the imidazolate-like character observed in $[\text{Fe}(\text{TPP})(2\text{-MeHIm})]_2 \cdot 2\text{-MeHIm}$.²¹

An attempted preparation of a bulk Mössbauer sample of $[\text{Fe}(\text{TPP})(2\text{-MeHIm})] \cdot (1,10\text{-phen})$ led to an interesting observation. The sample was prepared with same reaction conditions that were used for the form **B** preparation except for the use of 20 mL of chlorobenzene rather than 25 mL. The product was then crashed out of solution by the addition of 120 mL of hexanes. This yielded a microcrystalline solid, the Mössbauer spectrum of which displayed two quadrupole doublets overlapped at most temperatures. These additional Mössbauer data are given in Table S3 of the Supporting Information. The quadrupole splitting values for the major component showed very little temperature dependence, whereas the minor component showed strong temperature dependence. The minor component has quadrupole splitting and isomer shift values very similar to those for five-coordinate species, $[\text{Fe}(\text{TPP})(2\text{-MeHIm})] \cdot (1,10\text{-phen})$, as shown in Table 2. For the major component, the values are

very similar to those for low-spin $[\text{Fe}(\text{TMP})(2\text{-MeHIm})_2]$ ⁵⁶ and, thus, most likely to be six-coordinate $[\text{Fe}(\text{TPP})(2\text{-MeHIm})_2]$. However, $[\text{Fe}(\text{TPP})(2\text{-MeHIm})_2]$ has been only reported at low temperature,⁵⁷ suggesting small values for the binding constants. The observation led us to ask how much the hydrogen bonding by 1,10-phenanthroline could increase the binding constants and lead the formation of the six-coordinate species. We have therefore carried out further spectral studies to study this question.

Spectral Studies. Hydrogen bonding of 1,10-phenanthroline with coordinated imidazole in iron(III) and cobalt(III) porphyrinates has been reported to enhance the binding constants of imidazole and thus enhance the formation of six-coordinate species.^{22,23}

In solutions of our system, there are two reactions that must be considered:



Because the three possible iron(II) porphyrinates have substantially different absorption spectra,⁵⁸ UV-vis spectroscopy can be used to monitor the above reactions. The

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results of the spectroscopic titrations are shown in Figure 9. As seen, both the Soret band and visible bands show little change from spectrum **a** to spectrum **d**. Only in spectrum **e**, when the concentrations of 2-methylimidazole and 1,10-phenanthroline are both extremely large, do the spectra show moderate changes. The relative intensities of the two bands at 535 and 565 nm have become more equivalent suggesting an increased intensity of the 535 nm band. The intensity change is consistent with the observations of six-coordinate imidazole-ligated iron(II) porphyrinates.⁵⁸ Thus the change suggests that a small amount of six-coordinate species is formed. But in the Soret band region, there is no extra band or shoulder shown at 425 nm a typical position for a six-coordinate imidazole-ligated iron(II) species.⁵⁸ The spectra indicate that hydrogen-bonding by 1,10-phenanthroline to the coordinated imidazole probably causes an equilibrium shift to the right in reaction 2 and the formation of a small amount of six-coordinate species. However, the large amount of a six-coordinate species seen in the bulk Mössbauer sample seems understood only in terms of solubility issues at high iron(II) concentrations.

Summary. We have prepared new hydrogen-bonded species, [Fe(TPP)(2-MeHIm)]·(1,10-phen) and [Fe(OEP)(2-MeHIm)]·(1,10-phen), with 1,10-phenanthroline as the hydrogen-bond acceptor. Three new structures have been characterized and showed the formation of hydrogen bonds between 1,10-phenanthroline and the coordinated imidazole.

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The structures show that the geometric parameters of iron, such as the Fe–N_p and Fe–N_{im} bond lengths, and displacement of the iron atom out of the porphyrin plane are similar to those in imidazole-ligated species. Further studies of their Mössbauer spectra suggest they have the same electronic configuration as those of the imidazole-ligated species. Unlike the case of [Fe(TPP)(2-MeHIm)]₂·2-MeHIm,²¹ the hydrogen bonds in those new species have little or no effect on the geometric and electronic structures. These studies suggest there is a relatively large gradation of effects from hydrogen bonding on the five-coordinate high-spin, imidazole-ligated iron(II) porphyrinates.

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Supporting Information Available: Figures S1–S4 showing ORTEP diagrams of [Fe(TPP)(2-MeHIm)]·(1,10-phen) (form **B** molecule **2**), [Fe(OEP)(2-MeHIm)]·(1,10-phen) (molecule **1** and **2**), and [Fe(TPP)(2-MeHIm)]·(1,10-phen) (form **A**); Figure S5 showing atomic displacements from the mean plane of the 24-atom core; Table S1 showing analysis results of the core conformations by the normal structural decomposition method; Table S2 giving full details on the fits for applied field Mössbauer measurements of [Fe(OEP)(2-MeHIm)]·(1,10-phen); Table S3 showing variable temperature Mössbauer parameters for crashed out solid sample of TPP derivative; Tables S4–S21 giving complete crystallographic details, atomic coordinates, bond distances and angles, anisotropic temperature factors, and fixed hydrogen atom positions. This information is available as a PDF file. The crystallographic information files (CIF) are also available. This material is available free of charge via the Internet at <http://pubs.acs.org>.

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