Biomimetic Catalysis of a Porous Iron-Based Metal− Metalloporphyrin Framework

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S Supporting Information

[AB](#page-2-0)STRACT: [A](#page-2-0) [porous](#page-2-0) [me](#page-2-0)tal−metalloporphyrin framework, MMPF-6, based upon an iron(III)-metalated porphyrin ligand and a secondary binding unit of a zirconium oxide cluster was constructed; MMPF-6 demonstrated interesting peroxidase activity comparable to that of the heme protein myoglobin as well as exhibited solvent adaptability of retaining the peroxidase activity in an organic solvent.

As new types of functional materials, porous metal−organic
frameworks (MOFs) have experienced rapid development
space the post-decede¹ Their emenability to be decised with over the past decade.¹ Their amenability to be designed with specific functionality makes them stand out from traditional porous materials² an[d](#page-2-0) affords them great potential for various applications, such as gas storage/separation,³ CO₂ capture,⁴ sensors,⁵ and [ca](#page-2-0)talysis.⁶ Their tunable pore spaces and functional pore walls⁷ also provide a new [o](#page-2-0)pp[or](#page-2-0)tunity for biomim[et](#page-2-0)ic catalysis, whi[ch](#page-2-0), however, has rarely been explored.⁸ To develop porous M[O](#page-2-0)Fs as platforms for biomimetic catalysis, one can utilize their pore systems to directly immobiliz[e](#page-2-0) enzymes/proteins⁹ or selectively encapsulate the active centers of enzymes into the cavities of MOFs⁸ or one can regularly arrange the activ[e](#page-2-0) centers of enzymes into MOF frameworks using the derivatives of active center m[ol](#page-2-0)ecules as struts.¹⁰ The later strategy is particularly appealing to mimic heme proteins because iron porphyrin that is the active center [\(](#page-2-0)iron protoporphyrin IX) of heme proteins can be readily functionalized as struts for the construction of MOFs.¹¹ In a continuation of our work on the development of porous metal−metalloporphyrin frameworks (MMPFs),¹² h[ere](#page-2-0)in we report an iron-based MMPF, MMPF-6, which demonstrates interesting peroxidase activity comparable to tha[t o](#page-2-0)f the heme protein myoglobin (Mb).

Dark-red block crystals of MMPF-6 were obtained by reacting iron(III) meso-tetrakis(4-carboxyphenyl)porphyrin chloride (Fe-TCPP-Cl; Figure 1a) with zirconyl chloride under solvothermal conditions. Single-crystal X-ray studies revealed that MMPF-6 has the same structure as the recently reported MOF-545-Fe 13 in which the Fe-TCPP-Cl ligands connect with the in situ generated secondary building units (SBUs) of $\text{Zr}_6\text{O}_8(\text{CO}_2)_8(\text{H}_2\text{O})_8$ (Figure 1b) to form a threedimensional structure consisting of hexagonal and triangular one-dimensional (1D) channels that have diameters of ∼36 and ∼12 Å (atom-to-atom distance), respectively (Figure 1c). The phase purity of the bulk sample MMPF-6 was confirmed by powder X-ray diffraction (PXRD) studies (Figure S1 in the

Figure 1. (a) Iron(III) meso-tetrakis(4-carboxyphenyl)porphyrin chloride ligand. (b) $Zr_6O_8(CO_2)_8(H_2O)_8$ SBU. (c) Hexagonal and triangular 1D channels of MMPF-6. Color scheme: C, gray; O, red; Cl, green; Zr, turquoise.

Supporting Information, SI). The permanent porosity of MMPF-6 was assessed by N_2 sorption at 77 K (Figure 2), [which revealed a typical](#page-2-0) type IV isotherm¹⁴ and a Brunauer− Emmett–Teller surface area of 2100 m²/g comparable to that

Figure 2. N_2 sorption isotherms of MMPF-6 at 77 K (inset: pore-size distribution based upon the DFT model).

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reported in the literature.¹³ The pore-size distribution analysis based upon N_2 adsorption data at 77 K by the use of a density functional theory (DFT) [m](#page-2-0)odel revealed that the pore sizes of MMPF-6 are predominantly around 11 and 33 Å (Figure 2, inset), which are close to the channel sizes observed crystallographically.

It is well-known that heme is the catalytic center for he[me](#page-0-0) proteins, including cytochromes, peroxidases, myoglobins, and hemoglobins, and is capable of performing peroxidation of organic substrates by the use of hydrogen peroxide.¹⁵ The peroxidase activity of heme proteins can be assessed with some standard assays of selected substrates by monitoring th[e ra](#page-2-0)te of increase in absorbance at feature wavelengths of the oxidation products subsequent to the addition of peroxide, such as oxidation of 1,2,3-trihydroxybenzene (THB) to purpurogallin (420 nm) involving oxygen transfer^{16a} and oxidation of 2,2[']azinodi(3-ethylbenzothiazoline)-6-sulfonate (ABTS) to ABTS^{*+} (660 nm) involving electro[n t](#page-2-0)ransfer^{16b} (Scheme 1).

Scheme 1. Reaction Schemes for Oxidation of THB (Top) and ABTS (Bottom) by Hydrogen Peroxide Catalyzed by a Heme-Based Catalyst

The regular arrangement of hemes [iron(III) porphyrin macrocycles] with the five-coordinate iron(III) centers oriented toward the channels in MMPF-6 prompted us to evaluate the potential peroxidase activity of MMPF-6. Catalytic assays were conducted for MMPF-6 and compared with the heme protein Mb. Mb is an oxygen storage $protein$,¹⁷ and its peroxidase activity has also been widely studied,¹⁸ although the activity is lower than that of native peroxidase.¹⁹

Before catalytic assays, the conte[nt](#page-2-0) of the heme centers in MMPF-6 was measured to be 6.18 \times [10](#page-2-0)⁻⁴ mol/g by inductively coupled plasma mass spectrometry experiments on the activated MMPF-6 sample. The relatively lower measured content of the heme centers compared to that estimated from the single-crystal X-ray crystallography derived formula (7.98 × 10^{-4} mol/g) should be attributed to the incomplete removal of the high-boiling-point guest N,N-dimethylformamide molecules during the activation process. Free-base MMPF-6, MMPF-6(Fr), was also synthesized (Figure S1 in the SI) for control experiments.

As shown in Figure 3, without catalyst, the b[lan](#page-2-0)k reaction is going very slowly with rates of only 3.47×10^{-5} mM/s for THB oxidation and 2.87×10^{-6} mM/s for ABTS oxidation (Table 1) in 4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid (HEPES) buffer. MMPF-6 indicates a very fast initial rate [o](#page-2-0)f 1.39×10^{-3} mM/s for purpurogallin formation,²⁰ which is about one-third as fast as free Mb in HEPES buffer (Figure 3a and Table 1). Notably, MMPF-6 demonstrates [i](#page-2-0)mpressive activity not only for oxidation involving oxygen transfer but also for oxidatio[n](#page-2-0) involving electron transfer. In the context of ABTS oxidation, it exhibits a fast initial rate of 8.18 \times 10⁻⁴

Figure 3. Kinetic traces for oxidation of THB in (a) HEPES buffer and (b) an ethanol solution and oxidation ABTS in (c) HEPES buffer and (d) an ethanol solution.

mM/s for ABTS^{*+} formation, which is about one-fourth that for free Mb in HEPES buffer (Figure 3c and Table 1). We reasoned that the slower initial rate for MMPF-6 compared to free Mb should originate from diffusion of the substra[tes](#page-2-0) from solution into the channels of MMPF-6. MMPF-6(Fr) demonstrated much lower activities for both reactions compared to MMPF-6 (Figure 3a,c and Table 1), indicating insignificant contributions from the framework and zirconium oxide cluster. MMPF-6 can be reused for four cy[cl](#page-2-0)es without a significant drop in its peroxidase activity (Table 1) and can also retain its framework integrity, as confirmed by PXRD studies (Figure S1 in the SI), highlighting its heteroge[ne](#page-2-0)ous feature.

We also assessed the peroxidase activity of MMPF-6 in organic solvents, i[n w](#page-2-0)hich heme proteins are known to easily lose their activities.¹⁷ Catalytic assays performed in ethanol solutions indicated that MMPF-6 demonstrated an initial rate of 4.34 × 10<sup>-4<[/s](#page-2-0)up> mM/s for purpurogallin formation and 7.56 × 10[−]⁵ mM/s for ABTS•⁺ formation (Figure 3b,d), meaning an enhancement of ∼107 and ∼215 times, respectively, for the reactions without catalyst (Table 1). In comparison, free Mb exhibited a much slower initial rate (Table1 and Figure 3b,d) because of the agglomeration of [M](#page-2-0)b molecules in ethanol.¹ Slow initial rates were observed for MM[PF](#page-2-0)-6(Fr), furthering supporting the essential role of heme centers in MMPF-6 f[or](#page-2-0) the peroxidase activities.

In summary, a porous iron-based MMPF, MMPF-6, was constructed by self-assembly of the iron(III) meso-tetrakis(4 carboxyphenyl)porphyrin chloride ligand with the in situ generated $Zr_6O_8(CO_2)_8(H_2O)_8$ SBU under solvothermal conditions. MMPF-6 demonstrated interesting peroxidase activity comparable to that of the heme protein Mb in terms of the initial reaction rates in buffer solutions. MMPF-6 retained the peroxidase activity in an ethanol solution, highlighting its adaptability for organic solvents. This work paves a way to develop MOFs as a new type of biomimetic catalyst and also opens up new perspectives for mimicking heme-based proteins. Ongoing work in our laboratory includes the design and synthesis of new iron-based MMPFs for biomimetic catalysis.

Table 1. Summary of Catalysis Results of Oxidation of THB and ABTS in the Presence of 10 mM H_2O_2

^aDiluted to 5 μ M. ^bThe fourth cycle.

■ ASSOCIATED CONTENT

S Supporting Information

Experimental procedures and PXRD patterns. This material is available free of charge via the Internet at http://pubs.acs.org.

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Notes

The auth[ors declare no](mailto:sqma@usf.edu) competing financial interest.

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