## Dynamic Molecular Recognition in a Multifunctional Porphyrin and a Ubiquinone Analogue

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We have recently presented the effective intermolecular interaction between ubiquinone analogue and the porphyrin receptor $\alpha, \alpha, \alpha, \alpha$-meso-tetrakis(2-hydroxynaphthyl)porphyrin (3). ${ }^{1}$ Particularly, $\mathbf{3}$ has four convergent 2-hydroxynaphthyl groups fixed upon the porphyrin ring to obtain good affinity for ubiquinone analogues. ${ }^{2}$ In this communication, we report that the guest molecule tetramethoxy-p-benzoquinone (2) acts as a trigger of atropisomerization of meso-tetrakis(2-hydroxy-4-nonylphenyl)porphyrin (1) to enrich the particular $\alpha, \alpha, \alpha, \alpha$-isomer (1a) with induced fitting interaction ${ }^{3}$ as shown in Scheme I. This is the first example of the equilibrium of atropisomers shifting to the $\alpha, \alpha, \alpha, \alpha$-isomer in homogeneous solution. ${ }^{4}$


Porphyrin 1 was prepared in $14 \%$ yield by condensation of pyrrole and 2-methoxy-4-nonylbenzaldehyde in propionic acid and treatment with $\mathrm{BBr}_{3}$. Four alkyl groups were linked at the para positions of the meso-2-hydroxyphenyl groups to improve the solubility of the parent porphyrin in organic solvents. ${ }^{5}$ HPLC analysis of resulting porphyrin 1 showed a mixture of the four atropisomers $\alpha, \alpha, \alpha, \alpha(\mathbf{1 a}), \alpha, \alpha, \alpha, \beta(\mathbf{1 b}), \alpha, \alpha, \beta, \beta(\mathbf{1 c})$, and $\alpha, \beta, \alpha, \beta$ (1d) in the ratio 1.0:4.4:2.2:0.8, respectively. Silica gel column chromatography at ambient temperature could not separate the atropisomers of the mixture. The rate constant of rotation of one carbon-carbon bond of porphyrin-phenyl ring was determined to be $k=1.7 \times 10^{-5} \mathrm{~s}^{-1}$ at $25^{\circ} \mathrm{C}$ in $\mathrm{CHCl}_{3}$ by HPLC analysis.

The 1a and $\mathbf{1 b}$ bind specifically to $\mathbf{2}$ forming a complex of $1: 1$ stoichiometry ( $K_{1 \mathrm{a}}=6.0 \times 10^{3} \mathrm{M}^{-1}$, and $K_{1 \mathrm{~b}}=3.0 \times 10^{2} \mathrm{M}^{-1}$, respectively, in $\mathrm{CHCl}_{3}$ at $\left.25^{\circ} \mathrm{C}\right) .{ }^{6}$ It is likely that 1a selectively interacts with $\mathbf{2}$ via four efficient hydrogen bonds to maintain the cofacial porphyrin-quinone pair in the most stable form, and the

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## Scheme I



Table I. Relative Proportion of $\alpha, \alpha, \alpha, \alpha$-Isomer 1a among Four Atropisomers of $\mathbf{1}$ in the Presence of $\mathbf{2}$ at Equilibrium in $\mathrm{CDCl}_{3}{ }^{a}$

|  |  |  | proportion of 1a $(\alpha, \alpha, \alpha, \alpha)$ |  |
| :---: | :---: | :---: | :---: | :---: |
| entry | quinone 2 (equiv) | temp $\left({ }^{\circ} \mathrm{C}\right)$ | rel $^{b}(\%)$ | simulated ${ }^{c}(\%)$ |
| 1 | 1.0 | 25 | 58 | 50 |
| 2 | 1.0 | 43 | 39 |  |
| 3 | 1.0 | 9 | 72 |  |
| 4 | 0.5 | 25 | 36 | 34 |
| 5 | 2.0 | 25 | 69 | 64 |
| 6 | 3.0 | 25 | 78 | 71 |
| 7 | 5.0 | 25 | 80 | 76 |
| 8 | 10 | 25 | 80 | 80 |

${ }^{a}[1]=2.4 \times 10^{-3} \mathrm{M} .{ }^{b}$ The relative proportion was determined by the intensity of the ortho-proton resonance by $400-\mathrm{MHz}{ }^{1} \mathrm{H}$ NMR. ${ }^{c} \mathrm{~A}$ computational simulation of the relative proportion was calculated from the binding constants of $\mathbf{2}$ for $\mathbf{1 a}$ and $\mathbf{1 b}$ and the initial isomer ratios $(1 \mathrm{a}: 1 \mathrm{~b}: 1 \mathrm{c}: 1 \mathrm{~d}=1: 4: 2: 1)$.
subsequent atropisomerization is initiated to provide 1a for complexation with $2 .{ }^{7}$

Specific stabilization for the 1a-2 pair is observed by the proton resonances of $\mathbf{1}$ as shown in Figure 1. Although meso-phenyl proton signals in the ${ }^{1} \mathrm{H}$ NMR spectrum of 1 are not clearly distinguishable from each atropisomer (Figure 1a), the coexistence of 2 enables us to assign signals (Figure 1b). The ${ }^{1} \mathrm{H}$ NMR spectrum of a mixture of $\mathbf{1 a}$ and $\mathbf{2}$ indicates exclusive existence

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Figure 1. The $500-\mathrm{MHz}{ }^{1} \mathrm{H}$ NMR spectra of phenyl protons of 1 in $\mathrm{CDCl}_{3}$ at $25^{\circ} \mathrm{C}$ : (a) no quinone added; ( $\mathbf{b}, \mathrm{c}$ ) after addition of $\mathbf{2}$ to a $\mathrm{CDCl}_{3}$ solution of $\mathbf{1}([\mathbf{2}] /[\mathbf{1}]=$ ca 3.0 ), (b) after 0.4 h , (c) after 479 h ; (d) a mixture of isolated 1a and 2. The signal marked with an asterisk (*) is due to $\mathrm{CHCl}_{3}$.
of a $\mathbf{1 a - 2}$ complex ( $\sim 99 \%$ ) in Figure $1 d,{ }^{8}$ and the main peaks of Figure 1c are assignable to a 1a-2 complex. Thus, changes in the population of atropisomer-quinone pairs were monitored by the signal intensity of meso-phenyl protons (Figure 1b,c). For example, addition of 10 equiv of 2 to a $\mathrm{CDCl}_{3}$ solution of 1 caused the relative proportion of 1 a , including the complex and free
 $(500 \mathrm{~h})$ at $25^{\circ} \mathrm{C}$, whereas the relative proportions of $1 \mathrm{~b}, 1 \mathrm{c}$, and

[^2]1d decreased to $17 \%, 2 \%$, and $1 \%$, respectively. Table I summarizes the effect of temperature and concentration of quinone on the proportion of 1 a induced by complexation at equilibrium. (1) Atropisomerization to 1 a from the rest of isomers unambiguously increases as the temperature is lowered (entries 1-3).
(2) The proportion of 1 a is largely influenced by the addition of 2 in the range $0.5-2.0$ equiv (entries 1,4 , and 5 ), whereas the proportion of 1 a is almost saturated above 3.0 equiv of $\mathbf{2}$ (entries 6-8). Addition of 3 equiv of 2 results in the predominant existence of the 1 a isomer ( $78 \%$ of 1 ) as a cofacial complex (more than $97 \%$ of 1a). (3) Computational simulation of isomerization to 1a initiated by molecular recognition was carried out by using two higher binding constants of $2\left(K_{1 \mathrm{a}}\right.$ and $K_{1 \mathrm{~b}}$ ). ${ }^{10}$ In addition, we have assumed that the initial isomers populate in the statistical ratio ( $\mathbf{1 a}: 1 \mathrm{~b}: \mathbf{1 c}: 1 \mathrm{~d}=1: 4: 2: 1$ ) and atropisomerization among the free isomers proceeds at the same rate. The calculated percentage of the $\alpha, \alpha, \alpha, \alpha$-isomer for the complex 1a-2 and free host 1a shows good agreement with the observed value at a higher concentration of 2.
The unique system presented here may provide an example of the induced fitting type of molecular recognition, where specific intermolecular interaction between host and guest causes structural change in the host molecule to form a more stable complex. Elucidation of such dynamic molecular recognition is required to gain a freshinsight into the functions of enzymes and receptors. Further work on the kinetic property of present porphyrin-quinone adducts is in progress, and details on these will appear in future publications.

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Supplementary Material Available: Scheme for the synthesis of compound 1 , experimental procedures, and the details of the calculation of the relative proportion of 1a at equilibrium ( 5 pages). This material is contained in many libraries on microfiche, immediately follows this article in the microfilm version of the journal, and can be ordered from the ACS; see any current masthead page for ordering information.
(10) The binding affinities of $1 \mathbf{c}$ and $1 \mathbf{d}$ with $\mathbf{2}$ seem to be neglected compared with those of $\mathbf{1 a}$ and $\mathbf{1 b}$.


[^0]:    * To whom correspondence should be addressed.
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[^1]:    (6) Face to face complexations between 1a or 1b and 2 were confirmed by the downfield shift of the hydrogen-bonded OH of 1a or 1 b and upfield shift of the $\mathrm{OCH}_{3}$ in the ${ }^{1} \mathrm{H}$ NMR spectra in a similar manner as in previous work. ${ }^{1}$ These affinities were determined by titrimetric measurement of visible spectra with clear isosbestic points in the region $550-700 \mathrm{~nm}$. The titration was manipulated in a short time at $25^{\circ} \mathrm{C}$ after the isolation of each isomer. The isomerization of 1a was carefully examined by HPLC after measurement, and isomerization of 1a to the other three atropisomers was below $5 \%$.
    (7) Thermodynamic parameters of 18-2: $\Delta G^{\circ}=-5.5 \mathrm{kcal} / \mathrm{mol}, \Delta H^{\circ}=$ $-11.6 \mathrm{kcal} / \mathrm{mol}$, and $T \Delta S^{\circ}=-6.1 \mathrm{kcal} / \mathrm{mol}$, respectively, in $\mathrm{CHCl}_{3}$ (without EtOH as a stabilizer) at $25^{\circ} \mathrm{C}$.

[^2]:    (8) The 1 a isomer isolated at $4^{\circ} \mathrm{C}$ maintains its conformation for a time in the presence of 2 due to retardation of isomerization, since 1a interacts very strongly with quinone to form the stable complex.
    (9) Signals of 1a and 1a-2 could not be discriminated in the NMR time scale at room temperature due to fast exchange of free and bound quinone.

