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Control of the stoichiometry in host–guest complexation by redox chemistry of guests: Inclusion of methylviologen in cucurbit[8]uril†

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The binding stoichiometry of a host-guest complex can be effectively controlled by the redox chemistry of the guest: a 1:1 inclusion complex of methylviologen dication (MV²⁺) in cucurbit[8]uril (CB[8]) converts completely and reversibly to a 2:1 inclusion complex of cation radical (MV⁺⁺) in CB[8] upon the reduction of the guest.

Unveiling the principles governing the interplay of redox and molecular recognition is important not only for understanding biological systems but also for designing molecular devices. A number of host-guest systems involving redox active guests have thus been studied to elucidate the interdependence of redox processes and molecular recognition.¹ Cucurbituril (CB[6]), a macrocyclic cavitand comprising six glycoluril units, has a hydrophobic cavity accessible through two identical carbonyl-fringed portals.² Because of its rigid structure and capability of forming complexes with molecules and ions CB[6] has been widely used as a synthetic receptor¹ and as a building block for supramolecular assemblies.3 Our recent synthesis4 of larger cucurbituril homologues, cucurbit[n]uril (CB[n], n = 7and 8) containing seven and eight glycoluril units, respectively, has opened up new opportunities in supramolecular chemistry.5,6 In terms of cavity size, CB[7] and CB[8] are similar to β - and γ -cyclodextrins (β - and γ -CDs), respectively, but their inclusion behavior is often quite different. For example, our recent investigation^{7a} on inclusion behavior coupled with the redox chemistry of methylviologen (N,N-dimethyl-4,4-bipyridinium, MV) in CB[7] revealed that unlike β -CD, CB[7] prefers the charged species, methylviologen dication (MV² and cation radical (MV+·) to the fully reduced neutral (MV⁰) species as guests. Furthermore, dimerization of MV+· is effectively suppressed by forming a stable complex with CB[7] in aqueous solution and the redox process (MV^{2+}/MV^{+*}) of the MV²⁺-CB[7] complex occurs predominantly via a direct electron transfer pathway. This interesting discovery led us to extend our study to the larger host CB[8]. Here we report the unusual redox-coupled inclusion behavior of MV in CB[8] demonstrating the binding stoichiometry of a host-guest complex can be effectively controlled by the redox chemistry of the guest.

We previously reported that CB[8] forms a 1:1 host–guest complex with MV^{2+} (Scheme 1) as evidenced by ¹H NMR and



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Fig. 1 Cyclic voltammograms at scan rate 0.1 V s⁻¹ of a 0.5 mM solution of MV²⁺ in the absence (dotted line), in the presence of 0.25 equivalent (dashed line) and 1 equivalent (solid line) CB[8] in pH 7.0 phosphate buffer. SCE, saturated calomel electrode.

mass spectrometry, even though CB[8] has a cavity large enough to accommodate two MV²⁺ molecules.^{5a} The exclusive formation of the 1:1 complex is also confirmed by UV-visible spectroscopy. The absorption spectrum of MV²⁺ changes gradually with a set of isosbestic points as the concentration of CB[8] increases up to 1 equivalent (see ESI[†]). The observation of the isosbestic points supports that the 1:1 complex is exclusively formed from \hat{MV}^{2+} and CB[8]. The formation constant for the 1:1 complex MV2+-CB[8] determined by the photometric titration is $(1.1 \pm 0.3) \times 10^5$ M⁻¹ at 25 °C in water. The remarkably strong binding of MV²⁺ to CB[8] is in sharp contrast to the fact that γ -CD, which has a hydrophobic cavity with a comparable size, does not bind MV²⁺ appreciably.⁸ As in CB[7],⁷ the strong binding of MV²⁺ by CB[8] is presumably due to the favorable ion-dipole interaction between the positive charge of the guest and the portal oxygen atoms of CB[8] in addition to the hydrophobic effects.

Having established the binding stoichiometry and constant of the complex between CB[8] and MV²⁺, we investigated the interplay between the redox process of MV and its inclusion behavior in CB[8] by using cyclic voltammetry (Fig. 1). As well documented, MV²⁺ shows two reversible waves corresponding to the redox couples MV²⁺/MV⁺⁺ and MV⁺⁺/MV⁰ (dotted line).⁷ In the presence of 0.25 equiv. CB[8] (dashed line), however, two additional redox waves are observed beside the ones for free MV: a reversible wave appears at a potential ~160 mV more *positive* than that for the first redox couple of free MV, and a quasi-reversible wave occurs at a potential ~280 mV more *negative* than that for the second redox couple of free MV (Table 1). In the presence of 1.0 equiv. of CB[8] (solid line), these new waves completely replace the ones for free MV, but they become less reversible, particularly the second one.

This cyclic voltammetric behavior of MV in the presence of CB[8] is quite different from that in the presence of other hosts.

Table 1 Comparison of voltammetric parameters for free $MV^{2+},\,MV^{2+}-CB[7]$ and $MV^{2+}-CB[8]$

	$E^{1}{}_{\rm p}\;(\Delta E_{\rm p})^{a}$	$E^2{}_{\rm p}(\Delta E_{\rm p})^b$	$D^c/\mathrm{cm}^2~\mathrm{s}^{-1}$	Ref.
MV ²⁺ MV ²⁺ –CB[7] MV ²⁺ –CB[8]	-0.704 (64) -0.726 (66) -0.546 (65)	-1.014 (64) -1.164 (122) -1.290 (88)	6.6×10^{-6} 1.8×10^{-6} 1.7×10^{-6}	7 <i>a</i> 7 <i>a</i> This work
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^{*a*} Half-wave potential for the first reduction process expressed in volts *vs*. SCE (saturated calomel electrode). The value in parenthesis represents the potential difference in mV between the cathodic and anodic peak potentials measured at 0.1 V s⁻¹. ^{*b*} For the second reduction process. ^{*c*} Diffusion coefficient determined from the plot of $i_p vs$. $v^{1/2}$.

For example, β - or γ -CD little affects the first reduction wave but causes a large *positive* shift of the second reduction wave because both CDs interact only with fully reduced neutral species MV^{0.8} Furthermore, CB[7] induces a small *negative* shift of the first reduction wave ($\Delta E^{1}_{1/2} \sim -20$ mV) and a large *negative* shift of the second reduction wave ($\Delta E^{2}_{1/2} \sim -110$ mV) since CB[7] has a high binding affinity to MV²⁺ ($K = 2.0 \times 10^{5}$ M⁻¹) but a slightly smaller affinity to MV⁺⁺ and a much smaller affinity to MV^{0.7}

For better understanding of this unusual behavior of MV in the presence of CB[8] we carried out a spectroelectrochemical study. Fig. 2 shows the absorption spectra of MV+ generated by electrolysis of MV²⁺ in the absence (dashed line) and presence (solid line) of CB[8] (applied potential, -0.8 V vs. SCE). It is well known that MV+ exists as an equilibrium mixture of the monomeric and dimeric forms in aqueous solution. When equimolar CB[8] is present, however, only the characteristic absorption of the dimeric species is observed even at a low concentration such as 0.5 mM. Therefore, we conclude that the first redox wave of MV2+ in the presence of CB[8] corresponds to one-electron reduction of MV2+-CB[8] followed by rapid generation of the stable complex $(MV^+)_2$ -CB[8] as shown in Scheme 2.9 The large positive shift of the first reduction peak is apparently related to facile formation of the stable complex $(MV^{+})_2$ -CB[8]. From the spectroelectrochemical study the apparent dimerization constant of MV+ in the presence of equimolar CB[8] is estimated¹⁰ to be $2 \times 10^7 \,\mathrm{M^{-1}}$ (ESI[†]) which is about 10⁵ times larger than that of MV⁺ alone in aqueous media. Although CDs¹⁰ and surfactants¹¹ are known to enhance the dimerization of MV⁺⁺, such a large enhancement has never been observed. The remarkable stability of the dimer within CB[8] may be attributed to the unique properties of CB[8] as a host providing the dimer not only with a hydrophobic cavity of a right size but also with polar oxygen atoms at the portals for ion-dipole interaction with the guest. Most importantly, our work demonstrates that the 1:1 inclusion complex MV2+-



Fig. 2 Absorption spectra of MV⁺⁺ generated from a solution of MV²⁺ (0.5 mM) in phosphate buffer (pH = 7.0) using an indium tin oxide glass electrode (applied potential, -0.8 V vs. SCE). Solid and dashed lines represent spectra in the presence of 1 equiv. of CB[8] and in the absence of CB[8], respectively.



CB[8] converts completely and reversibly to the 2:1 inclusion complex of $(MV^{+})_2$ -CB[8] upon reduction of the guest, as illustrated in Scheme 2. To the best of our knowledge, this is the first example demonstrating that the binding stoichiometry of a host-guest complex can be effectively controlled by the redox chemistry of the guest. The large negatively shifted second redox wave of MV in the presence of CB[8] is apparently related to the reduction of $(MV^{++})_2$ -CB[8], but at this moment we do not clearly understand this process. We are currently investigating these electrochemical processes in more detail in conjunction with the mechanism of the electron transfer coupled with inclusion process.

In conclusion, we present a unique example demonstrating that the binding stoichiometry of a host–guest complex can be effectively controlled by the redox chemistry of the guest. This unprecedented redox control of the stoichiometry in host–guest complexation may provide a working principle for electrochemically controllable molecular machines.

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