

Electrochemically tuneable hydrogen bonding interactions between a phenyl-urea terminated dendrimer and phenanthrenequinone†

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Electrochemically tuneable hydrogen bonding interactions are described between a phenyl-urea terminated dendrimer and phenanthrenequinone.

The construction of hyperbranched macromolecular architectures featuring engineered non-covalent interactions is a burgeoning area within contemporary dendrimer chemistry.¹ In particular, the strength, directionality and selectivity of hydrogen bonding processes have proved to be particularly successful supramolecular interactions for modifying the periphery of dendrimer architectures.² Indeed, the reversibility of assemblies of this type has allowed the creation of responsive supramolecular dendrimers with architectures that can be assembled/disassembled under the control of an external stimulus.³ It is likely that responsive dendrimers constructed using supramolecular interactions will continue to be an attractive alternative to their covalently functionalised brethren, as the self-assembly method significantly reduces the lengthy synthetic routes required to fabricate elaborate functional macromolecules of this type.

The application of electrochemistry to modulate the efficiency of hydrogen bonded interactions is a rapidly developing field within supramolecular chemistry.⁴ For example, it has been shown that the electrochemical reduction of the phenanthrenequinone component of complex **1.2**, results in a significant increase in hydrogen bonding efficiency between these units.⁵ Moreover, it was also shown that the efficiency could be modulated in polar (DMF) and non-polar (CH₂Cl₂) environments, thereby nominating this system as an attractive building block for the construction of dendrimers with electrochemically controllable complexation properties.⁶ Here, we report the electrochemically tuneable hydrogen bonding interactions between phenyl-urea terminated dendrimer **4** and **1** (Fig. 1).⁷

Dendrimer **4**† was readily synthesised from commercially available polypropylenimine hexadecaamine dendrimer (DAB-Am-16) and excess phenyl isocyanate using a similar method to that previously reported.⁸ ¹H NMR spectroscopy was used to confirm the formation of hydrogen bonds between electroactive guest **1** and hosts **3** and **4**. The addition of aliquots of **1** to a solution of **3** or **4** in DMF-d₇ led to a broadening and small downfield shifts (~0.1 ppm) of the resonances for both urea protons, thereby indicating the formation of a low-affinity hydrogen bonded complex (see ESI†).

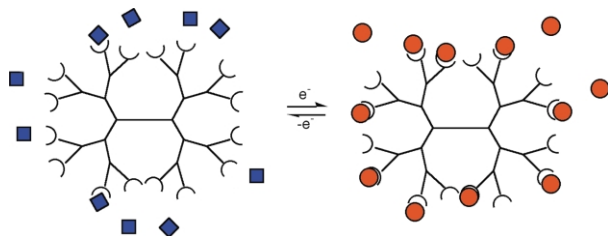
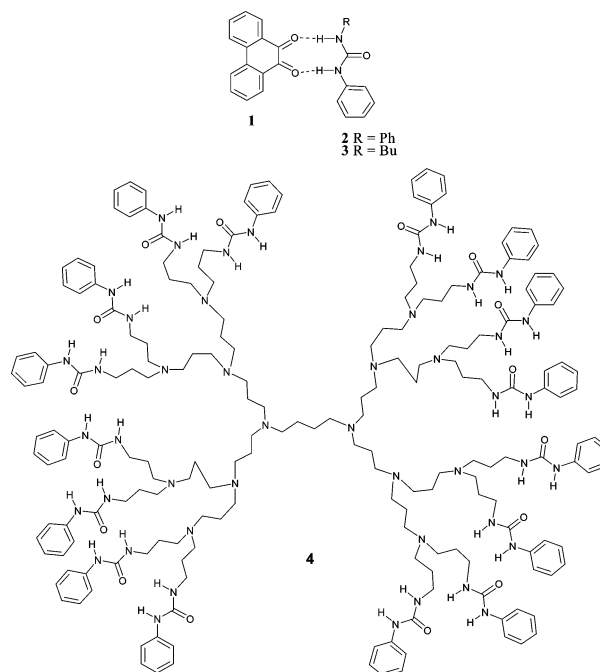


Fig. 1 Schematic representation of the electrochemically controlled complexation of dendrimer **4** with phenanthrenequinone **1**.

† Electronic supplementary information (ESI) available: ¹H NMR titration and cyclic voltammetry data. See <http://www.rsc.org/suppdata/cc/b2/b211370h/>



With hydrogen bond formation confirmed between the urea hosts and **1** in DMF, we next investigated their electrochemically controlled complexation using cyclic voltammetry (CV).⁹ The addition of an excess of dendrimer **4** to a solution of **1** (in DMF containing 0.1 M Bu₄NPF₆) immediately resulted in a +90 mV shift in the half-wave potential of the reduction wave of **1** (Fig. 2). This positive shift indicates a substantial stabilization (8.7 kJ mol⁻¹) of the radical anionic state of **1**, corresponding to over a 30-fold increase in binding affinity between this unit and the dendrimer periphery.¹⁰ A notable

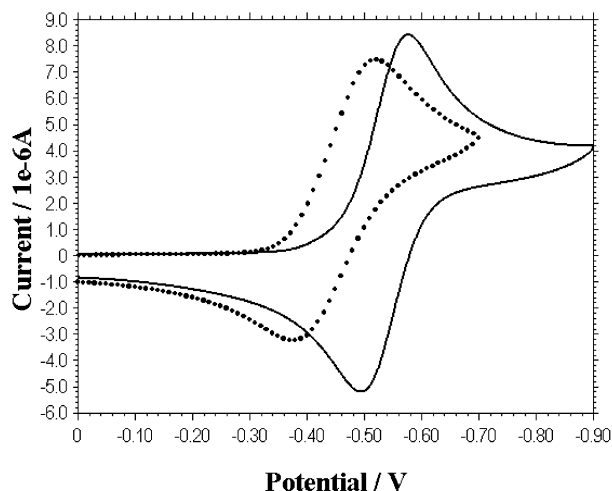


Fig. 2 CV data for **1** ($\sim 9 \times 10^{-4}$ M in DMF) recorded in the absence (—) of and in the presence (·····) of an excess ($\sim 6 \times 10^{-3}$ M) of urea dendrimer **4**.

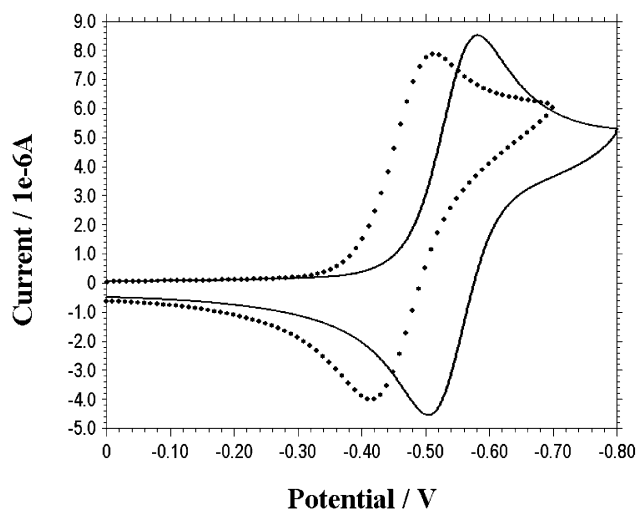


Fig. 3 CV data for **1** ($\sim 9 \times 10^{-4}$ M in DMF) recorded in the absence (—) of and in the presence (·····) of an excess ($\sim 9 \times 10^{-2}$ M) of urea **3**.

feature of the CV of **1** following the addition of an excess of dendrimer **4**, is the ~ 20 – 30% decrease in peak currents for this reduction wave, which is particularly evident during the re-oxidation of the $\mathbf{1}^{\cdot-}$ to the **1** state. This observation further supports complex formation between dendrimer **4** and **1**, by indicating that hydrogen bonding interactions with the relatively large dendrimer species slows down diffusion of **1** to the working electrode surface.¹¹ Evidence supporting the reduced reversibility of the redox wave of **1** being a consequence of the slower diffusion rate of the dendrimer/phenanthrenequinone complex, was obtained by measuring the effect scan rate has on the reversibility of the redox wave of **1**. At low scan rates (e.g. 25 mV s^{-1}), a reversible waveform was obtained for the reduction of **1** to $\mathbf{1}^{\cdot-}$ in the presence of excess **4**, however, at faster scan rates (e.g. 5 V s^{-1}) the waveform became considerably more irreversible (see ESI†).

To further corroborate electrochemically controlled host-guest complexation between **1** and **4**, we explored the changes in solution electrochemistry of **1** following the addition of an excess of model compound **3**.¹² A slightly smaller shift of $+83 \text{ mV}$ was observed for the redox wave of **1** following the addition of **3**. However, in contrast to the CV data obtained for dendrimer **4**, a relatively small decrease (~ 5 – 10%) in peak currents was typically observed (Fig. 3). This result is to be expected as host-guest complexation between **1** and **3** should not significantly perturb the diffusion rate of **1** to the working electrode.

In summary, we have established that the efficiency of hydrogen bonds formed between dendrimer **4** and **1** can be significantly enhanced upon electrochemical reduction of the latter to its radical anion state. In particular, by relating the perturbation of the redox potential of **1** to the thermodynamic data obtained for an analogous hydrogen bonded complex, the very weak hydrogen bonds formed between the neutral species ('off' state) can be increased 33-fold by reduction of **1** ('on' state). Therefore, this study paves-the-way for the development of new electrochemically controlled dendrimer architectures with biomimetic and device applications; the results of our investigations towards these goals will be reported in due course.

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Notes and references

† Selected data for **4**: Yield 60%. IR (KBr) ν (cm^{-1}) 3339, 2945, 2805, 1650, 1598, 1554, 1499, 1441, 1312, 1240, 753, 693; ^1H NMR (200 MHz, DMSO-d_6): δ 8.40 (br. 16H, PhNHCONH), 7.36 (d, 32H, $J = 7.9 \text{ Hz}$, Ph),

7.15 (t, 32H, $J = 7.7 \text{ Hz}$, Ph), 6.85 (t, 16H, $J = 7.2 \text{ Hz}$, Ph), 6.23 (br. t, 16H, PhNHCONH), 3.10 (br. 32H, NHCH_2), 2.33 (br. 84H, CH_2NCH_2), 1.51 (br. 56H, $\text{CH}_2\text{CH}_2\text{CH}_2$) 1.38 (br. 4H, $\text{NCH}_2\text{CH}_2\text{CH}_2\text{CH}_2\text{N}$); ^{13}C NMR (50 MHz, DMSO-d_6): δ 155.85, 140.87, 129.03, 121.50, 118.31, 52.15, 51.51, 38.64, 27.84; MS (MALDI-TOF): $m/z = 3592.7$ (M^+).

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- All electrochemical experiments were performed using a CH120A electrochemical workstation. The electrolyte solution was prepared from recrystallised Bu_4NPF_6 using spectroscopic grade dimethylformamide (0.1 M). A three electrode configuration was used with a Pt working electrode, a Ag/AgCl reference electrode and a platinum wire as the counter electrode. Scan rate = 100 mV s^{-1} (unless otherwise stated). $T = 25^\circ\text{C}$. The solution was purged for several minutes with N_2 prior to recording the electrochemical data; the voltammograms were recorded under an atmosphere of N_2 .
- We expect that the addition of an excess of dendrimer **4** to a solution of **1** would principally result in the formation of complex of stoichiometry **1.4**. Thus, we can estimate the change in K_a following the electrochemical formation of $\mathbf{1}^{\cdot-}\cdot\mathbf{4}$ using a thermodynamic cycle which can be expressed mathematically using: $K_a(\text{red})/K_a(\text{ox}) = e^{(nF/RT)(E_{1/2}(\text{bound}) - E_{1/2}(\text{unbound}))}$. $K_a(\text{red})$ and $K_a(\text{ox})$ are the association constants in the reduced and oxidised forms, and $E_{1/2}(\text{bound})$ and $E_{1/2}(\text{unbound})$ are the half-wave redox potentials in the receptor bound and unbound states. A $K_a(\text{ox})$ of 1 M^{-1} has previously been determined for the hydrogen bonded complex formed between 1-phenyl-3-propyl urea and **1** using simulated electrochemical data. As it was not possible to fully reach saturation in our electrochemical experiments, a lower limit for $K_a(\text{red})$ for $\mathbf{1}^{\cdot-}\cdot\mathbf{4}$ complex can be estimated to be 33 M^{-1} .
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- In order to ensure comparability between electrochemical measurements (i.e. maintaining the same number of urea recognition units per molecule of **1**), 16 molar equivalents (compared to the amount of dendrimer **4** added in the previous experiment) of model compound **3** was added to the CV cell.