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Dendritic nanodevices

Dendritic encapsulation as probed in redox active core dendrimers

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Abstract

The relationship between dendrimer primary structure, conformation, and core encapsulation behaviors is highlighted. This work has involved the synthesis of several novel types of redox-active core dendrimers. Assessment of diffusion measurements provides a first indication of molecular size. Conformation can be probed by exploiting electrochemical and magnetic properties of the core and correlation of these behaviors to computational models. *To cite this article: C.B. Gorman, C. R. Chimie 6* (2003).

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1. Introduction

The hyperbranched architecture of dendrimers has led to the proposal and subsequent study of these types of molecules for encapsulation. This type of encapsulation can involve small molecule guests within the dendrimer or covalent encapsulation of units within the molecule. In this latter class, the unit to be encapsulated is typically placed at the 'core' of the molecule – that is, at its topological center. Encapsulation in this sense implies that the dendritic arms around the core display themselves in space so as to efficiently sterically shield the core from the surrounding environment. Thus, the efficacy of encapsulation relies on a link between the hyperbranched, primary structure of the dendrimer and its conformation (e.g., its tertiary structure by analogy to a protein). Early theories of dendrimer conformation suggested that they adopted 'dense shell' or sterically congested conformations [1]. Later, it was argued that dendrimers tend to form con-

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formations with a uniform radial density and substantial distribution of the tips of the molecule throughout it [2, 3]. This latter conformation may or may not offer ideal conditions for encapsulation. Here, we highlight work that illustrates that dendrimers can be useful in encapsulation strategies, but mere hyperbranching is insufficient to provide ideal encapsulation.

Covalent encapsulation of a core moiety is typically studied in two ways. The photophysical behaviors of a core moiety can be probed as a function of the type and generation of the dendrimer [4–6]. Specifically, the degree of exciton quenching (e.g. intermolecular selfquenching of luminescence) can be attenuated or eliminated by placing the lumophore at the center of a dendrimer [7, 8]. Also, the degree of quenching of luminescence by dioxygen and other small quencher molecules can be attenuated in a similar manner [9–11]. These functions have been discussed in detail previously [4–6].

The other manifestation of core encapsulation is via changes in the electronic (electrochemical) properties of the dendrimer core. Specifically, as a redox-active

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core moiety is shielded, the relative rate of electron transfer to/from that core decreases. The dendrimer can also create a different type of microenvironment around this moiety as compared to the surrounding solvent. To the extent that this occurs, the thermodynamic redox potential of the redox active unit can change. Here, the former effect is emphasized, although the latter has been studied [12] and reviewed [4–6] as well.

Molecules that encapsulate a redox center have two potentially important functions in the study of both materials sciences and life sciences processes. First, the shielding of a redox center is the dominant structural effect of metalloproteins. To date, the general effects that structure has on redox potential are still unclear. Redox active core dendrimers offer the opportunity to elucidate general structural effects [4, 5, 12–25]. Second, controlling rate, driving force and direction of electron transfer are key elements in the design of molecular electronic devices. As will be shown, dendrimers offer a model system to show how an "insulating" macromolecular architecture affects these parameters. Dendrimers may be potentially useful in the design of rational, molecular electronics.

2. Synthesis of metal cluster core dendrimers

The ideal, redox active core dendrimer involves a core with well-understood electrochemical properties and a geometry that disposes the dendron arms about it in a uniform (symmetrical) fashion [20]. With respect to mimicking protein function, Diederich et al. first showed that dendrimers could be used to model protein environment. This group synthesized several classes of porphyrin-core dendrimers and showed that the dendrimer structure could affect redox potential as well as the kinetics of oxygen binding [13,14,16,26–30]. The facility of encapsulating biologically relevant clusters into dendrimers has further been illustrated by Aida et al. [31], who prepared dendrimer encapsulated copper(µ-oxo)dimers (found in methane monooxygenase and ribonucleotide reductase) and Basu et al. who prepared dendrimer encapsulated oxo-Mo(V)tetrasulfide units (found in DMSO reductase) [32]. Significant contributions have also come from Kaifer et al. [33-38] who have reported dendrimers based on ferrocenyl and viologen core moieties and Zheng et al., who

have reported dendrimers based on a Re⁶Se⁸ cluster core moiety [39–41].

The synthesis of dendrimers with these types of symmetric cores requires some type of ligand exchange reaction that is efficient. In general, it is very difficult or impossible to separate side products that result from the incomplete attachment of the requisite number of arms around the cluster. To this end, we have explored the conditions for efficient ligand exchange around two clusters, Fe₄S₄ and Mo₆Cl₈. Fig. 1 illustrates a general protocol for each type of ligand exchange. In the case of Fe_4S_4 , Holm et al. had previously shown that bulky aliphatic thiolates can efficiently be displaced by aromatic thiols with the loss of the initial component to vapor [42-45]. This scheme works well for the synthesis of iron-sulfur cluster core dendrimers - molecules with molecular weights of ca. 25 000 have been illustrated [18,19,24]. In the case of Mo_6Cl_8 , phenol derivatives can substitute for methoxy groups or potassium phenolate can substitute for triflate groups [23]. These types of reactions have had an important impact on the systematic study of dendritic encapsulation of redox active cores.

3. Dendrimer size in solution

The size of a dendrimer as estimated from a diffusion coefficient gives a first indication of conformation. Moreover, diffusion coefficients provide the element of mass transfer that is required to calculate electron transfer rates from voltammetry as illustrated below. To this end, we sought to measure diffusion coefficients of dendrimers. In Fig. 2, the general structures of flexible and rigid iron-sulfur cluster core dendrimers are shown. Each type of dendrimer has been synthesized for generations one through four. Two complementary techniques have been employed to measure diffusion coefficients for these molecules. The results for the flexible series are shown in Fig. 3. Taking advantage of the electroactive nature of the iron-sulfur cluster, diffusion measurements could be obtained by chronoamperometry. These values compared well to those obtained by pulsed field gradient spin echo NMR. Both techniques gave values expected for a systematic increase in molecular size with generation. Comparison of radii of gyration for these molecules calculated using the Stokes-Einstein equation to those derived from computations (below) indicated



Fig. 1. A general protocol for each type of ligand exchange used to form metal cluster core dendrimers.

that these molecules assumed globular, packed conformations [24].

4. Dendritic encapsulation – Effect on electron transfer rates

Heterogeneous electron transfer rates for iron–sulfur cluster core dendrimers were determined by fitting Osteryoung square wave voltammetry data to a quasireversible heterogeneous electron transfer model as described previously [24]. The results are presented in Table 1. One can observe that increasing generation decreases the rate of electron transfer in the flexible dendrimers. Thus, the effective encapsulation increases with increasing generation. This effect has been examined nicely in the very slow rate regime by using thin layer voltammetry on large dendrimers [46].

However, the decrease in rate does not reflect an increase in effective electron transfer distance that would be expected if the dendrimer were in an extended conformation. Examining an analogous series of rigid dendrimers, the electron transfer rate was found to decrease much faster with generation (or molecular weight). This result indicates that the primary structure of the dendrimer is important in biasing the conformation so as to maximize the effective distance (and thus rate) of electron transfer.

To further illustrate the effect of dendrimer structure on conformation, as evidenced by electron transfer rate, two dendrimer isomers were prepared and studied (Fig. 4). We and others [47–50] have hypothesized that primary structural elements that enforce backfolding should increase the degree of steric congestion around the core moiety of the dendrimer. This behavior should lead to more effective encapsulation. Indeed, as illustrated in Table 1, the backfolded isomer showed a lower rate of electron transfer than the extended isomer. This result is consistent with a more congested architecture and a higher effective distance for electron transfer in the former isomer as compared to the latter.



Fig. 2. Generation-3 examples of flexible and stiff Fe_4S_4 -cluster core dendrimers previously prepared in the Gorman group.

Concomitant with this observation is a shift in the redox potential of the backfolded isomer compared to the extended isomer. This result is consistent with the iron–sulfur cluster being found in a more hydrophobic microenvironment in the backfolded molecule as compared to the extended molecule. Thus, primary structure can be used to bias dendrimer conformation in an experimentally observable manner.



Fig. 3. Illustration of diffusion coefficients for flexible dendrimers of generations 1 through 4. These data were obtained using pulsed field gradient spin echo NMR (Stejskal–Tanner plots are shown at the top right) and chronoamperometry (Cottrel plots are shown at the bottom right).

Structure	$k_0 (\times 10^3 \mathrm{cm s^{-1}}) \mathrm{OSWV^a}$	$E_{1/2}$ (mV) OSWV or CV ^{a,c}	$\alpha OSWV^a$
G0Flex	7.43 (1.31)	-1344 (4)	0.57 (0.07)
G1Flex	6.15 (0.64)	-1359 (9)	0.54 (0.06)
G2Flex	3.29 (0.44)	-1374 (13)	0.33 (0.01)
G3Flex	0.76 (0.03)	-1374 (28)	0.29 (0.05)
G4Flex	0.13 (0.06)	NA ^b	0.47 (0.01)
G0Rig	6.24 (0.59)	-1351 (5)	0.56 (0.08)
G1Rig	4.12 (0.78)	-1322 (14)	0.54 (0.04)
G2Rig	2.35 (0.09)	-1324 (9)	0.37 (0.03)
3,2 (extended)	4.79 (0.92)	-1494 (4)	0.48 (0.03)
3,2B (backfolded)	0.64 (0.15)	-1618 (2)	0.50 (0.005)

Heterogeneous electron transfer rate constant (k_0), reduction potential ($E_{1/2}$) and α for the one electron redox couple [Fe₄S₄(S–Dend)₄]^{2–/3–}

^aThe magnitude of the 90% confidence interval is indicated in parentheses.

^bNot determined in a fit of an irreversible wave.

^c Thermodynamic redox potential.

Table 1

5. Dendrimer conformation: the link between primary structure and encapsulation behavior

Regrettably, no methods to directly observe dendrimer conformation have been successful. However, we have explored two methods that indicate some important conformational features of these molecules and help to explain the encapsulation phenomena described above.

The longitudinal relaxation time constants (T_1) of the protons in a series of dendrimers that alternatively had paramagnetic ($[Fe_4S_4(SR)_4]^{2-}$ R = dendron) and diamagnetic (tetraphenylmethane) cores were compared [21]. The T_1 values of the phenyl and benzyl protons in the paramagnetic core dendrimers were attenuated compared to analogous protons in the diamagnetic core dendrimers (Fig. 5). This observation indicated that protons in each set of topologically different repeat units (generation) of the dendrimer approach the core of the molecule closely in space. This conclusion is consistent with the computed radial density distributions of the different generations calculated from molecular dynamics simulations [25].

Using a conformational searching protocol, the number of low-energy conformers (within a Boltzman cutoff of 10 kcal mol⁻¹ at 500 K of the lowest energy conformer) were correlated with the relative offset of the core (Fig. 6) [24]. An offset of zero would find the core at the center of mass of the molecule. An offset of one would find the core at the edge of the molecule. One can see a substantially more offset core in all of the flexible models compared with all of the rigid models. In pictorial representations of the lowest en-



Fig. 4. Structures of two iron-sulfur cluster core dendrimer isomers prepared.



Fig. 5. T₁ values (in ms) of aromatic protons in an iron-sulfur cluster core dendrimer as compared to a diamagnetic analogue.

ergy conformations (Fig. 6, top), the core is observed to be very centrally located in the rigid molecule but virtually at the molecular edge in the flexible molecule. This model helps to explain why the effective electron transfer distance differs in these two types of molecules. This type of conformational search has been







Fig. 6. Relative core offsets (0 = core at center and 1 = core at edge) of flexible and stiff dendrimers from conformational search results presented previously [24]. Molecular models of conformation snapshots are shown above the histogram of each search result.

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applied to the dendrimer isomer pair as well and will be reported separately.

6. Conclusions

These data support several conclusions relevant to dendritic encapsulation. A simple dendrimer does encapsulate its core, but not necessarily in an efficient manner (at least for generations that are relatively accessible, synthetically). However, other structural elements can enforce a conformation in which the core is more closely held in the center of the molecule. Here, the relative rigidity and the use of backfolding have been illustrated as structural motifs to achieve this goal. Undoubtedly, other useful structural elements exist, and this gives an exciting purpose to the continued exploration of structure-property relationships that govern dendritic encapsulation.

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