

Template-Directed Synthesis of a Conjugated Zinc Porphyrin Nanoball

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Supporting Information Placeholder

ABSTRACT: We report the template-directed synthesis of a π -conjugated 14-porphyrin nanoball. This structure consists of two intersecting nanorings of 6 and 10 porphyrin units. Fluorescence up-conversion spectroscopy experiments demonstrate that electronic excitation delocalizes over the whole 3D π -system in less than 0.3 ps, if the nanoball is bound to its templates, or over 2 ps if the nanoball is empty.

Spurred by the discovery of buckminsterfullerene,¹ chemists have sought rational strategies for the synthesis of 3D π -conjugated geodesic cages. Stepwise chemical synthesis has not yet been used to prepare any fullerenes, except C₆₀ (ref. 2), but several fullerene-like ball-shaped π -conjugated hydrocarbons have been synthesized recently.³ The high dimensionality of these π -systems is expected to enhance their electronic delocalization, compared with 1D or 2D molecular semiconductors.⁴⁻⁶ Shape-persistent molecular cages are also in demand for their gas adsorption properties, which mimic those of zeolites.⁷ The best routes to molecular cages or capsules use reversible reactions that allow error correction, such as metal coordination⁸ or boronic ester condensation,⁹ but these reactions do not produce π -conjugated connections. In principle, reversible reactions such as imine formation,¹⁰ alkene metathesis¹¹ and alkyne metathesis¹² could be used to construct π -conjugated cages, but so far the cages made by these reactions lack long-range conjugation.^{7,10-12} Template-directed coupling, under kinetic control, is an alternative strategy for preparing large macrocycles and cages.¹³ Here we show how simple molecular templates can be used to synthesize the first fully π -conjugated porphyrin ball **b-P14** (Figure 1). This 14-porphyrin prolate ellipsoidal cage consists of two perpendicular intersecting conjugation pathways, containing 6 and 10 porphyrin units respectively. Structures of this type are valuable models for photosynthetic light-harvesting systems.¹³⁻¹⁵ When all 14 porphyrin units in **b-P14** are bound to templates, locking the conformation, excited state energy delocalization occurs over the whole system within 0.3 ps, whereas in the absence of the templates excitation is distributed over the ball with a time constant of about 2 ps.

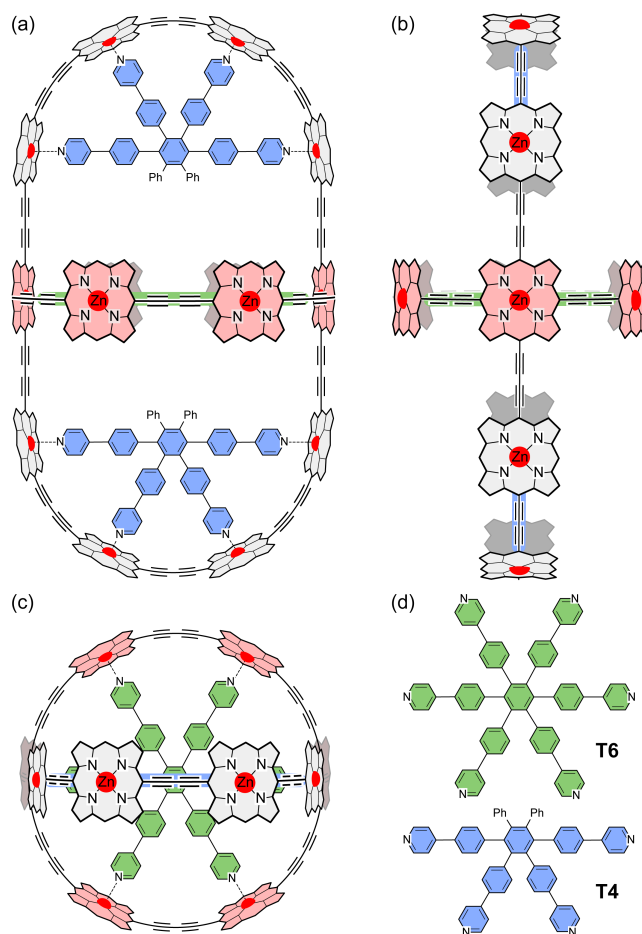


Figure 1. (a)–(c) Three orthogonal views of the structure of the 14-porphyrin ball template complex **b-P14·T6·(T4)₂** and (d) the templates **T6** and **T4**. The *meso*-aryl side groups on the porphyrins are omitted for clarity.

The porphyrin nanoball **b-P14** was synthesized as shown in Figure 2, using templates **T6** and **T4** (Figure 1). This route starts from a 6-porphyrin nanoring template complex with four terminal alkyne substituents, **c-P6(H)₄·T6**, which was prepared as reported previously.¹⁶ This ring was coupled to four porphyrin dimers **P2-CPDIPS**, followed by removal of the CPDIPS protecting groups, to give the extended nanoring **c-P6(P2)₄·T6**. The four-legged template **T4** was then used to

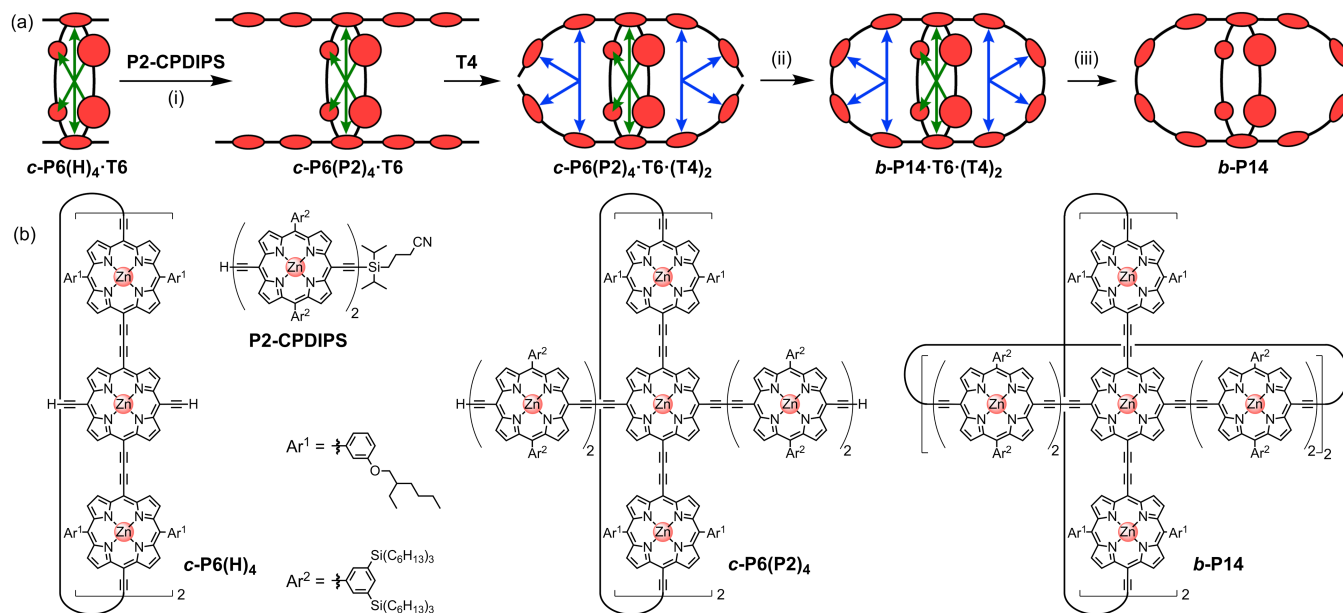


Figure 2. (a) Schematic synthetic route for **b-P14**. Reaction conditions: (i) **P2-CPDIPS**, Pd(PPh₃)₂Cl₂, CuI, 1,4-benzoquinone, CHCl₃/*i*-Pr₂NH, 41%; then TBAF, CH₂Cl₂, 97%; (ii) Pd(PPh₃)₂Cl₂, CuI, 1,4-benzoquinone, CHCl₃/*i*-Pr₂NH, 51%; (iii) DABCO, size exclusion chromatography, toluene/pyridine, 100%. (b) Chemical structures of **c-P6**, **P2-CPDIPS**, **c-P6(P2)₄** and **b-P14**.

close the 10-porphyrin ring, yielding the bicyclic nanoball **b-P14·T6·(T4)₂**. The templates can readily be displaced from this cage by high concentrations of a competing ligand;¹⁷ quinuclidine or DABCO remove both templates, giving **b-P14**, whereas pyridine selectively removes **T4**, giving **b-P14·T6**. Both **b-P14** and **b-P14·T6** were fully characterized (see SI). Two types of aryl solubilizing groups were used in this molecular design (Ar¹ and Ar², Figure 2) so as to confer high solubility while avoiding excessive steric congestion.

Gel permeation chromatography (GPC) confirmed the purity of the nanoball and showed that its molecular weight is in the expected range (ca. 20 kDa). The ¹H NMR spectrum of **b-P14·T6·(T4)₂** in CDCl₃ at 298 K is consistent with the expected *D*_{2h} gross symmetry and all the signals were assigned by COSY and NOESY spectroscopy, except for unresolved aliphatic multiplets in the 0.4–2.0 ppm region (see SI). This ¹H NMR spectrum shows that the nanoball consists of a mixture of conformers with ethylhexyl chains pointing into and out of the cavity; these rotamers are in slow exchange on the NMR timescale, with ca. 40% of the alkoxy-chains pointing towards the center of the ball. This ratio scarcely changes on removal of the **T4** templates. When all the templates are removed, the porphyrins of the 6-ring rotate rapidly, simplifying the NMR spectrum. Diffusion-ordered ¹H NMR spectroscopy (DOSY) experiments show that **b-P14·T6·(T4)₂** has a diffusion coefficient of $1.52 \times 10^{-10} \text{ m}^2 \text{ s}^{-1}$ (700 MHz, 298 K, CDCl₃), which corresponds to an effective hydrodynamic radius of 27 Å, calculated using the Stokes-Einstein equation for a spherical molecule.

The UV-vis-NIR spectrum of **b-P14·T6·(T4)₂** (Figure 3a, black curve) is essentially the sum of the absorption spectra of its two component rings, as modeled by **c-P6·T6** (the *D*_{6h} ring complex, with 3,5-bis(triethylsilyl)phenyl substituents)¹⁸ and **c-P10·(T5)₂** (where **T5** is the version of **T6** with five pyridyl sites),^{16–20} although the absorption spectrum of **b-P14·T6·(T4)₂** is slightly red-shifted, demonstrating greater π -conjugation (SI

Figure S82). When **b-P14·T6·(T4)₂** is treated with quinuclidine to displace the templates, two distinct denaturation processes are observed in the UV-vis-NIR titration (Figure 3): At quinuclidine concentrations of 3–30 mM (Figure 3b), the **T4** units are displaced leading to disappearance of the sharp peak at 883 nm;^{19–21} this peak is associated with the template-bound conformation of the 10-porphyrin ring. At quinuclidine concentrations of 30–300 mM (Figure 3c), the central **T6** unit is displaced causing disappearance of the distinctive three-finger pattern in the Q band of the 6-porphyrin ring component. All these spectral changes reflect the greater flexibility, and wider distribution of porphyrin-porphyrin dihedral angles, upon template removal.

Analysis of the denaturation binding isotherms^{13c,17,21} (Figure 3) reveals that the association constants for binding of **T4** and **T6** to **b-P14** to form **b-P14·T6·(T4)₂** are $(1.8 \pm 0.2) \times 10^{22} \text{ M}^{-1}$ for **T4** and $(5.5 \pm 1.2) \times 10^{37} \text{ M}^{-1}$ for **T6**, in toluene at 298 K. This analysis assumes that the denaturation processes for each template (**T4** and **T6**) are essentially all-or-nothing two state equilibria (i.e. intermediate partially denatured species do not build up to high concentrations); this assumption is supported by the isosbestic nature of the UV-vis-NIR titration and the good fits of the curves to the calculated isotherm for a two-state equilibrium (Figure 3b,c). The binding strength of **T6** is roughly an order of magnitude stronger in the ball than in a comparable 6-ring system,¹⁷ which can be attributed to the conformational preorganization provided by the 10-porphyrin ring.

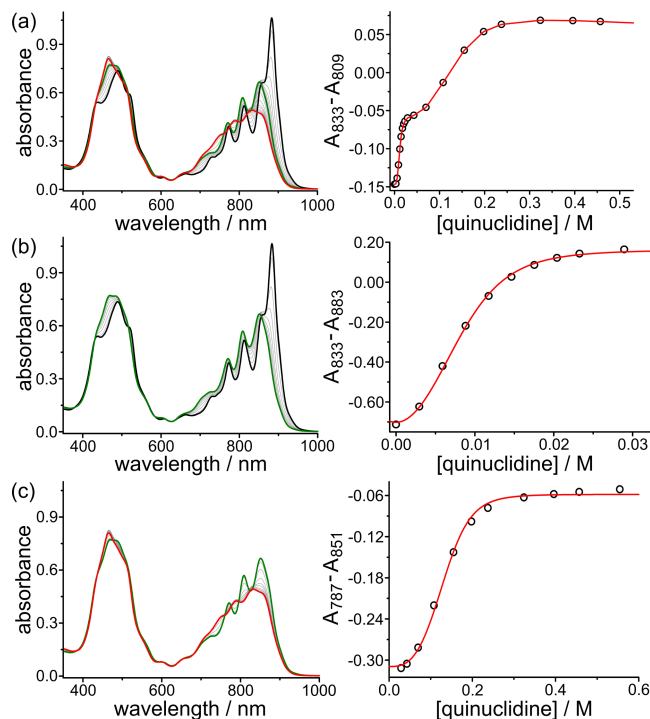


Figure 3. UV-vis-NIR denaturation titration of the **b-P14·T6·(T4)₂** complex with quinuclidine (toluene, 298 K). (a) Full titration, **T4** is displaced first (black to green) followed by the displacement of **T6** (green to red). The two phases of the titration are shown in (b) and (c), with the experimental (black circles) and calculated isotherms (red lines) shown on the right.

The UV-vis-NIR absorption and fluorescence spectra of **b-P14·T6·(T4)₂** and **b-P14** are compared in Figure 4. The more rigid conformation of the template-bound ball is reflected by its sharper and more red-shifted absorption and emission spectra.¹⁹⁻²¹ Fluorescence up-conversion spectroscopy experiments reveal that the excited states delocalize across the two perpendicular ring planes in the nanoball on an ultrafast timescale. The fluorescence anisotropy dynamics of **b-P14**, with and without templates, are shown in Figure 4c. The template complex, **b-P14·T6·(T4)₂** (black points), exhibits a constant and very low anisotropy ($\gamma = 0.02$) over the time range of this experiment (0–10 ps; time resolution 0.3 ps), showing that the excited state delocalizes in three dimensions within 0.3 ps. In contrast, **b-P14·T6** (green points) and **b-P14** (red points) show a fast initial drop in anisotropy within the first 5 ps from $\gamma = 0.1$ towards $\gamma \approx 0$. This fast depolarization resembles the anisotropy decay in porphyrin nanorings with >24 porphyrin units.¹⁵ The initial anisotropy of 0.1 suggests that upon excitation, an exciton is delocalized over a full ring and that both absorption and emission transition dipoles are polarized in the ring plane. After ultrafast relaxation, the exciton localizes and migrates rapidly around the entire porphyrin nanoball. Contributions from emission components polarized in both planes thus result in an anisotropy close to zero. Without the templates, exciton migration is slower, resulting in the observed anisotropy decays, whereas in **b-P14·T6·(T4)₂** the anisotropy decay is faster than the time resolution of the experiment. This behavior is very different from that of the nanorings **c-P6·T6**, **c-P6** and **c-P10**, which exhibit anisotropies of near 0.1 (remaining constant during 10 ps after excitation), in agreement

with theoretical predictions for an excited state that is delocalized over a 2D ring.^{15,16,18,20,22}

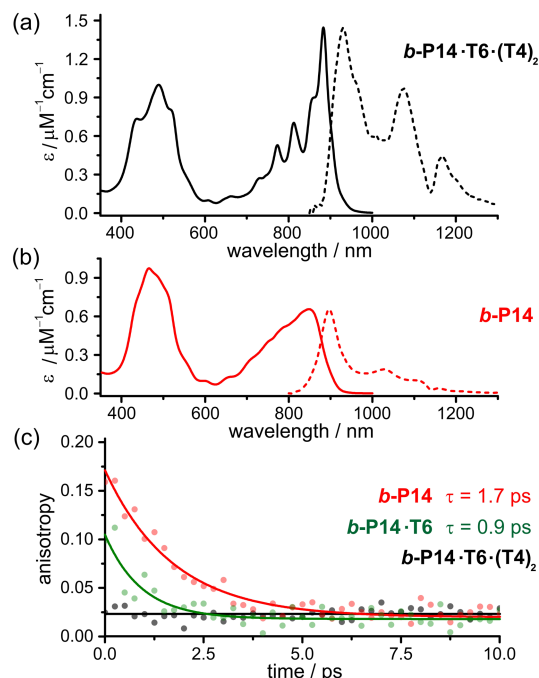


Figure 4. UV-vis-NIR absorption (solid line) and fluorescence (dashed line) spectra of (a) **b-P14·T6·(T4)₂** and (b) **b-P14** (toluene, 298 K). The dip at 1130–1170 nm corresponds to solvent absorption. (c) First order fits of the fluorescence anisotropy decay. Solutions of **b-P14·T6·(T4)₂** (black), **b-P14·T6** (green) and **b-P14** (red) were excited at 820 nm and emission was detected at 950 nm.

The prolate shape of **b-P14·T6·(T4)₂** and **b-P14** is reminiscent of C_{70} fullerene. While C_{60} fullerene has complete polarization memory loss with zero fluorescence anisotropy, C_{70} displays excitation-wavelength dependent anisotropy values ranging between -0.2 and 0.1 , because emission polarized in the x - y plane is energetically favorable as a result of geometrical deformation.²³ However, **b-P14·T6·(T4)₂** and **b-P14** do not display any significant changes in anisotropy with excitation wavelength in the range of 760–880 nm, probably because the absorption features of their two constituent rings broadly overlap and the emission may be polarized in both ring planes. DFT calculations (B3LYP/6-31G*) indicate that the HOMO of **b-P14** is distributed over the entire π -system (Figure 5), albeit with a higher density on the 6-ring and particularly the tetraalkynylporphyrins. The LUMO and LUMO+1 are located exclusively on the 6-ring and the 10-ring, respectively, with nearly identical energies. Time-dependent DFT calculations (B3LYP/6-31G*) were carried out to model the electronic excited states of **b-P14**. Natural transition orbital plots (Tables S5–S13) show that the two lowest energy singlet excited states (S_1 and S_2) are mainly localized in the 6-porphyrin and 10-porphyrin ring components, respectively. The transitions from S_0 – S_1 and S_0 – S_2 are dipole-forbidden ($f = 0$), whereas transitions to S_3 , S_4 and S_5 are allowed ($f = 0.13$, 5.12 and 0.83 , respectively) and these excited states are distributed over all 14 porphyrins. These calculations are in line with the observation that excitation delocalizes between the two rings within

the ball on a timescale of less than 300 fs. The dimensions of the calculated geometry of **b-P14** are 52.7 Å, 27.6 Å and 23.9 Å (measured as the diameter along the D_{2h} symmetry axes, without including aryl side chains).

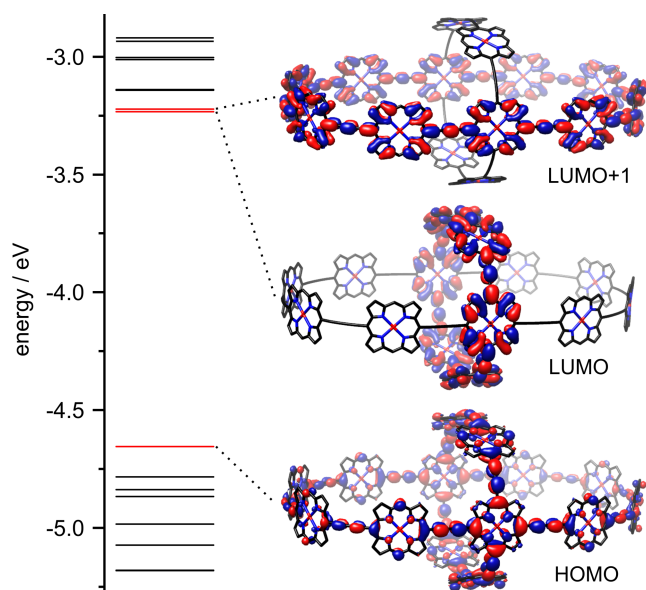


Figure 5. LUMO+1 (−3.22 eV), LUMO (−3.23 eV) and HOMO (−4.66 eV) of **b-P14** calculated at the B3LYP/6-31G* level of theory with Grimme's dispersion correction, GD3, shown at a density isovalue of 0.008 a.u. together with their corresponding energy levels. Aryl groups were replaced with hydrogen atoms to simplify the calculations.

In summary, we have synthesized a fully π -conjugated three-dimensional 14-porphyrin nanoball by a template directed approach. UV-vis-NIR titrations show that the templates bound within the cavity can be removed successively by the addition of a competing ligand. Fluorescence upconversion spectroscopy reveals ultrafast electronic delocalization between the two perpendicular ring planes in the porphyrin ball. The fluorescence anisotropy approaches zero, indicating that excitation rapidly migrates between the two ring components of the nanoball.

ASSOCIATED CONTENT

Supporting Information. Synthetic procedures, characterization data, binding studies, NMR assignments, fluorescence spectroscopy and calculated geometries. The Supporting Information is available free of charge at <http://pubs.acs.org>.

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