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Noncovalent Synthesis of Self-assembled Nanotubes through Decoupled Hierarchical Cooperative Processes

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ABSTRACT: Due to their wide number of biological functions and potential applications, self-assembled nanotubes constitute highly relevant targets in noncovalent synthesis. Herein, we introduce a novel approach to produce supramolecular nanotubes with defined inner and outer diameters from rigid rod-like monomers programmed with complementary nucleobases through two distinct, decoupled cooperative processes of different hierarchy and acting in orthogonal directions: chelate cooperativity, responsible for the formation of robust Watson–Crick H-bonded cyclic tetramers, and nucleation-growth cooperative polymerization.

INTRODUCTION
Noncovalent synthesis aims at the production of well-defined nanostructures, often mimicking those found in the natural world, and relies both on noncovalent interactions and cooperative effects between chemically programmed molecules.3–5 One of the most appealing noncovalent synthetic targets are self-assembled nanotubes, due to their nanoscale dimensions, with inner cavities in the attoliter regime, their large variety of biological functions, illustrated by tube-forming proteins like tubulin or gramicidin,4,5 and their potential applications6–12 in chemico- and size-selective encapsulation, transport, or catalysis. These cylindrical architectures can be formed from rationally designed molecules of different nature and through manifold approaches, that in many cases encompass multi-step pathways of increasing hierarchy.13–17

One of these approaches involves the combination of two hierarchical processes that are strongly coupled (Figure 1a; left). First, the association of dendron-shaped molecules with self-complementary H-bonding faces leads to rosette-like macrocycles and, as soon as these transient intermediates are formed, a stacking process is triggered in the orthogonal direction.18–21 The most popular building blocks employed here contain heterocyclic head groups like guanine,22,23 cyanuric/barbituric acid,24,25 melamine/diaminotriazine,26 their combination,27,28 or pyrimido[4,5-d]pyrimidine derivatives.29,30 This strategy can be implemented in aqueous or organic media with molecules carrying diverse external functions. However, it affords little control on pore dimensions and function, and the resulting nanotubes typically have small pores, able in the best case to accommodate ions or very small molecules.

Here we introduce a related strategy for the synthesis of self-assembled nanotubes that is based on a novel molecular design, involving rigid rod-like monomers that interact through their edges in a defined 90° geometry (Figure 1a; right). Specifically, our monomer structure (GC; Figure 1b) comprises complementary guanine (G) and cytosine (C) nucleobases linked by linear, π-conjugated spacers. The bases are also equipped with benzylic wedges substituted with long alkyl tails, so as to enhance solubility in apolar solvents, and with a peripheral amide group. This molecular design allowed to control self-assembly through two orthogonal cooperative events of different hierarchy, as shown in Figure 1c, that can be now efficiently decoupled. First, chelate cooperativity is responsible for the formation of unstrained cyclic tetramers through G:C Watson–Crick H-bonding interactions,3 as demonstrated previously with related molecules bearing bulky riboses.32–35 After these planar macrocycles are generated quantitatively, a polymerization process, that occurs through a nucleation-growth mechanism and is guided by π–π and H-bonding interactions along the stacking axis, can be triggered by a change in experimental conditions. Hence, both cooperative processes are independent and self-assembly can be controlled either at the cycle stage or at the polymer stage. Since the “supramonomer” is cyclic in nature, the resulting polymer is tubular, with precisely defined inner and outer diameters.
RESULTS AND DISCUSSION

Monomer-Cyclic Tetramer Equilibrium. The association state of GC is, as expected, strongly dependent on the H-bonding competing ability of the solvent environment. Highly polar solvents like DMF or DMAC are able to produce full dissociation at concentrations below 10^{-3} M. Moving to solvents of intermediate polarity,^{23} like THF, allowed us to monitor the monomer (GC)-cyclic tetramer (cGC) equilibrium at concentrations between 10^{-3} - 10^{-5} M by 1H NMR, absorption, CD and emission spectroscopies. Figure 2a shows the evolution of the GC aromatic 1H signals as a function of temperature at 10^{-4} M in THF-D_{8} (see Figures S1A-B for additional 1H NMR spectra at different temperatures and concentrations). As the temperature decreases from 323 to 268 K, the GC monomer signals (rods) decrease in intensity at the expense of the cGC signals (squares). H-bonding between G and C nucleobases was confirmed by the characteristic downfield shift of the G-amine and the C-amine proton signals to 13.2 and 10.3 ppm, respectively, as well as by NOESY cross-peaks between these two H-bonded protons (Figures 2b and S1C). It is interesting to note that both GC and cGC species are in slow exchange at the NMR timescale and that the shape and position of the 1H signals do not change much with temperature or concentration,^{23} which underlines the slow dynamics and all-or-nothing behavior of this macrocyclization process.

The 1H NMR data can be overlapped with optical spectroscopy techniques, and Figures 2c-e display, respectively, the absorption, CD and emission changes that occur at 10^{-4} M in THF as a function of temperature. Upon cooling, GC cyclotetramerization is promoted and a single, rigid conformation, that maintains the two Watson-Crick edges pointing in the same direction, is fixed.^{23-25} This is typically monitored by: 1) a red-shift in absorbance with a characteristic new maximum at 420 nm; 2) a decrease in emission intensity and a marked red-shift from 439 nm (GC) to 525 nm (cGC); and 3) a CD Cotton effect, positive for cS-GC and negative for cR-GC, with zero-crossing at 434 nm. Figures S1D-E contain additional temperature- and concentration-dependent spectra in THF.^{23-25}

Such spectroscopy changes can be used to determine the molar fraction of GC molecules associated as cGC, (γ), as a function of temperature (Figure 2f) or concentration (Figure 2g). The cyclization trends obtained from NMR and CD measurements at the same concentrations show an excellent agreement. Fitting the NMR data to a cyclotetramerization process afforded the corresponding equilibrium constants (K_GC; see S.1.), from which the effective molarity (EM) values could be calculated as K_{GC} = EM · K_{GC} in (Table 1a and S1), where K_{GC} is the reference association constant between nucleobases in THF.^{35}

Table 1. Thermodynamic parameters calculated for GC upon (a) cyclotetramerization, (b) polymerization as a function of solvent composition, and (c) polymerization as a function of temperature.

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<tr>
<td>K_{Gc}^{[a]}/M^3</td>
<td>K_{i}^{[b]}/M^3</td>
<td>EM^{[d]}/M</td>
<td>ΔH^{[eq]}/kJ·mol^{-1}</td>
<td>ΔS^{[eq]}/J·mol^{-1}·K^{-1}</td>
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<tr>
<td>1.1·10^{3}</td>
<td>3.4·10^{3}</td>
<td>2.3</td>
<td>-173.4</td>
<td>-311.2</td>
<td></td>
</tr>
<tr>
<td>b</td>
<td></td>
<td></td>
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<td></td>
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<tr>
<td>ΔG^{[eq]}/kJ·mol^{-1}</td>
<td>m^{[h]}/kJ·mol^{-1}</td>
<td>σ^{[k]}</td>
<td></td>
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<tr>
<td>-34.9 ± 0.5</td>
<td>90 ± 9</td>
<td>0.19 ± 0.03</td>
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<td>c</td>
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<tr>
<td>K_{i}^{[d]}/M^4</td>
<td>K_{i}^{[d]}/M^4</td>
<td>σ^{[l]}</td>
<td>ΔH^{[eq]}/kJ·mol^{-1}</td>
<td>ΔS^{[eq]}/J·mol^{-1}·K^{-1}</td>
<td></td>
</tr>
<tr>
<td>4.02·10^{4}</td>
<td>1.34·10^{5}</td>
<td>3.0·10^{-3}</td>
<td>-166.6 ± 3</td>
<td>-290 ± 10</td>
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[a] Reference GC association constant.  
[b] Cyclotetramerization constant.  
[c] Effective molarity calculated as: EM = K_T / K_{GC}.  
[d] Cyclotetramerization enthalpy and σentropy.  
[e] Gibbs free energy.  
[f] m parameter and [g] degree of cooperativity of the polymerization process observed by increasing V_{sec}.  
[h] Nucleation and [i] elongation constants.  
[j] Cooperativity factor, and elongation [k] enthalpy and [l] entropy of the polymerization process observed by decreasing T.

Using these calculated EM and K_{GC} values, we built speciation curves in which monomer, cyclic tetramer and small open oligomers were included. These curves, in which the relative distribution of species is represented as a function of overall concentration, illustrate graphically the all-or-nothing process observed in solution, where mostly GC monomer (green) and cGC macrocycles (purple) are in equilibrium, and the participation of small open
Figure 2. Cyclotetramerization. Changes observed in the (a) $^1$H NMR, (c) absorption (d) CD and (e) emission spectra as a function of temperature for 1.0·10$^{-4}$ M solutions of GC in THF-D$_8$ (a) or R-GC in THF (c-e). (b) NOESY NMR spectrum at $\tau_{nm} = 0$ ms of GC ([GC] = 2.0·10$^{-3}$ M, 298 K) showing cross-peaks between H-bonded G and C protons. (f-g) Representation of the molar fraction of GC molecules associated as cGC$_4$ cycles ($\chi_4$) calculated by $^1$H NMR (triangles) or CD (squares) as a function of $T$ the temperature at different concentrations or (g) the total concentration at 298 K. In (g) the relative abundance of GC, cGC$_4$ and open oligomers (like GC$_2$) was simulated (solid lines) using $K_{GC} = 1.1·10^3$ M$^{-2}$ and $EM = 2$ M.

As $V_s$ is increased, the spectroscopic features that characterize the cyclotetramerization process, as described above in THF and toluene, are first observed. For instance, upon monitoring this process by $^1$H NMR at a 3.0·10$^{-4}$ M concentration (Figure 3a), the residual monomer signals first disappear and the cGC$_4$ macrocycle is formed quantitatively. When this process was recorded by CD, absorption or emission spectroscopy at similar or lower concentrations (Figures 3b-d; green to purple spectra), we also saw the typical features of cGC$_4$ formation: a CD signal appears, a red-shifted absorption shoulder, and also a red shift and decrease in emission intensity as the heptane content is increased. Further increasing $V_s$ up to 0.6-0.9 (depending on concentration; see below), produced no spectroscopic change, indicating that the cGC$_4$ species is quantitatively stabilized in solution.

However, at even higher $V_s$, a distinct aggregation process is clearly detected (purple to pink spectra). In $^1$H NMR, very minor chemical shifts ($\Delta ppm < 0.05$) are observed and then the cGC$_4$ proton signals start to broaden and eventually disappear, which is characteristic of the formation of large aggregates. Unfortunately, as shown in Figure 3a, the moment the macrocycles start to aggregate, the proton signal of the peripheral amide also broadens and disappears, which prevented us to monitor intermolecular H-bonding interactions.

Turning to optical spectroscopy, an additional absorption red-shift to 425 nm is observed, while emission is further quenched and slightly blue-shifted. However, the most remarkable changes were detected in the CD measurements: at high $V_s$, the CD features evolve with clear isosbestic points to a new signal exhibiting a Cotton effect at 428 nm, that is positive for S-GC and
Figure 3. Supramolecular Polymerization. (a–d) Changes observed in the: (a) aromatic region of the 1H NMR spectra by increasing the volume fraction of cyclohexane-$D_{12}$ ($V_a$) in mixtures with THF-$D_8$ ([GC] $= 3.0 \times 10^{-4}$ M); (b) absorption (c) CD and (d) emission spectra as a function of the volume fraction of heptane ($V_e$) in mixtures with THF ([S-GC] $= 3.0 \times 10^{-3}$ M). (e–g) Changes in the S-GC CD signal at 429 nm at several concentrations as a function of (e) volume fraction of heptane ($V_e$) or (f) volume fraction of THF ($V_{THF}$) at 298 K or (g) temperature at $V_e = 0.99$ ($a_T$ = fraction of cyclotetramers, $a_N$ = fraction of nanotubes, $a_{agg}$ = fraction of aggregated species).

negative for R-GC (Figure S3E). Due to the similarity of these final spectroscopic features at high heptane contents to the ones measured for the initial dispersions in alkanes, this second sharp transition was attributed to a supramolecular polymerization process to yield (cGC$_4$)$_n$, which would be driven by $\pi$–$\pi$ stacking interactions between the large $\pi$-conjugated surface generated upon cyclization and by H-bonding interactions between the four peripheral amides (see Figure 1c). As a matter of fact, related G-C monomers lacking these amide groups at the C base did not undergo this second self-assembly process, and remained associated as cGC$_4$ even in pure heptane (see Figure S3F).

Both self-assembly stages, cyclotetramerization and polymerization, are displayed in Figure 3e by monitoring the evolution of the GC CD features as a function of $V_a$ at different concentrations (see also Figure S3G). It is clear that both processes are strongly dependent on the overall concentration. For instance, at $3.0 \times 10^{-4}$ M, we mainly start with a GC-cGC$_4$ equilibrium that is shifted to the cyclic species up to $V_a = 0.3$, while the polymerization transition is detected above $V_a = 0.65$. At $1.0 \times 10^{-5}$ M, in contrast, we observe the whole two-step self-assembly process: from the monomer in pure THF, to the cGC$_4$ cycle within the $V_a = 0.5$–0.9 plateau, and then to the polymer above $V_a = 0.095$.

Such cGC$_4$-(cGC$_4$)$_n$ transitions as a function of the solvent composition were fitted to an extended nucleation-elongation model developed by de Greef, Meijer and co-workers (Figures 3f and S3G, Tables 1b and S2), which allows the calculation of the Gibbs free energy gain upon monomer addition ($\Delta G_m$), the m parameter which characterizes the ability of the good solvent to associate with the monomer thereby destabilising the supramolecular aggregated species, as well as the equilibrium nucleation ($K_n$) and elongation ($K_e$) constants, whose ratio defines the cooperative parameter ($\sigma$). A detailed explanation is provided in section S3 of the Supporting Information.

We then turned to temperature-dependent studies in order to obtain complementary thermodynamic parameters for this polymerization process. We fixed solvent composition at $V_a = 0.97$ and analyzed the CD changes along cooling cycles within the 249–268 K range (Figures 3g and S3H). The non-sigmoidal curves obtained at three different concentrations could be fitted again to a cooperative nucleation–elongation model, in which the polymerization process can be divided in a nucleation and an elongation phase. The magnitudes $T_c$ (elongation temperature), $K_n$ and $K_e$ (nucleation and elongation constants), $\sigma$ (cooperativity factor), $\Delta H_n$ and $\Delta H_e$ (nucleation and elongation enthalpies), and $\Delta S^o$ (polymerization entropy) can be obtained from a non-linear least-square analysis of the experimental melting curves (Tables 1c and S3; see also the Supporting Information, section S3).

The degrees of cooperativity calculated in these experiments are lower than those determined before as a function of solvent composition, but this is not surprising, since each kind of analysis and experimental conditions differ substantially.

Characterization of the Self-assembled Nanotubes.

We then proceeded to characterize the final GC aggregates obtained after the polymerization process, in order to confirm their dimensions and tubular nature. However, solution measurements, like dynamic light scattering (DLS) or
small angle X-ray scattering (SAXS) at high heptane contents faced a challenging scenario, because of the evolution of a third hierarchical assembly level that involves the formation of large bundled agglomerates. In fact, after the polymerization process is complete at high \( V_a \), and in a timescale that depends on concentration (ranging from a few hours to several days), we observed that the solution became turbid and a precipitate appeared. When monitoring this process spectrosopically with time, the shape of the absorption and CD spectra did not change, but a gradual loss in absorption intensity and a baseline rise, attributed to scattering, was noted. Once this final aggregation state is reached, we noted quite marked kinetic effects, and the \((cGC)_n\) dispersions are rather inert to disaggregation, so solubilization by heating or dilution required unusually long times.

Despite these experimental difficulties, DLS measurements (Figure S4A) performed at several concentrations and THF:heptane ratios, thus targeting cycle \((V_a = 0.4)\) and polymer \((V_a = 0.99)\) states, were in agreement with the formation of large anisotropic aggregates at high alkane contents. On the other hand, SAXS experiments, obtained immediately after sample preparation at high \( V_a \), provided an indication of cylindrical organization of large particles. As shown in Figure 4a (black curve) the position of the 1st peak (marked with an arrow) indicates a regular dimension of 3.8 nm, attributed to the distance between bundled cylinder centers, while according to the vague second peak the packing appears hexagonal. However, these SAXS measurements, which required relatively high concentrations, were particularly sensitive to the precipitation process, and the patterns recorded evolved with time (compare black and blue curves in Figure 4a) eventually providing a picture of the smaller particles that remained in solution. In any case, fitting of the data acquired at different time lapses to a cylindrical core-shell model was consistent with a cylinder diameter of 4 ± 1 nm and a core diameter of about 1 nm.

Finally, we analyzed \((S/-R)-GC\) samples, drop-casted from diluted solutions with high heptane contents \((V_a > 0.9)\), by different microscopy techniques (SEM, AFM and TEM; see Section S4.3). SEM measurements (Figures 4b and S4C) confirmed the formation of networks of large fibrillar aggregates. A closer analysis by TEM (Figures 4c-d) revealed that these aggregates consist of heavily bundled longitudinal objects with a measured diameter of 3.9 ± 0.7 nm, which coincides with the hard aromatic section of the cyclic tetramer (Figure 4e). Organization into aligned nanotube bundles increases as the time in solution before deposition becomes longer (please, compare Figures 4d/S4F and 4e/S4G), and is plausibly driven by van der Waals interactions between interdigitated peripheral chains.

**ASSOCIATED CONTENT**

**Supporting Information.** Experimental details, compound synthesis and characterizations, additional spectroscopy and microscopy data and analysis. This material is available free of charge via the Internet at http://pubs.acs.org.
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Notes
The authors declare no competing financial interests.

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REFERENCES

(33) Romero-Pérez, S.; Camacho-García, J.; Montoro-García, C.; López-Pérez, A. M.; Sanz, A.; Mayoral, M. J.; González-


