

Il parco archeologico delle Torri Palatine

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Self-Sorted, Alternate/Random and Block Supramolecular Co-polymers via Sequence Controlled, Multicomponent Self-Assembly

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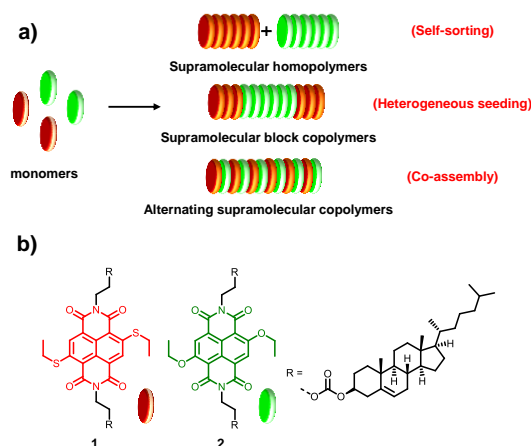
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ABSTRACT: Multicomponent supramolecular co-polymerization promises to construct complex nanostructures with emergent properties. However even with two monomeric components, various possible outcomes such as self-sorted supramolecular homopolymers, a random/alternating supramolecular co-polymer or a complex supramolecular block copolymer can occur, determined by their intermolecular interactions and monomer exchange dynamics and hence structural prediction is extremely challenging. Herein, we target this challenge and demonstrate unprecedented two-component sequence controlled supramolecular co-polymerization by manipulating thermodynamic and kinetic routes in the pathway complexity of self-assembly of the constitutive monomers. Extensive molecular dynamics simulations provided useful mechanistic insights into the monomer exchange rates and free energy of interactions between the monomers that dictate the self-assembly pathway and sequence. Fluorescent nature of core-substituted naphthalene diimide monomers has been further utilized to characterize the three sequences via Structured Illumination Microscopy (SIM).

Introduction

Sequence controlled polymerization is a vital phenomenon observed in natural systems. The genetic information is encoded by nucleobase sequences in DNA and protein's structure and function is determined by its amino acid sequence.¹ This sequence-control has been implemented in covalent polymers via elegant synthetic strategies resulting in emergent structural properties and multitude of applications.² On the other hand, supramolecular polymers, composed of dynamic non-covalent interactions between monomers, with predictive sequence would provide an opportunity to dynamically rearrange/reconfigure the composition similar to natural systems.³ However, supramolecular polymerization with even two components is challenging and the sequence outcome could be either self-sorted supramolecular homopolymers (AAA and BBB), a random or alternate supramolecular co-polymer (ABABAB) or complex supramolecular block co-polymer (AABBAA) whose sequence is determined by the free energy of intermolecular (homo AA, BB vs. hetero AB) interactions, monomer structure and monomer exchange dynamics.



Scheme 1. Two-component supramolecular polymerization. (a) Schematic representation of two-component supramolecular polymerization into self-sorted supramolecular homopolymers, heterogeneously seeded supramolecular block copolymers and co-assembled supramolecular alternating copolymers. (b) Molecular structures of **1** and **2** used for the two-component supramolecular polymerization.

In this study, we have demonstrated an unprecedented two-component sequence controlled supramolecular co-

polymerization into the three unique sequences by exploiting the delicate balance between the kinetic and thermodynamic parameters of self-assembly. We further manipulate these parameters to illustrate dynamic reconfiguration of these sequences into one another. Self-sorted supramolecular polymers have been achieved by high homorecognition (AA, BB > AB) between monomers, designed with structural mismatch or low monomer exchange dynamics.^{4,7} Adams and coworkers reported pH-programmed self-sorting of gelator monomers with different pK_a.⁵ On the other hand structural mismatch between donor and acceptor chromophores leading to orthogonal supramolecular homopolymers have been reported.⁶ We have reported that chirality driven mismatch can also be used to construct self-sorted systems even with structurally similar monomers.⁷ In contrast, alternating supramolecular co-polymers require stronger heterorecognition between monomers (AA, BB < AB) and has been preferentially attained by strong hetero-complementary interactions such as electrostatic and donor-acceptor charge-transfer interactions. Charge-transfer supramolecular polymers reported by our group, Zhang and others are excellent examples of alternating supramolecular polymers with electronically complementary donor and acceptor monomers.⁸ Besenius and coworkers employed complementary electrostatic interactions from lysine and glutamic acid residues to obtain alternating arrangement at neutral pH.⁹ Oren and coworkers has utilized strong charge-transfer ternary complexes with cucurbit[8]uril to manifest alternate host-guest supramolecular polymer.¹⁰ These self-sorted or alternating supramolecular co-polymers have been achieved mostly under thermodynamic conditions, where the interaction energies between the monomers determines the sequence.

However supramolecular block co-polymer arrangement has been a grand challenge in the field.¹¹ This is due to specific requirements such as structural similarity between monomers, low exchange dynamics and delicate balance between homo- and hetero-recognition between monomers. However, mechanistic understanding¹² and development of living supramolecular polymerization has aided in addressing this challenge.^{11b,13,14} Meijer and coworkers have constructed supramolecular block co-polymers under thermodynamic control with monomers exhibiting reactivity ratio difference analogous to the synthesis of block co-polymers via chain-growth polymerization in covalent polymers.¹⁵ In another approach Yagai and coworkers have demonstrated co-assembly between two monomers under thermodynamic control to result in a supramolecular co-polymer, where the monomer composition varies gradually during the polymerization and along the polymer chain.¹⁶ Supramolecular block co-polymerization under kinetic control requires sequential addition of dormant monomers undergoing nucleation-elongation growth. However, this is challenging, as minor difference in structure alters the energy landscape significantly. Hence, Sugiyasu and Takeuchi's group¹⁷ achieved block sequence by crystallization-driven self-assembly approach similar to Manners and coworkers elegantly established in block co-

polymer assemblies.¹⁸ Aida and coworkers synthesized p-n junction via seeded assembly between electronically different chromophores.¹⁹ Recently, Würthner and coworkers presented the synthesis of supramolecular co-polymers of core-substituted perylene bisimide using kinetically control seeded supramolecular polymerization which is more closer to the conventional multicomponent living polymerization leading to block co-polymers.²⁰

So far, the three sequences have been synthesized separately owing to their orthogonal and unique requisites. Thus, achieving the three sequences from the same monomer pair and their dynamic reconfiguration is still elusive. In order to achieve precise sequence control, it is necessary to move through the complete energy landscape of self-assembly via thermodynamic and kinetic routes. To realize this, herein we introduce an unprecedented two-component supramolecular polymerization with sequence control to access self-sorted, alternate and block co-polymer arrangement via pathway complexity (Scheme 1a).²¹

To attempt this, we chose our previously studied molecular design based on carbonate cholesterol appended naphthalene diimide²² which due to long-range dipole-dipole interactions followed a cooperative mechanism, necessary for seeded supramolecular polymerization. Secondly, to have minimum structural perturbation and to probe the polymerization process, we symmetrically disubstituted the naphthalene diimide core with ethane thiol (**1**) and ethoxy (**2**) groups to give red and green emissive dyes respectively (Scheme 1b). These molecules have unique optoelectronic properties as well as different absorption and emission profiles aiding to the orthogonal probing of their self-assembly via spectroscopic and super-resolved fluorescence microscopy characterization (SIM).²³

Supramolecular Polymerization of **1** and **2**

Core-substituted NDI (cNDI) monomers **1** and **2** have distinct absorption (450-600 nm and 400-500 nm, respectively) and emission profiles (570-750 nm and 450-570 nm, respectively), which allowed the orthogonal probing of its self-assembly characteristics (Figure S1-S3). Supramolecular polymerization of these monomers could be induced in the toluene-MCH solvent mixture (5 % toluene/MCH) and was probed spectroscopically. **1** and **2** formed red ($\lambda_{\text{max}} = 625$ nm) and green ($\lambda_{\text{max}} = 480$ nm) emissive, one-dimensional supramolecular polymers respectively as evident from the structured illumination microscopy (SIM) and transmission electron microscopy (TEM) images (Figure 1a,b, S3). Detailed absorption, steady-state and time-resolved fluorescence and excitation spectroscopy studies further revealed J-type slipped organization of chromophores in these stacks. The monomer **2**, in addition to its green emission from self-assembled chromophores, also showed a weak, static excimer emission in the red wavelength region with a maximum at 634 nm (Figure S2c,d). Further, both cNDI chromophores are organized in a helical manner, as evident from the exciton coupled circular dichroism spectra (Figure S1c, S2b), biased by the peripheral chiral cholesterol groups.¹⁹ The good spectral overlap

between the absorption spectrum of **1** and emission spectrum of **2**, further enables the probing of the co-polymerization process via Förster Resonance Energy Transfer (FRET) (Figure S4).

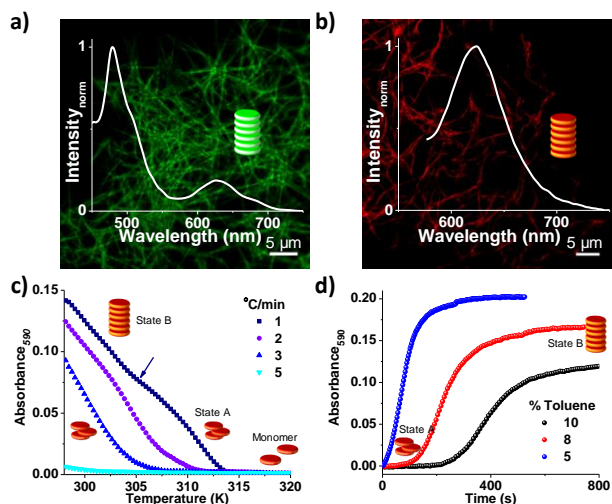


Figure 1. Self-assembly of **2** and **1**. (a) and (b) Emission spectra of **2** ($\lambda_{ex} = 442$ nm) and **1** ($\lambda_{ex} = 480$ nm) showing fluorescent assembled state with corresponding SIM image showing red (channel I) and green (channel II) fluorescent one-dimensional fibers (5 % toluene/MCH). (c) Dependence of pathway complexity of **1** on cooling rate, an increase in cooling rate leads to the preferential formation of kinetic trapped **State A** over thermodynamically stable **State B** (8 % toluene/MCH). (d) Time-dependent absorption changes at 590 nm of a fast cooled (5 K/min) solution of **1** from 363 K to 298 K depicting self-assembly of **1** into **State B** via sigmoidal nucleation-elongation kinetics. ($[1] = [2] = 5 \times 10^{-5}$ M, $l = 10$ mm)

Pathway complexity of **1** and **2**:

Temperature and time-dependent spectroscopic analyses of **1** and **2** in toluene/MCH solvent mixture provided mechanistic insights into the supramolecular polymerization process. Interestingly, temperature-dependent cooling curves of **1** (363 K to 298 K) recorded at 1 K/min and monitored at 590 nm absorption band (5 % toluene/MCH and 5×10^{-5} M), depicted a two-state transition corresponding to a gradual conversion of monomer to an intermediate self-assembled **State A** finally to the self-assembled **State B**, corresponding to the supramolecular polymeric state (Figure 1c, S5). **State A** was stabilized more in higher toluene percentages (10 and 8 % of toluene) and further, a fast cooling (≥ 5 K/min, ≥ 5 % toluene/MCH) resulted in exclusive formation of **State A** (Figure 1c, S6). Interestingly, self-assembled **State A** transformed to final **State B** with time and time-dependent absorbance changes monitored at 590 nm showed a non-linear, sigmoidal transformation with a lag phase and nucleation-elongation growth mechanism (Figure 1d, S7-S10). An increasing percentage of toluene followed an enhancement in t_{lag} (lag time) and t_{50} (time required for completion of 50 % of the process) due to more stabilization of **State A** at higher percentage of toluene. Also, the presence of hysteresis between the cooling and heating profiles (Figure S6b) indicated **State A** to be a non-

equilibrium metastable state *en route* to the thermodynamically stable **State B** of the supramolecular polymer.

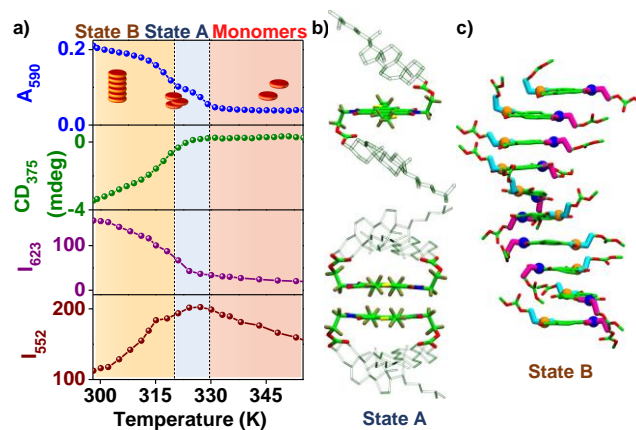


Figure 2. Characterization of metastable **State A** and thermodynamically stable **State B**. (a) Comparison of absorbance at 590 nm, CD at 375 nm and emission changes ($\lambda_{ex} = 480$ nm) at 623 nm and 552 nm against temperature. Absorbance and emission at 552 nm show two-step transition. The pink regime indicates molecularly dissolved state, blue region shows the region where **State A** occurs and the orange region shows conversion to final thermodynamically stable **State B**. (b) Side view of the metastable locked chair conformation (top) and side view of the locked dimer conformation (bottom) optimized at the PM6 // BLYP-D3/DZVP level of theory (**State A**) and (c) snapshot of chair decamer obtained from MD simulations (**State B**). Cholesterol hydrogens are not shown for clarity. ($[1] = [2] = 5 \times 10^{-5}$ M, 5 % toluene/MCH)

To get further insight into the pathway selectivity in the self-organization of **1** and the metastable **State A**, we further performed concentration and temperature-dependent spectroscopic studies and Molecular Dynamics (MD) simulations. The formation of **State A** resulted in insignificant changes in absorption and emission spectra compared to monomeric state and remains CD inactive, suggesting the formation of less-ordered small aggregates or a conformationally locked, metastable states for the monomers of **1** prior to the nucleation process (Figure 2a, S5). The transition to **State B** was accompanied with a significant broadening of the absorption spectrum and appearance of the new red-shifted aggregate absorption band at 590 nm and the emission band at 625 nm implying the formation of an extended self-assembled structure with exciton-coupled π -stacked chromophores. Gas phase quantum chemical calculations and all-atom molecular dynamic simulations with explicit solvation suggested the formation of locked monomers and dimers as **State A**, which convert to extended supramolecular polymers (**State B**) via conformational changes in the monomer structure (Figure 2b-c). Interestingly, molecule **2** also shows a similar hysteresis and kinetically controlled nucleation-elongation growth at higher solvent percentages of good solvent, toluene (>30 % toluene/MCH) (Figure S12-S14). A decrease in lag time with an increase in the concentration of monomers designates an on-pathway formation of **State A** (Figure S9).

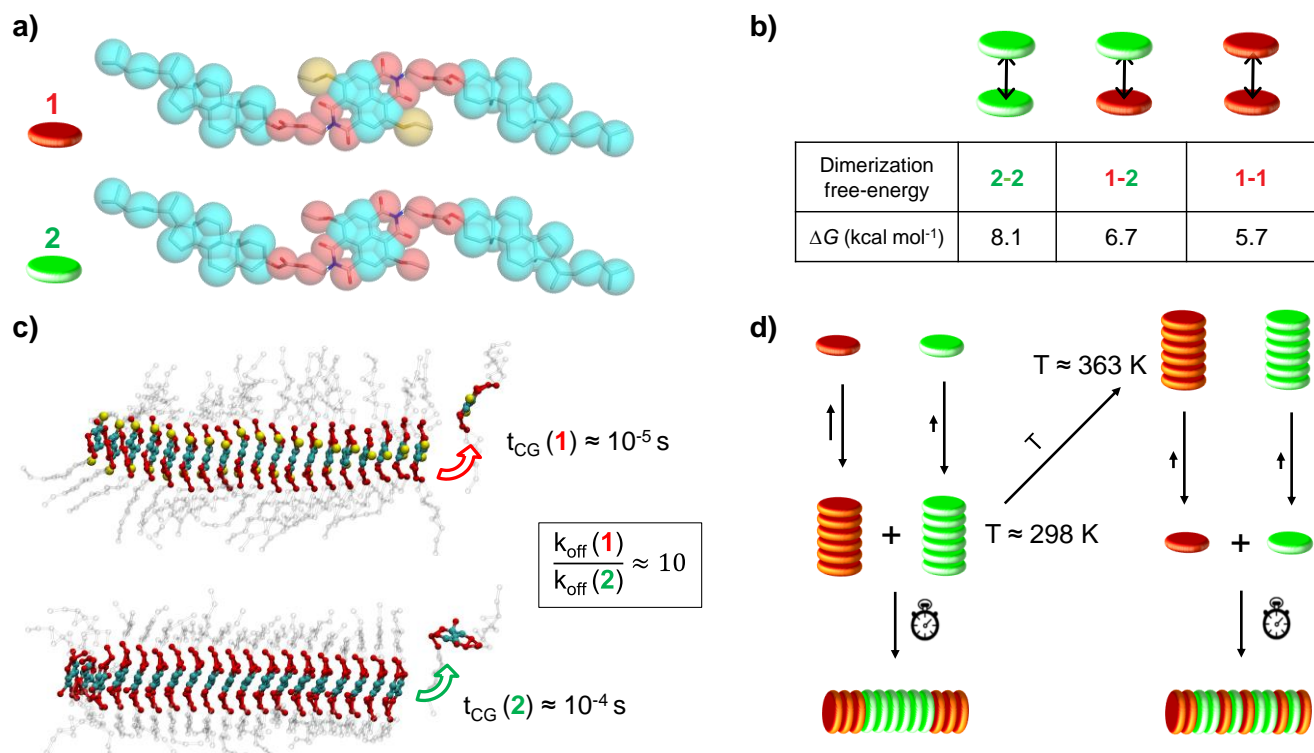


Figure 3. Coarse-grained (CG) molecular simulations of systems **1** and **2**. (a) CG models of monomers **1** and **2** (resolution of the CG models $\sim 5\text{\AA}$). (b) The dimerization free-energies, ΔG , obtained from WT-MetaD simulations allow to compare the strength of monomer-monomer interactions (**1-1** vs. **1-2** vs. **2-2**). (c) Infrequent WT-MetaD simulations exploring the mechanism and kinetics of monomer exchange from the tips of fibers **1** and **2**. Monomer exchange is found to be ~ 10 times faster in fiber **1** than in fiber **2**. (d) Model for the formation of blocks of **1** and **2** at room temperature (kinetically trapped state), while increasing the temperature to $\sim 90^\circ\text{C}$ moves the equilibrium toward the monomers, which then tend to self-assemble in a random fashion.

Fibers stability and monomer exchange kinetics of **1** vs. **2**:

Molecular simulations are extremely useful to obtain a molecular level understanding of dynamic supramolecular polymers.^{24,25,26} In particular, by combining fine coarse-grained (CG) models and enhanced sampling approaches it is possible to explore the monomer-monomer interactions, the monomer exchange kinetics and the relative stability of assemblies.^{17,27} Hence we have built fine CG models for monomers **1** and **2** (having resolution $\sim 5\text{\AA}$), which are shown in Figure 3a. Using the same approach used recently for modeling similar supramolecular assemblies,^{17,24-27} the CG models, which are based on the MARTINI CG force field,²⁴ have been then refined in order to be consistent with the behavior of the systems seen at the all-atom level (see Computational Methods in the Supporting Information for details). We used well-tempered metadynamics (WT-MetaD) simulations²⁹ to explore the stacking/de-stacking of monomers in explicit cyclohexane solvent. This allowed us to compare the free-energy of binding between different monomers, *i.e.* **1-1** vs. **1-2** vs. **2-2** (see table in Figure 3b). The **2-2** interaction (8.1 kcal mol⁻¹) was found to be

stronger than the mixed **1-2** interaction (6.7 kcal mol⁻¹) and of the **1-1** interaction (5.7 kcal mol⁻¹). Accordingly, fiber **1** was found to be more dynamic than fiber **2**. Starting from pre-equilibrated stacks containing 20 monomers of **1** or **2**, we used infrequent WT-MetaD simulations to activate and study the event of monomer exchange from the tips of the two fibers.^{17,27} At this scale, the exchange of monomers out from the assemblies is a rare event, the kinetics of which can be reliably reconstructed from multiple biased WT-MetaD exchange runs (see also Methods in the SI).^{17,27,30,31} These simulations provided characteristic monomer exchange timescales (t_{CG}) in the order of $\sim 10^{-4}$ s for fiber **2** vs. $\sim 10^{-5}$ s for fiber **1** (Figure 3c). While these exchange timescales are obtained from approximated CG models, and should be thus considered qualitatively, these still maintain a useful comparative value, indicating that the monomer exchange rate (k_{off} , which can be calculated as $1/t_{CG}$) is ~ 10 times faster in fiber **1** than in fiber **2**. Altogether, these data allowed us to propose the scheme for the formation of block vs. alternated/random supramolecular polymers of **1** and **2** as shown in Figure 3d. At room temperature (298 K) monomer exchange in/out these fibers is rare and fibers **1** and fibers **2** tend

a) Alternate copolymers

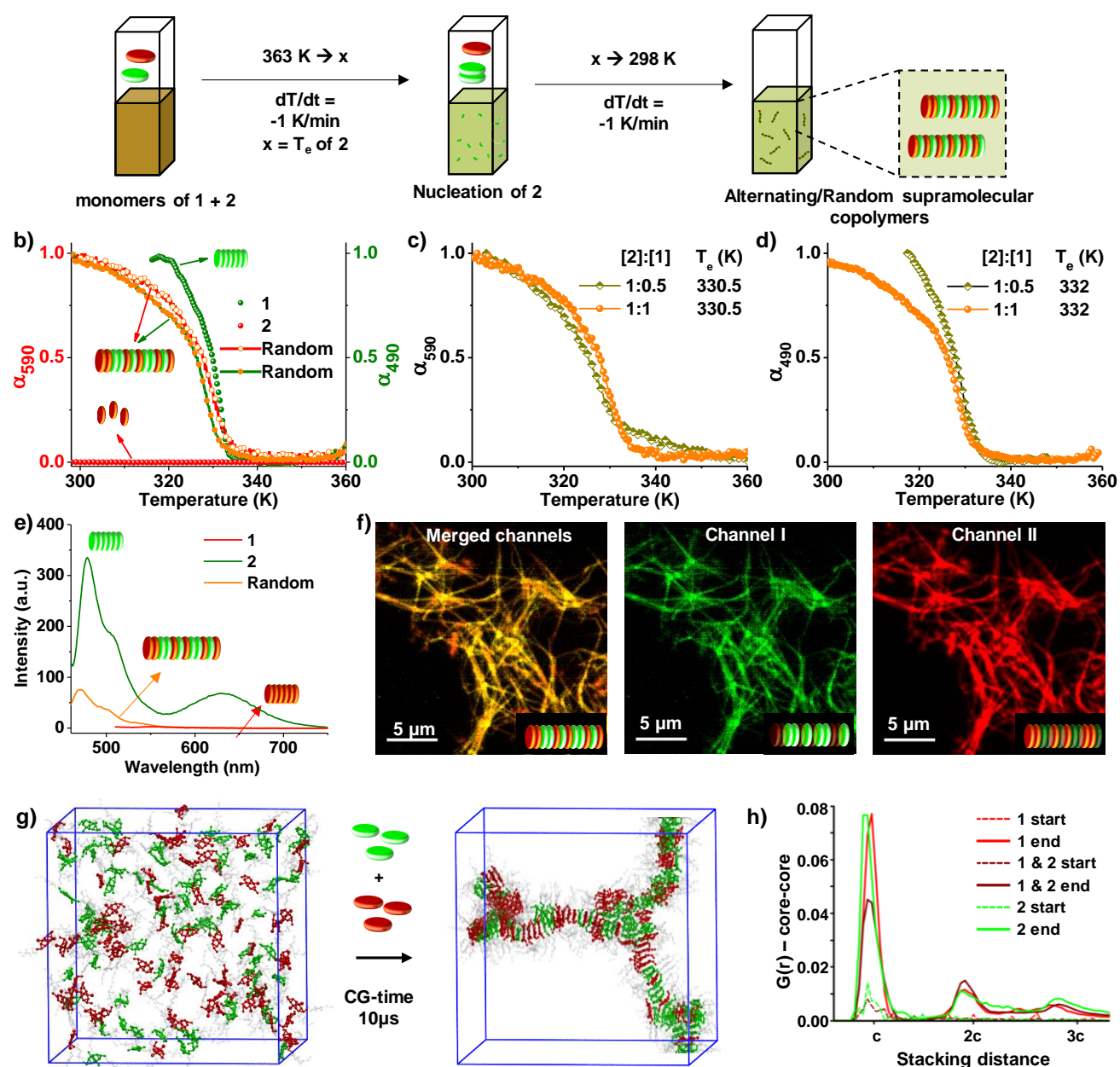


Figure 4. Experimental and simulation studies of supramolecular alternating/random co-polymers. (a) Schematic representation showing the preparation of alternating/random supramolecular co-polymer under thermodynamic control. (b) Cooling curves of homopolymers of **1** and **2** and the copolymer, showing heterogeneous nucleation under thermodynamic control, with nucleation and growth of **1** on *in situ* formed nuclei of **2**, when monitored at aggregated band of **1**. ($[1] = [2] = 2.5 \times 10^{-5} \text{ M}$). Temperature dependent cooling curves of (c), (d) co-assembled solution of **1** and **2** keeping the concentration of **2** constant and with varying concentrations of **1**, which shows a constant T_c when monitored (c) at 590 nm (corresponds to aggregate band of **1**) and (d) at 490 nm (corresponds to aggregate band of **2**) showing formation of nuclei of **2**, followed by the growth of both **1** and **2** to form an alternating/random co-polymer. ($[2] = 2.5 \times 10^{-5} \text{ M}$). (e) Quenched donor emission with the absence of excimer band depicting the formation of supramolecular alternating/random co-polymers. ($\lambda_{ex} = 442 \text{ nm}$, $l = 10 \text{ mm}$) (f) SIM image displaying completely overlapping red and green emission from **1** and **2** respectively confirming the formation of supramolecular alternating/random co-polymers. ($[1] = [2] = 2.5 \times 10^{-5} \text{ M}$, 5 % toluene/MCH). (g) Self-assembly of **1** and **2** monomers (75 + 75) during 10 μs of CG-MD. During the run, the **1** and **2** monomers mix randomly together, in the form of short red and green segments and monomers suggesting the formation of random supramolecular copolymers. (h) Radial distribution functions ($g(r)$) of the cores of the monomers at the start and end the CG-MD. (Channel I: $\lambda_{ex} = 488 \text{ nm}$, $\lambda_{coll} = 495\text{-}575$; Channel II: $\lambda_{ex} = 561 \text{ nm}$, $\lambda_{coll} = 570\text{-}650$).

to bind between them forming non-dynamic and stable blocks (the interaction **1-2** is quite strong). However, when temperature is increased to 363 K, the equilibrium moves toward the monomers and hence fibers **1** and **2** disassemble with time. From such a disassembled state, the mixing of monomers then become quite random, consistent with the rather similar **1-1** vs. **1-2** vs. **2-2** monomer-monomer interactions (Figure 3b, table). This has been supported by means of experiments and self-assembly simulations (*vide infra*).

Controlled supramolecular polymerization:

As discussed above, during a two-component supramolecular polymerization, there are three possible outcomes: formation of supramolecular homopolymers (self-sorting), supramolecular alternating/random co-polymer and supramolecular block co-polymer (Scheme 1). Based on the essential requirements as mentioned above, we thought of using molecules **1** and **2** exhibiting kinetically controlled growth and low structural mismatch for a sequence controlled supramolecular polymerization. By exploiting, thermodynamic and kinetic routes of sample preparation and the molecular level insight obtained from MD simulations of monomer exchange dynamics and interaction energies, we envisage to obtain the three structurally distinct supramolecular polymers by utilizing the same molecules. We use extensive spectroscopic and microscopic probing and MD simulations to characterize these sequence-controlled supramolecular assemblies.

Alternate/Random Supramolecular Co-polymers under Thermodynamic Control

First, we investigated the temperature-dependent supramolecular co-polymerization of an equimolar mixture of **1** and **2** monomers, by slowly cooling (1 K/min) from 363 K to 298 K (Figure 4). The growth process of both **1** and **2** can be selectively probed by monitoring their aggregation band at 590 nm and 490 nm, respectively. Homopolymerization of individual monomers (2.5×10^{-5} M) in 5 % toluene/MCH solvent mixture, by cooling at a rate of 1 K/min, showed distinct self-assembling behavior. Monomer **1** gets trapped in the metastable **State A**, whereas **2** undergoes a cooperative supramolecular polymerization with an elongation temperature (T_e) of 332 ± 1 K (Figure S15a). Thus, we envisage that, on supramolecular co-polymerization, strong homo-recognition between monomers **1** and **2** would lead to self-sorted supramolecular homopolymers, while strong hetero-recognition between the monomers would be required to construct supramolecular alternating co-polymers. Temperature-dependent supramolecular co-polymerization of 1:1 mixture of **1** and **2** from 363 K to 298 K at 1 K/min showed cooperative growth of both **1** and **2** (when monitored exclusively at their individual aggregation bands) with the identical T_e of 332 ± 1 K matching well with the T_e of pure **2** (Figure 4b) suggesting the co-assembly of monomers. At a constant concentration of **2**, the variation of concentration of **1** did not result in any change in their elongation temperature (Figure 4c,d, S16). Further, the variation of monomer composition (**1:2** ratio) while keeping the total concentration at 2.5×10^{-5} M showed a

change in T_e proportionally with the concentration of **2** (Figure S17) and matched well with the T_e of pure **2** of similar concentrations. These observations hint towards the heterogeneous nucleation of monomers of **1** triggered by **2**, rather than the cooperative growth of a hetero-dimer, which would have resulted in maximum T_e at 1:1 monomer composition. However, the resultant solution showed a new CD spectrum different from individual components with no bisignation at the absorption maximum of **2**, elucidating an alternating/random arrangement of monomers rather than blocky supramolecular polymerization (Figure S18b). The absence of excimer emission of **1** and significant fluorescence quenching of **2** due to energy transfer depicts a negligible domain formation of **2** (Figure 4e, S18). Further, the heating curve of the alternating/random supramolecular co-polymer showed significantly high stability with a melting temperature (T_m) of 340.2 K, which is 30 K higher than the homo supramolecular polymer of **1** (Figure S19). Finally, the microscopic visualization of these supramolecular polymers showed no distinct red and green emissive fibers and a complete spatial overlap validates the formation of the alternating or random supramolecular copolymers (Figure 4f, S20). We envisage that reactivity ratio ($(\Delta G_{1-1} + \Delta G_{2-2}) / 2 \times \Delta G_{1-2}$) of monomers **1** and **2** is close to unity, where disassembled **1** and **2** tend to mix at the level of individual monomers or as small groups of **1** and **2**, leading to an alternate/random arrangement of monomers under thermodynamic conditions.¹⁵ This is consistent with the rather similar **1-1** vs. **1-2** vs. **2-2** monomer-monomer interaction energies obtained from the simulations (Figure 3b) and with the mixing fashion that in such conditions is entropically more favorable (random mixing). This has been further confirmed by means of a coarse-grained molecular dynamics simulation (CG-MD) where 75 + 75 initially disassembled **1** and **2** monomers self-assemble spontaneously during 10 μ s of CG-MD in explicit cyclohexane solvent molecules. During this simulations, **1** and **2** monomers mix in a quite random fashion, where red and green monomers alternate as short segments and/or individual monomers into the generated copolymers as evident from Figure 4g (see also supplementary movies). Notably, the size of the **1** and **2** domains in the mixed stack is related, together with a favored random mixing, in consistent with the fact that **1-1** vs. **1-2** vs. **2-2** interactions, though very similar, are not perfectly identical. This is also supported by the radial distribution functions $g(r)$ of the cores at the start and end of the CG-MD simulations (dotted vs. solid lines in Figure 4h). In general, all the $g(r)$ peaks increases during the CG-MD, which is intrinsically related to the formation of ordered stacks (high-probability to find cores at stacking distance c in the self-assembled structures). However, it is interesting to note that the first peak, $g(c)$, is higher in the red and green curves (accounting for **1** and **2** monomers only), compared to the dark red curve which accounts for all monomers. This demonstrates that small groups (i.e. dimers, trimers, etc.) of red or green are more present in the self-assembled structure than a perfect mixing of alternated **1** and **2** monomers (a perfect monomeric

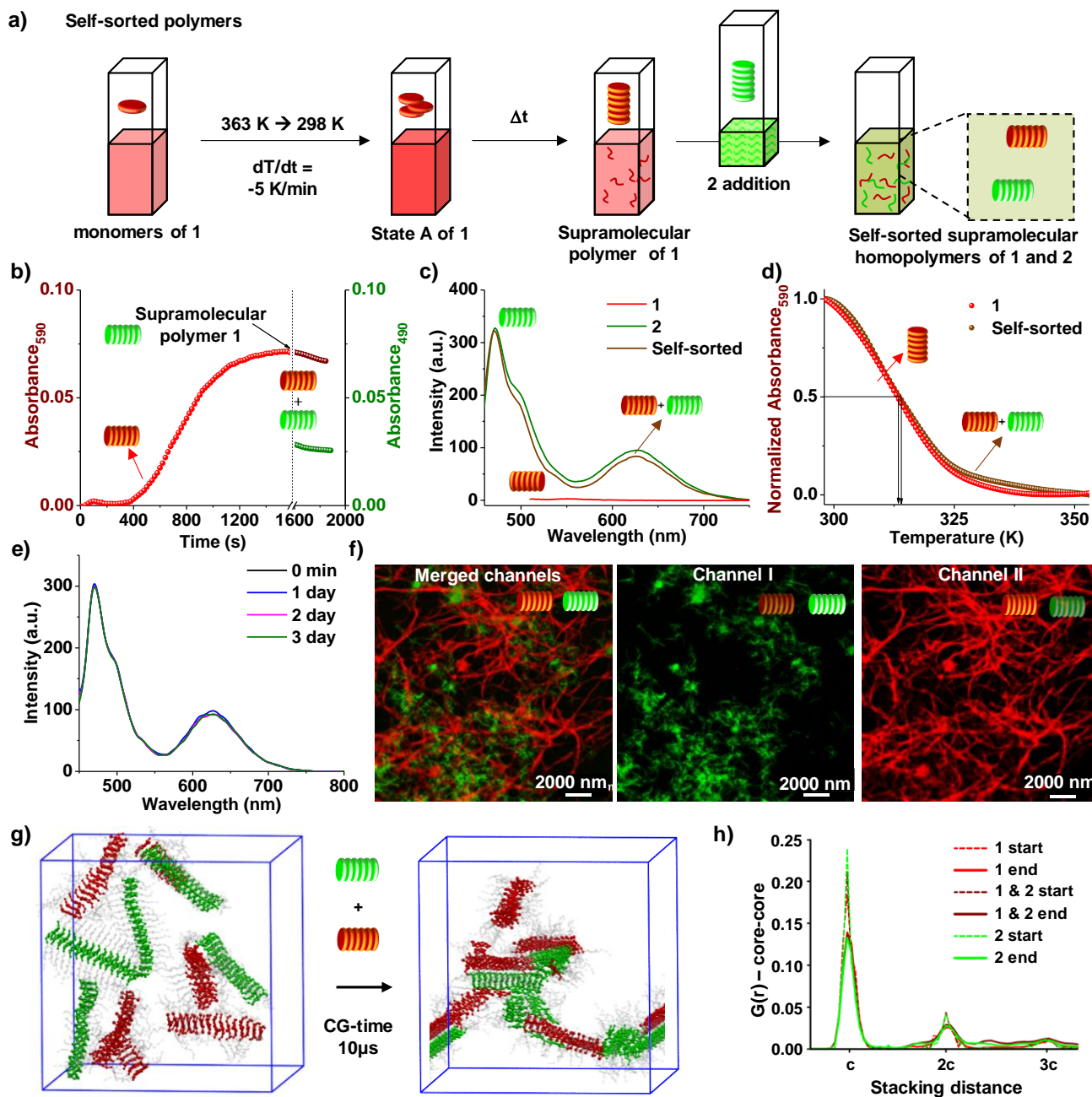


Figure 5. Experimental and computational characterization of self-sorted supramolecular homopolymers of **1** and **2**. (a) Schematic representation showing the preparation of the self-sorted homopolymers. (b) Time-dependent absorbance changes monitored at the aggregate bands of **1** (590 nm) and **2** (490 nm). Addition of supramolecular polymer of **2** to a kinetically grown supramolecular polymer of **1** (after 1600 s), shows insignificant changes in the supramolecular organization of **1** and **2**. (c) Emission spectra of self-sorted and homopolymers of **1** and **2** ($\lambda_{\text{ex}} = 442$ nm, $l = 10$ mm), showing negligible energy transfer in the self-sorted state as evident from the lack of quenching of both aggregate and excimer emission of the donor **2** ($\lambda_{\text{ex}} = 442$ nm, $l = 10$ mm). (d) The heating curve (monitored with the absorbance at 590 nm) of the solution of self-sorted supramolecular polymers of **1** and **2** showing similar stability of **1** compared to the pure supramolecular polymer of **1** (heating rate = 1 K/min) (e) Time-dependent emission spectra ($\lambda_{\text{ex}} = 442$ nm, $l = 10$ mm) of the self-sorted fibers, which does not show any changes with time suggesting the low dynamics of the monomers. (f) Corresponding SIM images showing spatially segregated supramolecular polymers of **1** and **2** confirming the self-sorted arrangement. ($[1] = [2] = 2.5 \times 10^{-5}$ M, 5 % toluene/MCH, $l = 10$ mm). (g) Self-assembly of 5 + 5 preformed stacks each containing 15 monomers of **1** and **2** during 10 μs of CG-MD. During the run, the **1** and **2** stacks interact and self-assemble, yet remaining quite stable. (h) Radial distribution functions ($g(r)$) of the cores of the monomers at the start and end the CG-MD. (Channel I: $\lambda_{\text{ex}} = 488$ nm, $\lambda_{\text{coll}} = 495\text{--}575$; Channel II: $\lambda_{\text{ex}} = 561$ nm, $\lambda_{\text{coll}} = 570\text{--}650$).

mixing would provide a higher $g(c)$ peak for the dark-red curve).

These observations show a marked difference from a recent report by Meijer and coworkers, where prevailing homo over heterorecognition leads to the formation of supramolecular block co-polymer under thermodynamic control.^{11,15} Altogether our results strongly demonstrate that such different behavior between supramolecular systems based on similar concepts is most likely the consequence of homo vs. hetero monomer-monomer interactions in the system. In fact, a perfect segregation in stable blocks starting from disassembled monomers would imply stronger homo vs. hetero monomer-monomer interactions in the system. However, since our current system shows a rather similar hetero vs. homorecognition under thermodynamic conditions, heterogeneous nucleation under kinetic control and at low monomer dynamics (at low temperature) appears to be the only way to obtain supramolecular block co-polymers (vide supra).²⁰

Kinetically Stable, Self-Sorted Supramolecular Homopolymers

Supramolecular random co-polymerization under thermodynamic conditions prompted us to investigate the monomer exchange dynamics at room temperature and kinetic stability of the stacks by probing the spectroscopic properties of a mixture of homo-polymers of **1** and **2**. To investigate this, an equimolar mixture of **1** and **2** stacks was synthesized by the post-synthetic mixing of the supramolecular homopolymers of **1** and **2**, made individually by the kinetically controlled nucleation-growth (Figure 5a). Independent spectroscopic probing of the **1** and **2** monomers in the homopolymers, did not show any absorbance changes at 590 nm for **1** and 490 nm for **2**, hinting towards the absence of any structural reorganization upon mixing (Figure 5b). Further, the final absorption and CD of this mixture matches well with the summation of the two individual supramolecular polymers (Figure S21), hinting to a self-sorted fibers. In addition, absence of energy transfer depicted by insignificant quenching of donor (**2**) emission in the mixture characterizes that the homopolymers of **1** and **2** are spatially self-sorted (Figure 5c, S22). The heating curves of self-sorted assemblies and individual homostacks match well and show an identical melting temperature (T_m) of 313 ± 1 K reiterating that the supramolecular homopolymers are spatially segregated (Figure 5d, S23). Low exchange dynamics of the monomers at 298 K were further probed via energy transfer for three days, which did not show any change in the emission profile of the donor-acceptor mixture (Figure 5e, S24). These kinetically stable, self-sorted supramolecular homopolymers were microscopically visualized by selective excitation of **1** and **2** (561 nm and 488 nm respectively) using SIM where spatially segregated green and red emissive supramolecular polymers were observed (Figure 5f, S25). This is again consistent with the low monomer exchange dynamics seen for these assemblies from the simulations (Figure 3). This was also further supported by a CG-MD simulation where we inserted 5 + 5 preformed stacks of **1** and **2** monomers, respectively. During the CG-MD run, the stacks were seen to

interact in a stable way, but with very limited internal mixing of monomers (Figure 5g, see also supplementary movies). The persistence of the **1** and **2** stacks is also evident in the $g(r)$ of the cores in the system, which demonstrate that in this case the first $g(c)$ peak, very high from the beginning, changes minimally during the run following to a very limited rearrangements in the stacks (Figure 5h).

Block Co-polymers under Kinetic Control

Random co-polymerisation of the monomers during the thermodynamic supramolecular co-polymerization (vide supra), suggests that construction of two-component supramolecular block co-polymers requires a kinetically controlled living supramolecular polymerization process. Low monomer exchange dynamics at room temperature, kinetically realizable metastable monomer states, nucleation-growth mechanism of self-assembly and tendency of heterorecognition due to low monomer structural mismatch between **1** and **2**, as evident from the detailed probing of the kinetic and thermodynamic aspects (vide supra) of the monomers, are perfectly suited for the construction of block supramolecular structures via heterogeneous nucleation of the second monomer on the seeds or stacks of the first molecule. Molecule **1** exhibits seeding characteristics as displayed by immediate non-sigmoidal growth of metastable **State A** of **1** upon the addition of pre-grown seeds of **1** during the lag phase (Figure S26).¹⁴ Due to kinetically controlled nucleation-growth of **1** with a longer lag phase compared to **2**, along with the higher kinetic and thermodynamic stability of **2** (Figure 3), latter was introduced as a seed to the metastable monomers of **1** to trigger the heterogeneous nucleation (Figure S15). A self-assembled segment of monomers **2** (seeds), was synthesized by sonicating a solution of kinetically grown **2** (5×10^{-4} M) for 5 minutes (5 % toluene/MCH) as described in the experimental section. Further metastable state of **1** (**State A**) was synthesized by the fast cooling (5 K/min) of its monomeric solution at 363 K in 5 % toluene/MCH. The addition of seeds of **2** to **State A** of **1** resulted in an instantaneous non-sigmoidal growth depicting heterogeneous nucleation and supramolecular co-polymerization, which is evident from the changes in absorbance monitored selectively at 590 nm corresponding to the aggregate band of **1** (Figure 6b,c). Variation of monomer to seed concentration showed consistent heterogeneous nucleation on the introduction of seed (Figure S27). Increase of monomer concentration with a constant seed concentration showed a linear increase in polymerization rate without any lag phase (Figure 6c, S27a,b). In addition, the variation of seed concentration with constant monomer concentration showed a linear increase in polymerization rate with an increase of seed concentration (Figure S27e,f). These kinetic features refer to the seeded supramolecular polymerization between **1** and **2**.²⁰ Absorption and CD spectra of the resultant co-assembled solution show that individual aggregation characteristics of both molecules are retained (Figure S27c, d). Further, persistent excimer emission confirms the presence of homopolymeric domains of **2** in the resultant structure (Figure S28). In addition, the self-assembled emission

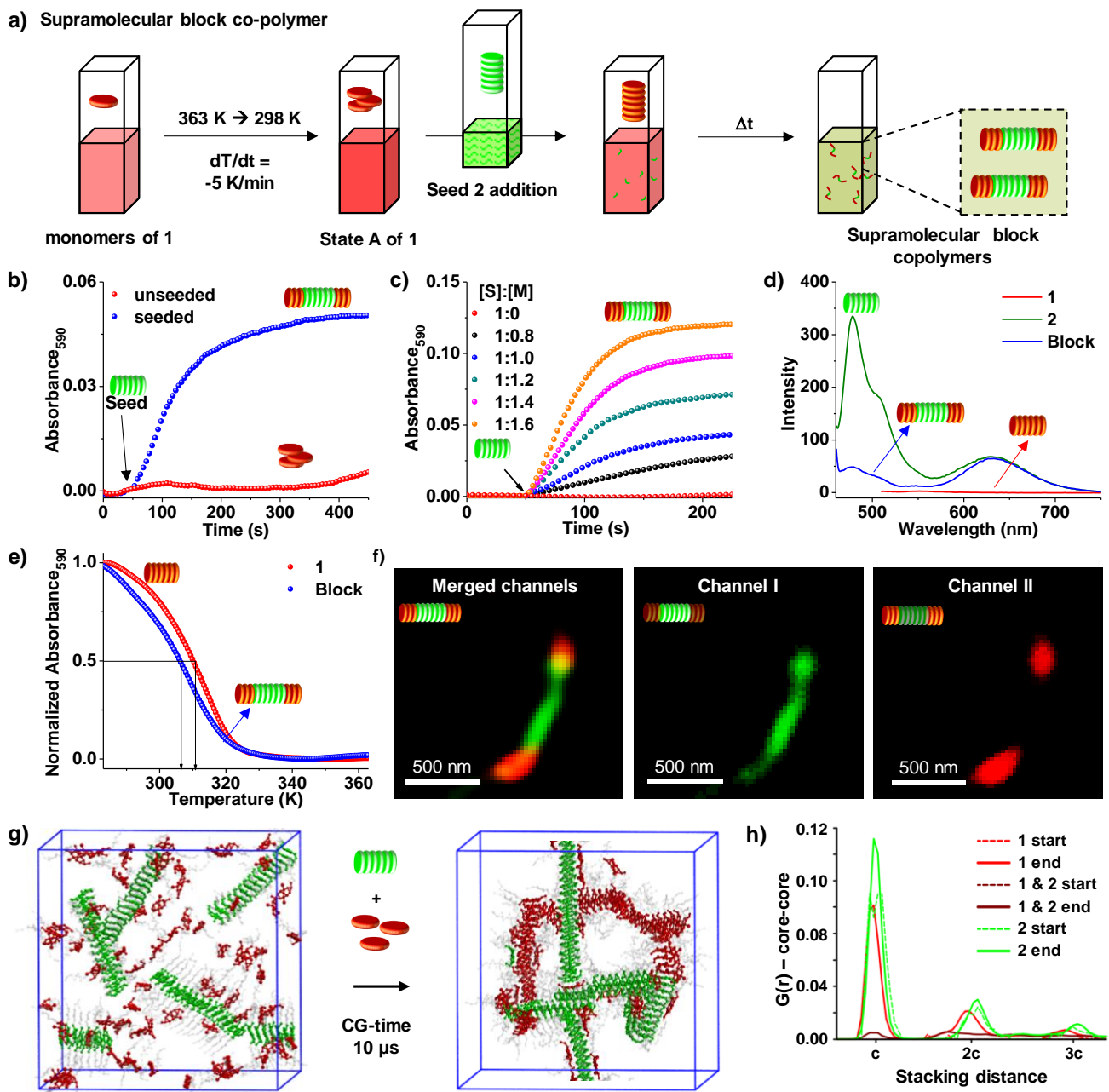


Figure 6. Experimental and computational characterization of heterogeneously seeded supramolecular block co-polymers. (a) Schematic illustration presenting the synthesis of the supramolecular block co-polymers. (b) Time progressive change in absorbance after addition of seeds of **2** (blue curve) at the lag phase of **1** (red curve) monitored at 590 nm (aggregation band of **1**) showing heterogeneous seeding of **1** on **2** supramolecular polymers ($[1] = [2] = 2.5 \times 10^{-5}$ M in the final solution). (c) Time-dependent absorbance changes at 590 nm corresponding to the growth of **1** upon heterogeneous nucleation with constant seed and varying monomer ratio ($[2] = 2.5 \times 10^{-5}$ M in the final solution). (d) Emission spectra ($\lambda_{ex} = 442$ nm) displaying a significant donor emission quenching due to energy transfer from **2** to **1** illustrating an efficient interaction between supramolecular polymers of **1** and **2** and thus suggesting a block formation. (e) Heating curves (1 K/min heating rate) of supramolecular block co-polymer and homopolymer **1** (monitored at aggregate band of **1** at 590 nm), showing a decreased stability of segments of **1** in the co-assembled structure. ($[1] = [2] = 2.5 \times 10^{-5}$ M). (f) SIM images depicting supramolecular block co-polymers ($[1] = [2] = 2.5 \times 10^{-5}$ M, 5 % toluene/MCH). (g) Self-assembly of 75 initially disassembled monomers in the presence of 5 preformed stacks each containing 15 monomers of **2** (seeds) observed during 10 μ s of CG-MD. During the run, the **1** monomers self-assemble between them using the **2** stacks as nucleation seeds. (h) Radial distribution functions ($g(r)$) of the cores of the monomers at the start and end the CG-MD. (Channel I: $\lambda_{ex} = 488$ nm, $\lambda_{coll} = 495$ –575; Channel II: $\lambda_{ex} = 561$ nm, $\lambda_{coll} = 570$ –650).

of **2** at 480 nm is significantly quenched upon co-polymerization, suggesting an efficient energy transfer to **1**, which also hints towards the close proximity of the self-assembled domains of **1** and **2** (Figure 6d, S28). All these observations corroborate to the formation of a supramolecular block co-polymer with an alternating sequence of self-assembled segments of **1** and **2**. The melting curves of the resultant co-assembled stacks showed increased stability of **1** and a negligible difference in the stability of **2** illustrating a block structure (Figure 6e, S29). The supramolecular block co-polymers were visualized by merging green and red emission channels in SIM images showing a spatial correlation of green-emitting fibers terminating with two red-emitting segments (Figure 6f, S30). Due to sonication of supramolecular polymers of **2** to prepare seed, the sizes of final supramolecular block co-polymers are smaller than that of alternating and self-sorted supramolecular polymers as seen before. Importantly, no significant changes were observed in the spectroscopic features of these block structures, even after 3 days illustrating its low dynamics and high kinetic stability (Figure S31). However, both self-sorted homopolymers and supramolecular block co-polymers, formed under kinetically controlled manner, when heated to 363 K and cooled at 1 K/min, reconfigured to the thermodynamically more stable alternating/ random supramolecular co-polymers as expected (Figure S32-S33).

These experimental evidences were again supported with molecular simulations. We equilibrated by means of a 10 μ s CG-MD simulation a CG system containing 75 initially dispersed monomers of **1** in the presence of 5 pre-formed stacks of **2** (seeds) similar to a heterogeneous nucleation experiment. In this run, we could observe spontaneous self-assembly of red monomers (**1**) using the green stacks (**2**) as nucleation sites (Figure 6g, see supplementary movies). Thus the simulations show that the self-assembly of **1** is seeded by the assemblies of **2**, in consistent with the experiments. Also, at the same time, this CG-MD shows no appreciable internal rearrangements or dynamic equilibrium of the formed assemblies. The $g(r)$ peaks, absent initially for the assemblies of **1** ($g(r)=0$), increase at the end of the CG-MD simulation. However, the $g(r)$ of the monomers of **2** also increases, while the mixing appears to be negligible (Figure 6h). This simulation confirms that the **1** monomers indeed self-assemble onto the **2** seeds, but no internal reshuffling of **1** and **2** monomers is evidenced in the generated assemblies in the timescale accessible by these simulations.

Conclusions

In conclusion, through appropriate usage of kinetic and thermodynamic pathway complexity²¹ of molecular self-assembly, we have accomplished an unprecedented sequence control in the supramolecular copolymerization of two core-substituted naphthalene diimide π -conjugated monomers to yield self-sorted, alternate/random and block supramolecular polymers. Further the characteristic and distinct optical properties of both cNDI monomers which are highly sensitive to the intermolecular interactions along with the induced circular dichroism from the peripheral chiral groups could be explored for the *in situ*

spectroscopic probing of the monomer sequence during the kinetically and thermodynamically driven supramolecular co-polymerization process. Further, orthogonal, green and red fluorescence of the monomers **1** and **2** in the supramolecular co-polymers provided a handle to probe the monomer exchange dynamics in the stacks and also helped to uniquely characterize these multi-component structures by visualized under super-resolved structured illumination microscopy. Detailed molecular dynamic simulations provided mechanistic insights into the strength of the inter-monomeric interactions and provided insight into the relative rate of the monomer exchange dynamics in these assemblies, which was crucial for rationalizing the results from heterogeneous nucleation experiments. Thermodynamic co-polymerization of monomers **1** and **2** resulted in the alternate or random sequence of monomers, due to the comparable **1-1**, **2-2** and **1-2** interaction energies between the monomers. On the other hand, self-sorted homopolymers of **1** and **2** could be realized by exploiting low monomer exchange dynamics or high kinetic stability of these assemblies. Finally, the most challenging block supramolecular co-polymers with domains of **1** and **2** have been synthesized by heterogeneous seeded growth.

Synthesis of block-supramolecular structures through kinetically controlled heterogeneous nucleation strategy presented here is particularly interesting as it is reminiscent of the macro-initiator approach in classical living polymerization by the sequential addition of monomers. This strategy is expected to give pure blocks with a control on the length, compared to the thermodynamically driven cooperative supramolecular co-polymerization based on the reactivity ratio of monomers. We plan to investigate the dispersity and length control of block segments in near future similar to what has been achieved in single component supramolecular polymers.^{14,32} Further, supramolecular block co-polymerization of optoelectronically active π -conjugated monomers as described here, would be the way forward to synthesize axial organic heterostructures with a pure donor-acceptor interface to extract interesting functions analogous to well-studied axial inorganic heterostructures. Moving towards higher complexity in terms of multicomponent systems and careful utilization of equilibrium and non-equilibrium states can render much wider regime of structural and functional states of supramolecular polymers.

ASSOCIATED CONTENT

Supporting Information. Extended computational methods and procedures. This material is available free of charge via the Internet at <http://pubs.acs.org>. For instructions on what should be included in the Supporting Information as well as how to prepare this material for publication, refer to the journal's Instructions for Authors.

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