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#### Supporting Online Material

www.sciencemag.org/cgi/content/full/313/5783/77/DC1  
Materials and Methods  
Fig. S1  
References and Notes

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# Probing the Solvent-Assisted Nucleation Pathway in Chemical Self-Assembly

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Hierarchical self-assembly offers a powerful strategy for producing molecular nanostructures. Although widely used, the mechanistic details of self-assembly processes are poorly understood. We spectroscopically monitored a nucleation process in the self-assembly of  $\pi$ -conjugated molecules into helical supramolecular fibrillar structures. The data support a nucleation-growth pathway that gives rise to a remarkably high degree of cooperativity. Furthermore, we characterize a helical transition in the nucleating species before growth. The self-assembly process depends strongly on solvent structure, suggesting that an organized shell of solvent molecules plays an explicit role in rigidifying the aggregates and guiding them toward further assembly into bundles and/or gels.

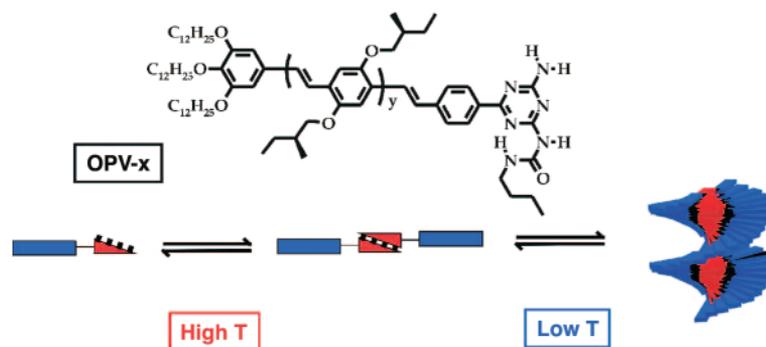
Chemical self-assembly offers a very attractive approach for constructing complex, supramolecular nanostructures. Hierarchical processes, typical of chemical self-assembly, spontaneously produce ordered ensembles of single or multiple molecular components and are ubiquitous in chemistry, physics, materials science, and biology (1). For example, a large variety of molecules has been reported that form gels through three-dimensional networks of (bundles of) fibers (2). Highly versatile biomaterials have been produced by self-assembly of peptide amphiphiles (3), and (semi)conducting tubes and rods have been achieved from properly chosen  $\pi$ -conjugated oligomeric building blocks (4). In many cases, the resulting fibrillar structures are helical, and a preferred handedness is obtained by the introduction of stereocenters into the building blocks (5). Ensuring that the components aggregate in a specific motif, however, remains a formidable task; molecular components are easily trapped in kinetically stable arrangements of varying topology

(6). The application of spectroscopy in combination with scattering and microscopy techniques has provided a reasonable view of the final self-assembled structures, yet a thorough understanding of the processes leading up to these structures remains elusive. Unravelling such structural pathways is crucial for the widely sought goal of extending rational synthesis into the nanoscale regime (7).

Although the growth of fibrillar structures typically requires nucleation, prior studies have not focused on the nature and properties of the nuclei. This lack of data is in sharp contrast to

the abundant studies on the crystallization of small molecules (8) or polymers (9), and on the aggregation of proteins (10). For instance, notable insight into the formation of actin filaments, microtubules, and viral capsids by a nucleation-growth mechanism was obtained after the pioneering work of Klug (11) and Caspar (12). More recently, this insight was used to study pathologies stemming from amyloid deposits and diseases related to protein aggregation (13). Phenomenologically, the different phases of protein aggregation should be very similar to those involved in chemical self-assembly. Two models are commonly invoked to describe protein aggregation. Isodesmic self-assembly (10), also called multistage open association or ladder- or free-association model, is noncooperative, and the association constant is independent of the size of the object. Nucleated self-assembly, also called nucleation growth or initiation elongation, is characterized by a size-dependent association constant that gives rise to cooperative kinetics. We have sought to characterize such processes in chemical synthetic systems.

The fibrillar structures we examined comprise oligo(*p*-phenylenevinylene) derivatives (OPV-*x*) with chiral side chains capped on one end by a tridodecyloxybenzene and on the other by a ureidotriazine tailored for self-complementary fourfold hydrogen bonding (structure shown in Fig. 1) (14, 15). Previous



**Fig. 1.** Molecular structure of the oligo(*p*-phenylenevinylene) derivatives OPV-*x* [for OPV-3, *y* (number of dialkoxybenzene units) = 1; OPV-4, *y* = 2; and OPV-5, *y* = 3] and schematic representation of the self-assembly process, with blue blocks representing the OPV backbone and red wedges representing the hydrogen-bonding end groups.

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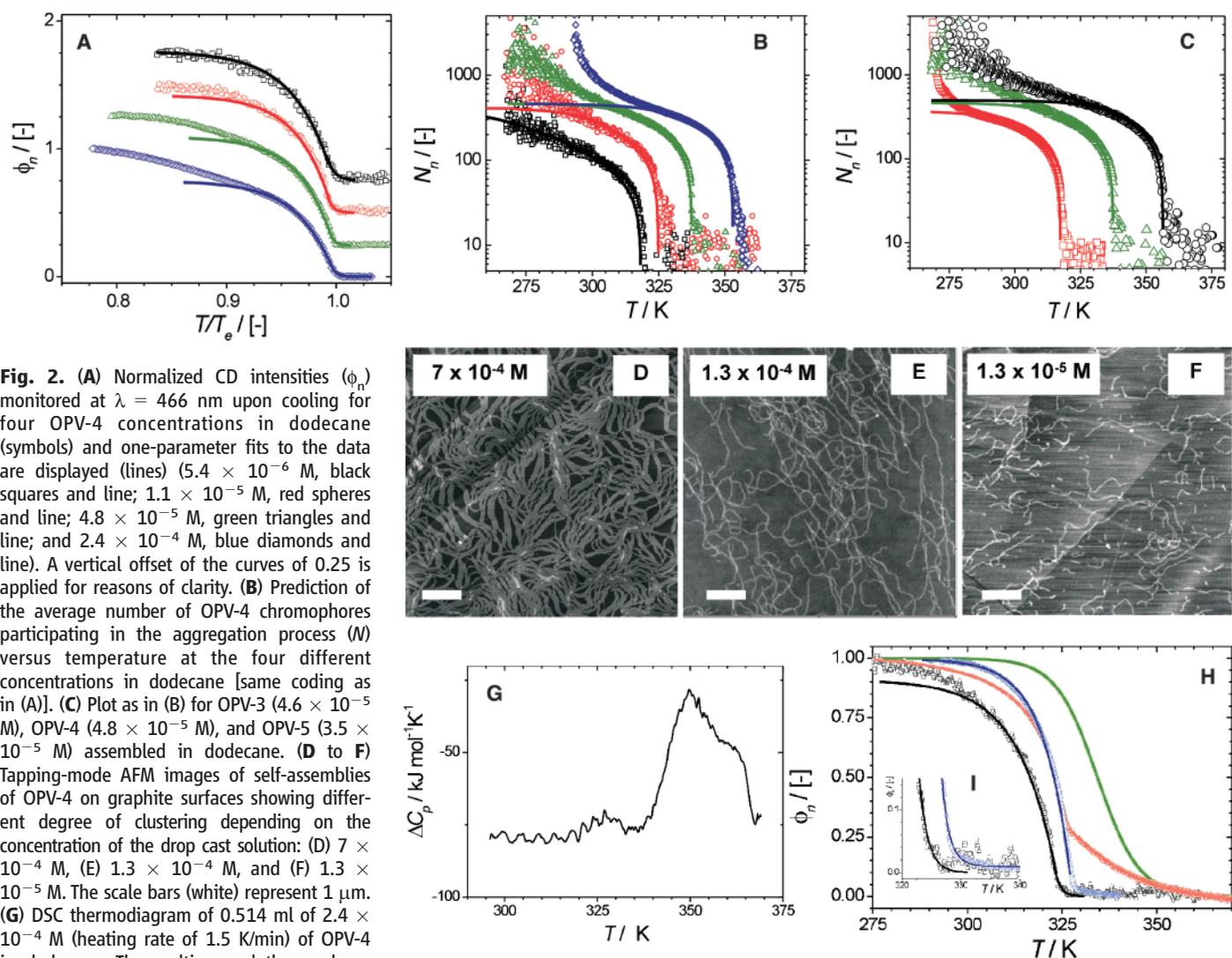
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temperature-dependent ultraviolet and visible absorption (UV/vis), fluorescence, and circular dichroism (CD) measurements (fig. S1) (14, 15) revealed two different states of the  $\pi$ -conjugated chromophores in dodecane solution: discrete monomeric or hydrogen-bonded dimeric species at high temperature and helical aggregates at low temperature (Fig. 1). The hydrogen-bonded dimers have been studied in detail with scanning tunneling microscopy (STM) (16) and  $^1\text{H}$  nuclear magnetic resonance (NMR) spectroscopy (14), whereas the fibrillar structural dimensions have been measured by small angle neutron scattering (SANS) and atomic force microscopy (AFM) (15). For example, OPV-4 exhibited a fibril persistence length of 150 nm and diameter of 5.8 nm in dodecane (table S1). The diameters of the fibrils were in close agreement with molecular modeling studies (table S1). Fluorescence microscopy revealed a per-

pendicular chromophore orientation within the fibrils (17).

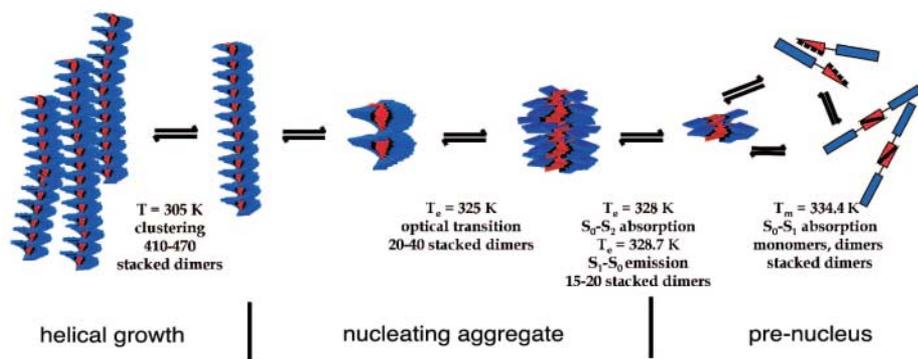
By improving our spectroscopic sensitivity, we reveal in this study distinct hierarchical stages that govern the formation process of helical OPV-4 aggregates. Optical probing is a powerful technique on account of the extreme sensitivity of the  $\pi$ -conjugated chromophore to conformational, orientational, and supramolecular states of the OPV compound. We have analyzed the data with a model similar to the Oosawa-Kasai model for helical assembly of proteins in solution (18–21), which yields thermodynamic information such as the mean size of the assemblies as well as the equilibrium constants. Moreover, we uncovered the explicit role of the solvent molecules in the formation of the supramolecular stacks. Our findings can be generally applied to many chemical self-assembly processes of ordered quasi-one-dimensional stacks.

By using CD spectroscopy at a specific wavelength, we could monitor the optical changes characterizing the crossover from the monomeric state of OPV-4 to the aggregated state in dodecane. A Peltier-temperature programmer for thermostating the samples was used to cool the solution slowly in small steps (from 375 K to 275 K, with a rate of 0.5 K/min and a resolution of one data point per 0.1 K). The slow cooling is necessary to suppress kinetic (i.e., nonequilibrium) effects on the self-assembly (fig. S2, A and B). We tracked the growth in the intensity of the  $\lambda = 466$  nm CD band for four different OPV-4 concentrations and consistently observed a sharp increase at a specific temperature (fig. S2C), indicative of the transition from monomers into helical aggregates with a preferred handedness. This curve is not sigmoidal, and hence it cannot be described by an isodesmic model. However, it does follow expected Oosawa-Kasai behavior



**Fig. 2.** (A) Normalized CD intensities ( $\phi_n$ ) monitored at  $\lambda = 466$  nm upon cooling for four OPV-4 concentrations in dodecane (symbols) and one-parameter fits to the data are displayed (lines) ( $5.4 \times 10^{-6}$  M, black squares and line;  $1.1 \times 10^{-5}$  M, red spheres and line;  $4.8 \times 10^{-5}$  M, green triangles and line; and  $2.4 \times 10^{-4}$  M, blue diamonds and line). A vertical offset of the curves of 0.25 is applied for reasons of clarity. (B) Prediction of the average number of OPV-4 chromophores participating in the aggregation process ( $N_n$ ) versus temperature at the four different concentrations in dodecane [same coding as in (A)]. (C) Plot as in (B) for OPV-3 ( $4.6 \times 10^{-5}$  M), OPV-4 ( $4.8 \times 10^{-5}$  M), and OPV-5 ( $3.5 \times 10^{-5}$  M) assembled in dodecane. (D to F) Tapping-mode AFM images of self-assemblies of OPV-4 on graphite surfaces showing different degree of clustering depending on the concentration of the drop cast solution: (D)  $7 \times 10^{-4}$  M, (E)  $1.3 \times 10^{-4}$  M, and (F)  $1.3 \times 10^{-5}$  M. The scale bars (white) represent 1  $\mu\text{m}$ . (G) DSC thermogram of 0.514 ml of  $2.4 \times 10^{-4}$  M (heating rate of 1.5 K/min) of OPV-4 in dodecane. The melting endotherm shows transition temperatures at 351.0 K and 361.7 K. (H) Transition curves based on UV/vis [ $\lambda = 490$  nm (green) and  $\lambda = 335$  nm (blue)], fluorescence ( $\lambda = 500$  nm), and CD ( $\lambda = 466$  nm) spectra for OPV-4 ( $1.1 \times 10^{-5}$  M). Fits to

the data are shown on the basis of isodesmic or cooperative self-assembly. (I) A closer view of the nucleation regime based on the UV-vis ( $\lambda = 335$  nm) and CD data.



**Fig. 3.** Schematic representation of the hierarchical self-assembly of OPV-4 in solution based on the measurements presented in Fig. 2. First, monomers form dimers via quadruple hydrogen bonding. Upon cooling, about 10 to 15 dimers are brought together via an isodesmic pathway, forming disordered stacks. Upon further cooling, the molecules in the preaggregates become more restricted in relative position via a cooperative process ( $T_e = 328$  K). In the next step, the preaggregates undergo a coil-helix transition to form a chiral nucleus of about 28 dimers, at which point the elongation-growth pathway sets in. Lastly, the cooperative stack length is reached and clustering of the assemblies occurs.

for thermally activated equilibrium polymerization (19), in which the non-isodesmic helical assembly (characterized by an activation step and subsequent propagation steps) is preceded by the isodesmic assembly of nonhelical structures. The modified model is a mathematically more tractable version of a self-assembled Ising chain theory introduced to describe the helix-coil transition in supramolecular polymers (22, 23). In the simplified model, the two kinds of assemblies are linked through equilibrium between nonhelical and helical assemblies of a critical size (19). According to the model, the number average aggregation number of the helical assemblies  $N$  obeys the following relationship:

$$K_e = \exp[h_e(T - T_e)/RT_e^2] \\ = 1 - N^{-1} + K_a N(N - 1) \quad (1)$$

with  $K_e$  as the equilibrium constant of the elongation process,  $K_a$  the equilibrium constant of

the activation step,  $T$  the absolute temperature,  $h_e$  the molecular enthalpy of the elongation process, and  $R$  the gas constant. The fraction of molecules in assemblies,  $\phi_n$ , obeys approximately

$$\{1 - \exp[h_e(T - T_e)/RT_e^2]\} \quad (2)$$

in the helical polymerized regime  $T < T_e$  and approximates

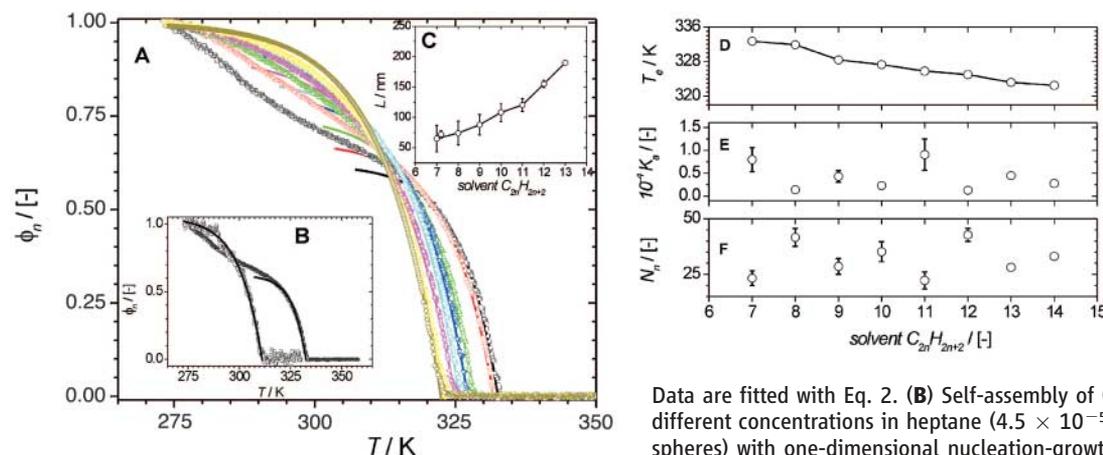
$$\{K_a^{1/2} \exp[(2/3)K_a^{-1/2} - 1] h_e(T - T_e)/RT_e^2\} \quad (3)$$

near the polymerization temperature  $T_e$  [Supporting Online Material (SOM) Text]. Applying this model to our data, we can estimate  $N$ , the mean number of dimers in the aggregate, from Eq. 1;  $T_e$ , the temperature at which elongation sets in,  $\phi_n$ , the normalized fraction of molecules in the aggregate, and  $h_e$ , the enthalpy of bond formation from both Eqs. 2 and 3; and

$K_a$ , the equilibrium constant between the active and nonactive state, from Eq. 3; active meaning the state that can initiate the elongation.

The normalized CD curves (Fig. 2A) are well fit by Eqs. 2 and 3, showing unambiguously that our model description is internally consistent and that helical aggregates are not present above the characteristic  $T_e$  (i.e., for values of  $T/T_e$  in excess of unity) but form only after the sharp nucleation step. The  $T_e$  decreases upon diluting the sample, revealing a linear relationship in the Van't Hoff plot (fig. S2D), with  $\Delta H = -100$  kJ/mol and  $\Delta S = -215$  J/mol K. Although there is a good correlation between experiment and theory in the case of the most dilute solution (5.4  $\mu$ M) (Fig. 2A), at higher concentrations the theory underestimates the CD data at low temperatures (i.e., low  $T/T_e$ ) (Fig. 2A). This result suggests an additional process that is not considered in the current polymerization theory. Interestingly, the fits for the different concentrations show that the temperature at which theory deviates from experiment ( $T_{\text{sat}}$ ) rises with increasing concentration. This deviation is also visible in a plot of  $N$  versus temperature (Fig. 2B). At lower temperatures, the average columnar length estimated from the fitted model is remarkably constant for  $N$  values from 410 to 470. Assuming a  $\pi$ - $\pi$  stacking distance of 0.35 nm (15), this  $N$  range corresponds to a length of 145 to 165 nm. These values are comparable with the values found by SANS and AFM [150 and 125 nm, respectively (table S1)]. With these data in hand, the deviation from one-dimensional growth at higher concentrations can be attributed as the clustering of stacks (both laterally and along the growing direction, Fig. 2, D to F). The data suggest this clustering is isodesmic in nature (fig. S3).

At the helical polymerization temperature  $T = T_e$ , the theory allows for the calculation of  $K_a$ : We find values of  $2.0 \times 10^{-4}$ ,  $0.46 \times 10^{-4}$ ,  $0.31 \times 10^{-4}$ , and  $0.15 \times 10^{-4}$ , in order from lowest to highest starting OPV concentra-



**Fig. 4.** Solvent-dependent CD measurements of OPV-4 in eight alkane solvents. (A) The normalized aggregate fraction of self-assembled OPV-4 in alkane solvents versus  $T/T_e$  [ $4.5 \times 10^{-5}$  M  $C_{2n}H_{2n+2}$ , where  $n$  values were 7 (black), 8 (red), 9 (green), 10 (blue), 11 (cyan), 12 (magenta), 13 (yellow), and 14 (dark yellow)] based on monitoring CD intensity at  $\lambda = 466$  nm upon cooling.

Data are fitted with Eq. 2. (B) Self-assembly of OPV-4 versus temperature at two different concentrations in heptane ( $4.5 \times 10^{-5}$  M, squares, and  $4.5 \times 10^{-6}$  M, spheres) with one-dimensional nucleation-growth fits. (C) Cooperative length ( $L$ ) for OPV-4 in the different solvents, (D)  $T_e$  (accuracy is 0.1 K based on the fit), (E)  $10^4 K_e / [H]$ , and (F)  $N_n / [H]$  are plotted versus number of carbons in alkane solvent (fig. S9). Error bars are based on the fit.

$K_a$  (error based on the determination of  $\phi_{\text{sat}}$ ), and (F)  $N$  are plotted versus number of carbons in alkane solvent (fig. S9). Error bars are based on the fit.

