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Tuning the structure of 1,3,5-benzene tricarboxamide self-assemblies through stereochemistry†

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A heterochiral 1,3,5-benzene tricarboxamide (BTA) monomer, derived from valine dodecyl ester, forms long rods in cyclohexane whilst its homochiral analogue assembles into dimers only at the same concentration. This highly original assembly behaviour is related to the destabilization of the dimeric structure containing the two heterochiral monomers as corroborated by a combined experimental and computational study.

The non-symmetrical spatial arrangement of functional organic molecules in self-assembled systems yields chiral supramolecular structures with new functions compared to the simple, non-associated, building blocks.¹ Controlling the structure and chirality of synthetic assemblies in solution is of prime importance for their applications in chiral recognition and sensing,^{1e,2} in separation processes,³ or as chiroptical switches,^{1e,4} optoelectronic materials,^{1e,5} and asymmetric catalysts.⁶

The aggregation of an enantiopure monomers into a well-defined chiral one-dimensional (1D) nanostructure in solution is usually programmed at the molecular level and executed during the self-assembly process.⁷ However, as the assembly relies on dynamic non-covalent interactions, complex self-assembly pathways also exist for which a single monomer may aggregate into morphologically-different chiral supramolecular structures (finite entities or polymeric 1D filaments, helices, nanotapes, nanotubes...). In such cases, tuning experimental conditions such as concentration,⁸ temperature,⁹ time (kinetics),¹⁰ solvent,¹¹ light^{9,12} and additives¹³ results in the selection of a predominant aggregate. Changing the stereochemistry of the interacting monomers, for example by using a scalemic mixture of monomers, mainly serves in the purpose of unveiling chirality amplification^{1b} and self-sorting¹⁴

processes. Racemic mixtures of monomers may assemble into fibrillar networks different from those formed by their homochiral stereoisomers¹⁵ but rarely leads to drastic changes in the nature of the chiral nanostructures (*e.g.* finite vs long helical aggregates).¹⁶ Efforts are required to design supramolecular assemblies which can adopt various well-defined chiral nanostructures by simply changing the stereochemistry of its interacting components.

Heterochiral monomers, *i.e.* monomers containing chiral centres of opposite chirality, inherently conduct to heterochiral assemblies since these monomers cannot be deracemized at the supramolecular level.^{15c,17} Contrary to our expectations, the heterochiral monomer of a BTA derived from valine dodecyl ester (**BTA (S,S,R)-Val**, Chart 1) assembles into long helical rod-like objects while the corresponding *homochiral monomer* (**BTA (S)-Val**) only forms dimers at the same concentration. In this work, the factors that conduct the heterochiral monomer to form longer assemblies than its homochiral analogue have been rationalized at the molecular level.

Previously, we unravelled the self-assembly behaviour of homochiral ester BTA monomers derived from α -amino esters (ester BTAs).^{8b,8c} The nature of the dominant hydrogen-bonded chiral assemblies in solution, ester-bonded dimers or helical stacks, can be controlled by the nature of the substituent (R^1 in Chart 1) and the experimental conditions (concentration, temperature). In particular, **BTA (S)-Cha** and **BTA (S)-Val** exclusively form dimers between 50 μ M and 90 mM. The molecular arrangement of these dimers, as depicted in Chart 1, has been proposed previously based on spectroscopic analyses and Molecular Mechanics/Molecular Dynamics (MM/MD) calculations.^{8c} We now report the X-ray crystal structures of **BTA (S)-Cha^{t-Pr}** (Figs. 1 and S1) and **BTA (S)-Cha^{t-Bu}** (Fig. S2), which confirm our hypotheses. These structures are similar and show that the discrete dimers adopt a D_3 symmetry with several important structural features: i) a crossed hydrogen-bond pattern involving the amide N-H and ester C=O ($H\cdots O$ distances ranging from 2.154(3) to 2.277(3)

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†Electronic Supplementary Information (ESI) available: General procedures, synthesis and characterization of ester BTAs, Fig. S1 to S15, Table S1 and crystallographic data. See DOI: 10.1039/x0xx00000x

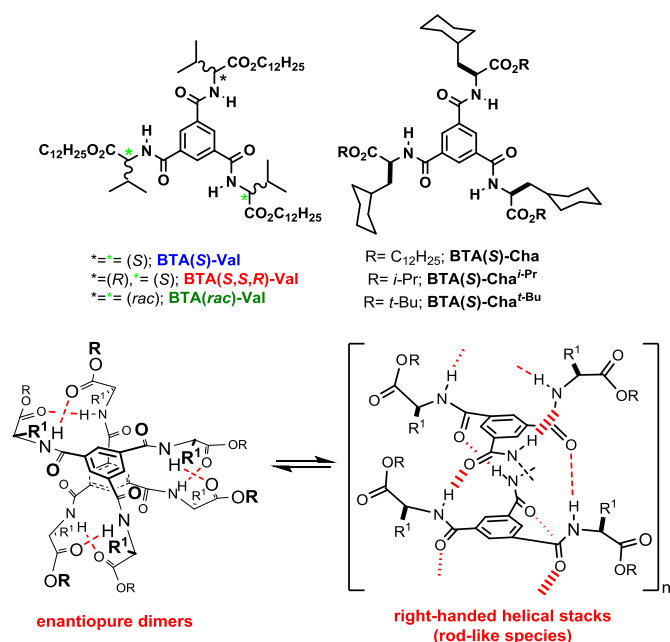


Chart 1 Chemical structures of the 1,3,5-benzene tricarboxamide (BTA) monomers investigated in this study (top). Molecular structures of the enantiopure rod-like and dimeric hydrogen-bonded species formed by BTAs derived from (*S*) α -amino esters (ester BTAs, bottom). **BTA (rac)-Val** as obtained by synthesis with racemic valine consists of a mixture of **BTA (S,S,R)-Val** (37.5%), **BTA (R,R,S)-Val** (37.5%), **BTA (S)-Val** (12.5%) and **BTA (R)-Val** (12.5%).

Å), ii) a distance between the centroids of the two stacked rings consistent with an aromatic interaction (3.468(1) and 3.502(1) Å), and iii) a (*P*) planar chiral arrangement of the aromatic rings induced by the (*S*) chirality of the peripheral amino ester groups. As a consequence of the geometrical constraints imposed by the hydrogen bonds, the alkyl ester moieties (*i*-Pr or *t*-Bu) and the cyclohexyl groups on the same BTA arm are directed in opposite directions and lie almost perpendicularly to the BTA aromatic rings. This leads to an alternate distribution of the cyclohexyl and alkyl ester groups at both upper and lower rims of the dimer (Figs. S1 and S2). Also, each aromatic C-H hydrogen of one BTA is close to the atoms of one side chain belonging to the other BTA in the dimer. This is reflected by short distances between the aromatic C-H hydrogens and the ester carbonyls belonging to the other BTA unit (ranging from 2.352(3) to 2.640(3) Å) as well as between the aromatic hydrogens and other atoms (see caption in Figs. S1 and S2).

These X-ray structures will allow for a precise understanding of the factors determining the stability of the dimeric structure formed by ester BTAs, but before, it is important to verify that the main structural features of the dimeric structure in the solid state are maintained in solution. **BTA (S)-Chaⁱ-Pr** displays virtually identical FT-IR spectra in the solid state and in solution (Fig. S3). Also, the ¹H NMR spectrum of **BTA (S)-Chaⁱ-Pr** in C₆D₁₂ (Fig. S5) shows a single set of signals demonstrating that enantiomerically-pure dimeric assemblies are present in solution. The aromatic rings adopt a (*P*) planar chirality as confirmed by the -/+/- pattern of the CD signals in the 200-260 nm region (Fig. S4).^{8c} This complete induction of the supramolecular chirality (observed both in the crystalline and solution states) is in accordance with the

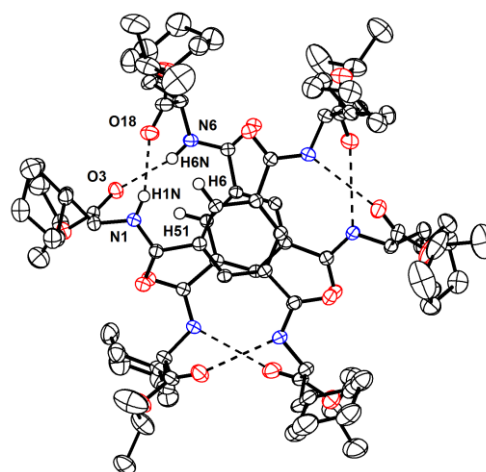


Fig. 1 X-ray crystal structure of **BTA (S)-Chaⁱ-Pr** in **BTA (S)-Chaⁱ-Pr · 0.25CH₃CN · 0.5H₂O**. H atoms (except H1N, H6N, H6 and H51) and solvent molecules omitted for clarity. Ellipsoids are represented at 30% probability level. Selected distances (Å): H1N-O18: 2.186(3); H6N-O3: 2.204(3); H6-O18: 2.482(2); H51-O3: 2.352(3).

energy difference between the (*P*) and (*M*) diastereomers calculated to be larger than 10 kcal.mol⁻¹ for a related ester BTA.^{8c} Finally, ROESY analyses were performed in C₆D₁₂ for **BTA (S)-Chaⁱ-Pr** in order to isolate atomic contacts that are specific to the dimer (Fig. S6). Remarkable dipolar couplings occur between the aromatic C-H hydrogens (all equivalent) and: i) the hydrogen atom attached to the α -carbon, ii) the methine hydrogen of the *i*-Pr group, and iii) one methyl of the *i*-Pr group. These contacts occur between the aromatic hydrogens of one BTA and the side chain of the other BTA within the dimeric structure. The NOE contacts observed in the 2D ROESY experiment are consistent with the distances between these atoms in the X-ray structure of **BTA (S)-Chaⁱ-Pr** (all inferior to 4.5 Å, Fig. S1). Taken all together, these analyses reveal that the dimeric structure is the same in the solid-state and in solution.[‡]

The tight interlocking of the two homochiral molecules in this dimeric structure offers the obvious prospect of trying to control its stability through stereochemical modifications. It seems likely that imposing a chiral mismatch in the molecular structure should destabilize the dimer.

In order to experimentally probe this stereochemical effect, **BTA (S,S,R)-Val** was synthesized. First, **BTA (rac)-Val** was prepared from trimesoyl chloride and a racemic mixture of the dodecyl ester of valine (see the S1[†]). Then, **BTA (S,S,R)-Val** was isolated from **BTA (rac)-Val** by means of preparative chiral HPLC (*ee* and *de* > 99.5%).[†] At low concentration (\leq 15 mM) in cyclohexane, **BTA (S,S,R)-Val** forms assemblies that resemble **BTA (S)-Val** dimers as shown by their similar FT-IR (10 mM, Fig. S3) and CD spectra (2 mM, Fig. S4). Notably **BTA (S,S,R)-Val** show the same CD bands than **BTA (S)-Val** but of lower intensity as expected for a heterochiral system. The discrete nature of these assemblies is established by the fact that FT-IR (Fig. S8) and ¹H NMR (Fig. S9) analyses of **BTA (S,S,R)-Val** recorded at various concentrations are unchanged up to 15 mM.[‡]

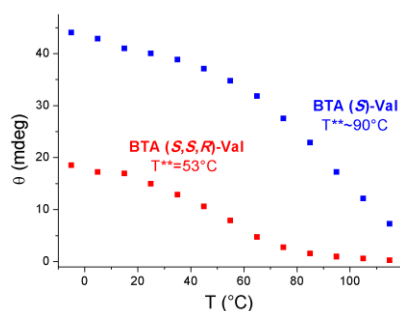


Fig. 2 Ellipticity (θ , 225 nm) versus temperature for **BTA (S,S,R)-Val** and **BTA (S)-Val** (methylcyclohexane, 25 μ M). The transition temperatures (T^{**}) between dimers and monomers are determined at the inflection point of the curves. These experiments have been performed in a closed quartz cell which explains why some of the measured temperatures exceed the boiling point of methylcyclohexane. For the full CD spectra see Fig. S13.

To gain information on the structure and stability of the potential dimers of **BTA (S,S,R)-Val**, we performed MM/MD calculations.⁵⁵ Three possible molecular arrangements were obtained, depending on the relative position of the (*R*) ester side arms within the dimers (I-III, Fig. S10), and submitted to MD simulations of 10 ns. These structures are of C_2 symmetry and have similar energies: accordingly we cannot ascertain the molecular arrangement of the heterochiral monomers in the dimers. Most importantly for the present study, they are all significantly less stable than the **BTA (S)-Val**^{Me} dimer (by 5.2 to 5.7 kcal.mol⁻¹, Table S1). An analysis of the energetic factors contributing to the destabilization of the dimers of **BTA (S,S,R)-Val**^{Me} indicates that the hydrogen-bond energy decreases, but by only 0.5 to 1.4 kcal/mol, *i.e.*, by 4 to 11% of the initial value. Most of the penalty actually arises from the valence energy term (Table S1 and Figs. S11 and S12). In other terms, the chiral mismatch in the hydrogen-bonded arms slightly lowers the strength of some of the hydrogen bonds, but more importantly it imposes on the molecular backbone geometrical constraints that increase the energy of the dimeric structure. Experimental evidence for the lower stability of the dimers of **BTA (S,S,R)-Val** compared to those of **BTA (S)-Val** was obtained by means of variable-temperature CD analyses, to probe the transition between dimers and monomers in methylcyclohexane (25 μ M). For **BTA (S,S,R)-Val**, the intensity of the CD signals rapidly decreases upon raising the temperature and only a vanishingly small CD signal is present at 105°C, which corresponds to the monomer (Fig. S13).^{8c} In contrast, full dissociation of the dimers is not observed for **BTA (S)-Val**. Plotting the ellipticity (measured at 225 nm) as a function of the temperature yields a transition temperature of 53°C for **BTA (S,S,R)-Val** and of *ca.* 90°C for **BTA (S)-Val** (Fig. 2). The fact that the dimeric structure of **BTA (S,S,R)-Val** is less stable than that of **BTA (S)-Val** has interesting consequences at higher concentrations. At 90 mM, FT-IR shows that **BTA (S,S,R)-Val** almost exclusively forms the classical stacks stabilized by hydrogen bonds to the amide C=O, whereas **BTA (S)-Val** is still assembled into dimers (Fig. 3). As a result, the viscosity of the former solution increases markedly with the concentration (Fig. 4). For instance, a 30 mM (4.0 wt%)

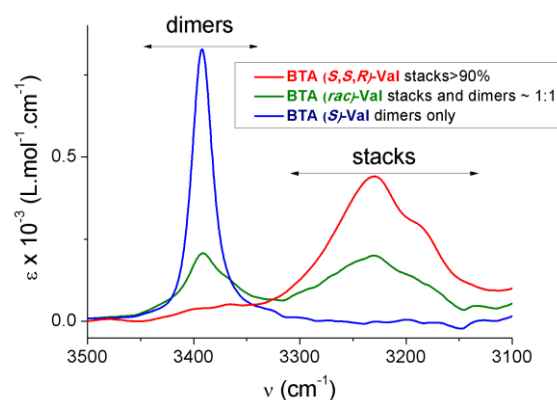


Fig. 3 FT-IR spectra of **BTA (S,S,R)-Val**, **BTA (rac)-Val** and **BTA (S)-Val** in cyclohexane (90 mM, 20°C). Zoom on the N-H region. For more details see Fig. S14.

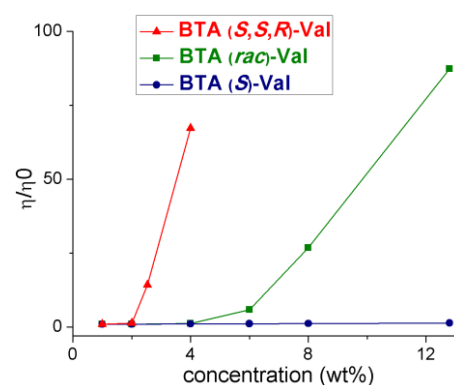


Fig. 4 Relative viscosity of **BTA (S,S,R)-Val**, **BTA (rac)-Val** and **BTA (S)-Val** in cyclohexane versus concentration (20°C).

cyclohexane solution of **BTA (S,S,R)-Val** is 60 times more viscous than that of **BTA (S)-Val**. This indicates that the drastically different self-assembly behaviour exhibited by the heterochiral monomer (**BTA (S,S,R)-Val**) compared to the homochiral one (**BTA (S)-Val**) has strong consequences on their macroscopic properties.

We finally turn our attention to the assembly properties of **BTA (rac)-Val**, which contains a large proportion of heterochiral monomers and is much easier to prepare than the pure heterochiral monomer. Following a statistical distribution of the enantiomers, **BTA (rac)-Val** is “racemic by synthesis” and is thus composed of the heterochiral monomers (**BTA (S,S,R)-Val** and **BTA (R,R,S)-Val**) and the homochiral monomers (**BTA (S)-Val** and **BTA (R)-Val**) in a 3:1 ratio. As expected, the viscosity of a **BTA (rac)-Val** solution is intermediate between those of **BTA (S,S,R)-Val** and **BTA (S)-Val** (Fig. 4). For example, a 90 mM (12.9 wt%) cyclohexane solution of **BTA (rac)-Val** is 87 times more viscous than a solution of **BTA (S)-Val**, while the **BTA (S,S,R)-Val** solution is too viscous to be measured with our capillary viscometer. **BTA (rac)-Val** forms an approximately 1:1 mixture of helical stacks and dimers at this concentration as deduced from FT-IR analyses (Fig. 3). These helical stacks are long, as indicated by the slope of the SANS intensity (Fig. S15) which is maintained down to the lowest q values measured, meaning that a significant fraction of the stacks are longer than 200 Å.

In conclusion, the structure of the chiral BTA assemblies can be tuned by modifying the stereochemistry of the BTA monomer. The heterochiral and the homochiral monomers both form a dimeric structure at low concentration, which transforms into long helical stacks only in the former molecule as a result of the lower stability of the chirally mismatched dimers. The strategy can potentially be applied to other types of monomers that self-assemble into two (or more) competing chiral supramolecular structures in solution.

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Notes and references

‡ BTA (S)-Val forms the same dimers as BTA (S)-Cha^{i-Pr}, as evidenced by similar spectroscopic analyses (including similar NOE contacts, Fig. S7).

§ FT-IR analyses of dimers of BTA (S,S,R)-Val (Fig. S8) show the presence of several superimposed N-H absorption bands in the region between 3350 and 3450 cm⁻¹ which we attribute the hydrogen bonds between the amide N-H and the ester C=O of different strength and to free N-H bonds. ¹H NMR analyses of dimers of BTA (S,S,R)-Val (Fig. S9) are consistent with a C₂-symmetric species since two different sets of signals in a 2/1 relative intensity are present. These analyses do not allow us to ascertain the exact molecular arrangement of the monomers in the BTA (S,S,R)-Val dimers.

§§ Calculations were made on BTA (S,S,R)-Val^{Me} and BTA (S)-Val^{Me} with methyl ester groups instead of dodecyl ester groups in order to alleviate the computational cost.

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