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ARTICLE TYPE

Estimating the shape and size of supramolecular assemblies by variable temperature diffusion ordered spectroscopy

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Reported is characterization of the self-assembly of π conjugated oligomers, molecules studied recently in photovoltaic devices, using variable temperature diffusion ordered spectroscopy (VT-DOSY). Iterative fitting of 10 diffusion coefficient versus temperature data to a modified Stokes-Einstein equation, molecular modelling, and comparison to non-assembling model compounds, has allowed estimation of assembly size, shape, and molecularity.

Diffusion ordered spectroscopy (DOSY) is a diffusion nuclear ¹⁵ magnetic resonance (NMR) technique that is useful for aggregate molecularity, size, and shape determination.¹ In a number of cases, DOSY NMR has been used to rigorously characterize synthetic supramolecular assemblies.^{1d, 1f, 2} Although not often coupled with variable temperature (VT) NMR, the combined

- ²⁰ techniques report on dynamic assembly processes in solution.^{2d, 2f} Reported here is an approach using VT-DOSY to understand the self-assembly of organic π -conjugated oligomers, molecules studied recently in thin film photovoltaic devices, mutually by π stacking and hydrogen bonding.³ We believe that this ²⁵ characterization method is underutilized in a community where
- 25 characterization method is underutilized in a community where drawing relationships between molecular/supramolecular structure and ultimately device function is challenging but central to rational materials design.

We recently demonstrated how the photovoltaic power ³⁰ conversion efficiency of a branched oligothiophene (**BQPH**; Figure 1):fullerene blend was enhanced (twofold) through hydrogen bond promoted self-assembly relative to control devices fabricated from chromophores incapable of hydrogen bonding (e.g., **BQPME**).³ Elegant previous work of Lehn and

- ³⁵ Zimmerman inspired the molecular design that features the phthalhydrazide (**PH**) heterocycle, a building block shown to form robust hydrogen-bonded assemblies in solution and on surfaces, including cyclic trimers (i.e., (**PH**)₃).⁴ In the context of **BQPH**, we suspected that its putative H-bonded rosettes (i.e.,
- ⁴⁰ (**BQPH**)₃) might further organize into π -stacked columnar nanostructures (i.e., [(**BQPH**)₃]_n), an appealing chromophore arrangement (if extendable to thin films) for bulk heterojunction photovoltaics. Indeed, columnar arrangements of (**PH**)₃-based assemblies have been characterized before.^{4a} Herein, we provide
- ⁴⁵ supporting evidence of this mechanism of assembly in solution for **HexBQPH** by VT-DOSY NMR by employing a modified Stokes-Einstein equation that allows derivation of molecular size in solution for objects with non-spherical shape.^{1e} Facilitating the





analyses is comparison of the solution behavior of **HexBQPH** to ⁵⁵ two molecules incapable of H-bonded assembly, **HexBQPME** (a model of the monomer)³ and **HexB**⁵ (a covalent analogue of the H-bonded trimer).

A ¹H NMR spectrum of **HexBOPH**³ (10 mM) in the hydrogen bond promoting solvent toluene- d_8 (see Figure S1) shows two 60 broadened peaks at $\delta = 12.78$ and 13.75 ppm at 27 °C. The signals are consistent with the -NH and -OH protons of HexBQPH, respectively, in its lactim-lactam tautomeric form. That the peaks are well-separated and shifted far downfield suggest both H-bonded assembly and slow monomer-aggregate 65 exchange on the NMR time scale.^{4a} Also observed by ¹H NMR are significantly broadened and upfield shifted (up to 0.5 ppm) thiophene –CH resonances, indicative of π -stacking. In contrast, HexBQPME displays sharp peaks in the ¹H NMR spectrum under the same conditions (not shown). Upon warming through 70 90 °C, the -NH/-OH peaks of **HexBOPH** coalesce at ~ 85 °C (but remain significantly deshielded) and the aromatic signals sharpen (completely at ~ 75 °C) and shift downfield. The results are consistent with a transition from π -stacked. H-bonded assemblies to discrete H-bonded assemblies in solution.3



assemblies.

Prior to VT DOSY studies, an analogous 1-D VT ¹H NMR study was performed with comparator HexB (see Figure S2). The design of this compound was suggested by computational ¹⁰ modelling of the anticipated trimeric assembly of HexBQPH, that gave its approximate molecular dimensions (see Figures S3-S5). Preparation of phenyl cored thiophene dendrimer⁵ HexB came via three-fold Suzuki coupling of the boronic ester of **HexBQ**³ with 1,3,5-tris(5-bromothiophen-2-yl)benzene (see 15 Scheme S1).⁶ Compared to HexBQPH under identical conditions, the ¹H NMR spectrum of HexB shows some broadening and upfield shifting (up to ~ 0.15 ppm) of the thiophene protons (in the 6.9-7.3 ppm region) and benzene core protons (at 7.1 ppm) at room temperature; again the aromatic 20 proton peaks sharpen and move on average downfield upon raising the temperature. While the result may speak to some aggregation through π -stacking, the less pronounced chemical

- shift changes suggest weaker $\pi \pi$ association for **HexB** versus HexBQPH (vide infra).
- Obtained from DOSY NMR experiments are translational 25 diffusion coefficients (D) which are related to molecular size through the well-known Stokes-Einstein equation:

$$D = \frac{k_B T}{6\pi\eta r_h} \tag{1}$$

- where $k_{\rm B}$ is the Boltzmann constant, T is the temperature, $r_{\rm h}$ is the ³⁰ hydrodynamic radius, and η is the viscosity of the solvent.^{1e, 7} Equation 1 assumes that the diffusing entity is a sphere (Figure 2a) and is large relative to the van der Waals volume of the solvent.^{1e} Given these assumptions, the equation does not necessarily best describe the diffusion of the molecular entities 35 considered here, including the self-assembled aggregates of
- HexBQPH. Therefore, the equation has been augmented as discussed by Macchioni and coworkers^{1e} to include shape- and size correction factors. In modified form (Equation 2):

$$D = \frac{k_B T}{c f_s \pi \eta r_h} \tag{2}$$

$$_{40} c = \frac{6}{1 + 0.695 \left(\frac{r_{vdW}}{r_h}\right)^{2.234}}$$
(3)

$$f_{S} = \frac{\left[\sqrt{1 - {\binom{b}{a}}}\right]^{2}}{\left(\frac{b}{a}\right)^{\frac{2}{3}} \ln \left[\frac{1 + \left(\sqrt{1 - {\binom{b}{a}}}\right)^{2}}{\frac{b}{a}}\right]}$$
(4)

$$f_{S} = \frac{\sqrt{\left(\frac{b}{a}\right)^{2} - 1}}{\left(\frac{b}{a}\right)^{2} \tan^{-1} \sqrt{\left(\frac{b}{a}\right)^{2} - 1}}$$
(5)



Figure 3. Linearized [•] variable temperature diffusion data for 45 HexBQPME (24 mM; DMSO-d₆). The linear regression fit is shown for data derived from the prolate []] spheroid model and iteratively determined values of a, b, and $r_{\rm h}$.

where c is a size correlation factor between $r_{\rm h}$ of the diffusing species and the van der Waals radius $(r_{\rm vdW})$ of the solvent^{1e, $\overline{8}$} 50 (Equation 3), and f_s is a system-derived shape friction correction factor (Equations 4 and 5).^{1e, 9} The diffusing entities represented by f_s are mathematically modelled as prolate (cigar-shaped; Equation 4 and Figure 2b) or oblate (pancake-shaped; Equation 5 and Figure 2c) spheroids and are parameterized by the axial (a) ss and equatorial (b) axes of the respective spheroids (Figure 2).^{1e, 9} Alternatively, f_s can be modelled as a cylinder (Figure 2d) parameterized by the length (L) and diameter (d) (see the Supporting Information for details).¹⁰ An important consequence of using Equations 2 and 3 is that $r_{\rm h}$ now represents the 60 hydrodynamic radius of a sphere with equivalent volume to that

of the spheroid used to calculate f_{s} .^{1e}

Given Equation 2, for a monomeric species a plot of the diffusion coefficient D versus (T/η) should result in a straight line with a slope inversely related to c, f_s , and r_h . One can then use an $_{65}$ iterative fitting procedure to estimate physical parameters a and b(or L and d) from the data, given a priori knowledge and shape/size constraints offered by molecular modelling (see Figures S3-S5). The latter provides critical guidance and can afford, for example, initial values for a and/or b (L and/or d), an ⁷⁰ approximation of the molecule's aspect ratio (i.e., 2a/2b), and a starting ellipsoid shape model (Figure 2) for the fitting. The interdependence of several parameters (e.g., c is derived from $r_{\rm h}$; $f_{\rm s}$ is derived from b/a) means that only a, b, and $r_{\rm h}$ need to be iteratively varied. Combinations are sought that arrive at a 75 derived (fitted) slope that matches the experimental one (typically

within 0.2%). Then, the values of a and b can be proportionally varied such that the calculated volume of the spheroid matches (typically within 1%) a spherical value derived from $r_{\rm h}$. Worth noting, since the ratio of a to b directly determines f_s , proportional

⁸⁰ changes to both do not affect this parameter. Also clear from the treatment is that the diffusion behavior fitted via one model cannot be fitted using another model and that when selected, the prolate model will result in $f_{\rm s} < 1$, whereas the oblate model will result in $f_s > 1$. Worth noting is that compensatory effects s involving c and f_s can give rise to a cf_s product ~ 6 and apparent

agreement with the classical Stokes-Einstein equation.^{1e}

Page 2 of 4

1



Figure 4. Linearized variable temperature diffusion data for **HexBQPH** (20 mM; toluene- d_8). The slope change at ~ 60 °C is evident from the intersection of the solid and dashed linear regression fits.

The approach is illustrated in Figure 3 through an evaluation of **HexBQPME** in DMSO-*d*₆. Worth noting, temperature gradients within our samples were minimized by the use of a well designed heating and cooling system, and a convection correction was ¹⁰ implemented within pulse sequencing to ensure accurate measurements of diffusion coefficients.¹¹ Diffusion constants were recorded and plotted against (T/η) (Figure 3, dark circles); a linear fit provides the slope (i.e., $1.23 \times 10^{-15} \text{ m}^3/\text{kg}$) which is inversely related to *c*, *f*_s, and *r*_h. An initial estimate of *a* (13 Å) is ¹⁵ entered based on molecular modelling (see the SI for calculations), and an iterative fitting to determine a reasonable combination of *a*, *b*, and *r*_h ensues. In one fit (open squares), the prolate model yields values of *a* = 13 Å, *b* = 7.4 Å, and *r*_h = 9.0 Å, ultimately resulting in *f*_s = 0.69 and *c* = 5.8 (Table 1). Use of

²⁰ the oblate model for **HexBQPME** (which changes the value of f_8) does not fit the experimental data (data not shown).

Two solvent types were selected for variable temperature DOSY studies of **HexBQPH**. Results from toluene- d_8 , an assembly promoting solvent, can be evaluated against the VT-²⁵ NMR data discussed above. Studies in assembly suppressive (i.e., H-bond competitive) solvents, such as dimethyl sulfoxide- d_6 (DMSO- d_6), allow direct comparison with the monomeric controls, **HexBQPME** and **HexB**. Diffusion coefficients were

first collected for **HexBQPH** in DMSO- d_6 . Satisfyingly, fits of ³⁰ the diffusion coefficient versus temperature data using the prolate spheroid parameters provides good agreement (Table 1) with **HexBQPME**, expected given the similar molecular dimensions of the two based on modelling (see Figures S3–S5). The slightly larger apparent size of **HexBQPH** might be explained through its

strong hydrogen bonding with the solvent. Similar evaluation of **HexB** (in THF- d_8 due to solubility) using the oblate parameters provides values of *a* and *b* consistent with its molecular dimensions derived from modelling and an unaggregated species. Next, variable temperature diffusion data was collected in

⁴⁰ toluene- d_8 for **HexBQPME**, **HexBQPH**, and **HexB**. Evidence for the aggregation of **HexBQPH** is immediately provided through its small diffusion coefficient $(1.03 \pm 0.01 \times 10^{-10} \text{ m}^2/\text{s})$ relative to **HexBQPME** $(5.90 \pm 0.06 \times 10^{-10} \text{ m}^2/\text{s})$ at 22 °C.

 Table 1. Size and shape parameters for HexBQPH, HexBQPME, and

 45
 HexB monomers based on VT-DOSY data.

Molecule	Model ^c	$a (\text{\AA})^d \\ [a_{ ext{calcd}}]^e$	$b (\text{\AA})^d \\ [b_{ ext{calcd}}]^e$	p^{f}	$r_{\rm h}({\rm \AA})^d$	$f_{\rm s}$	с
HexBQPME ^a	Р	13 [13]	7.4	1.8	9.0	0.69	5.8
HexBQPH ^a	Р	15 [12]	8.1	1.9	10	0.71	5.8
HexB ^b	0	4.7	32 [24]	6.9	17	1.3	5.9

^{*a*} Based on diffusion coefficients measured in DMSO- d_6 (**HexBQPH** = 27 mM; **HexBQPME** = 24 mM). ^{*b*} Based on diffusion coefficients measured in THF- d_8 (**HexB** = 30 mM). ^{*c*} O = oblate; P = prolate. ^{*d*} Estimated fitting error: $\pm 10\%$ for *a*, *b*, and $r_{\rm h}$. ^{*e*} Calculated parameters are estimated from 50 molecular modelling (see the SI for details). ^{*f*} *p* = (semimajor axis)/(semiminor axis).

Table 2. Size and shape parameters for **HexBQPH**, **HexBQPME**, and **HexB** in toluene- d_8 based on VT-DOSY data.^{*a*}

Molecule	Model ^d	a (Å) ^e	$(\text{\AA})^e$	$L(\text{\AA})^e$	d (Å) ⁶	$r_{\rm h}$ $({\rm \AA})^e$	$f_{\rm s}$	с
HexBQPME	Р	14	7.8			9.5	0.69	5.7
$HexBQPH^b$	0	26	28			27	1.0	6.0
$HexBQPH^b$	С			38	51	27	1.0	6.0
HexBQPH ^c	0	5.0	26			15	1.2	5.9
HexBQPH ^c	С			7.0	51	15	1.2	5.9
HexB	0	4.5	30			16	1.3	5.9
HexB	С			7.8	58	17	1.2	5.9

⁵⁵ ^a Based on diffusion coefficients dtermined at the following concentrations: HexBQPH = 20 mM; HexBQPME = 20 mM; HexB = 18 mM. ^b Based on data acquired at low temperature (21.8–52.6 °C). ^c Based on data acquired at higher temperature (58.8–92.2 °C). ^d C = cylinder; O = oblate; P = prolate. ^e Estimated fitting error: ± 10% for *a*, *b*, 60 *L*, *d*, and *r*_h.

Linearization of the diffusion versus temperature data and iterative fitting was first performed on **HexBQPME**. Using the prolate spheroid model, values of a = 14 Å, b = 7.8 Å, and $r_h = 9.5$ Å could be derived (Table 2), values which are similar to ⁶⁵ **HexBQPH** and **HexBQPME** in DMSO-*d*₆ (Table 1). The data is consistent with the conclusion that **HexBQPME** is monomeric in both DMSO-*d*₆ and toluene-*d*₈, and that **HexBQPH** is monomeric in DMSO-*d*₆.

Upon linearization of the diffusion coefficient data for ⁷⁰ **HexBQPH** (Figure 4), bimodal behaviour is found reflected by the distinct slope change observed at approximately 60 °C. The temperature, based on 1-D VT ¹H NMR data (vide supra), is consistent with a transition from π -stacked aggregates to discrete H-bonded aggregates.¹² That the dynamic assembly process ⁷⁵ likely involves conversion of columnar π -stacks of trimers (i.e., [(**HexBQPH**)₃]_n) to non- π -stacked trimeric assemblies (i.e., (**HexBQPH**)₃) comes through iterative fitting of the two temperature ranges (21.8–52.6 °C and 58.8–92.2 °C) and comparison to **HexB**. Using first the oblate model for ⁸⁰ parameterization, the low temperature regime returns *a* = 26 Å and *b* = 28 Å (r_h = 27 Å) while the high temperature regime provides *a* = 5.0 Å and *b* = 26 Å (r_h = 15 Å). The equatorial radius (*b*) is very similar across both ranges and further similar to the calculated end-to-end length of **HexBQPH** (25 Å; see SI for details). Shown is that upon increasing the temperature, **HexBQPH** experiences a significant size change *only* in one dimension (from a = 26 Å to a = 5.0 Å); this is consistent with s columnar growth. Confirmation that the high temperature

- assembly is highly represented by a discrete trimer comes through comparison with **HexB**. A plot of *D* versus (T/η) for **HexB** in toluene-*d*₈ gives a linear fit, suggesting that there is little extended aggregation of this species under these conditions (vide
- ¹⁰ supra). Iterative parameter fitting using the oblate spheroid model provides a = 4.5 Å, b = 30 Å, and $r_h = 16$ Å. These values indeed match well with both the modelled **HexB** structure (see Figure S5) and the high temperature values for **HexBQPH**.
- To verify our analysis, the diffusion data for **HexBQPH** and ¹⁵ **HexB** in toluene- d_8 were alternatively subjected to iterative fitting using a cylindrical model (i.e., treatment of the diffusing entity as a cylinder). Consistent results are obtained (Table 2), where for **HexBQPH** only a change in the length (*L*) of the cylindrical aggregates is observed as a function of temperature (21.8–52.6 ²⁰ °C: *L* = 38 Å, 58.8–92.2 °C: *L* = 7 Å), while the diameter (*d*)
- remains roughly the same. The **HexB** cylindrical parameters (L = 7.8 Å; D = 58 Å) are both consistent with those derived from the **HexB** oblate fitting and for **HexBQPH** at higher temperatures.
- The molecularity of the **HexBQPH** aggregates can be estimated by appropriately scaling the volumes derived from the $r_{\rm h}$ values.^{1e} Monomeric **HexBQPH** and **HexBQPME** have $r_{\rm h} \sim$ 9.5 Å (and equivalent sphere volume of ~ 3600 Å³) and are assigned a molecularity of 1. Both spheroid fitting routines of **HexBQPH** in toluene- d_8 ($r_{\rm h} = 27$ Å) yield an average
- ³⁰ molecularity of 23 on the basis of equivalent sphere volume for the low temperature regime, which corresponds to an average supramolecular assembly of ~ 6 trimeric discs. The r_h value of 15 Å (from both fitting routines) for the high temperature regime equates to an average molecularity of 3.9. This value, when
- a_{35} coupled with the shape parameters *a* and *b* (or *L* and *d*) and molecular modelling results, strongly supports the presence of a discotic trimeric aggregate.

Conclusions

- In conclusion, an iterative method for size and shape ⁴⁰ approximation involving variable temperature diffusion measurements has been exploited for the characterization of organic π -conjugated chromophore assembly in solution. Diffusion coefficients were collected via DOSY NMR, linearized, and fitted to a modified Stokes-Einstein equation
- ⁴⁵ using prolate spheroid, oblate spheroid, and cylindrical models. **HexBQPH** self-assembly showed a bimodal temperature response in toluene- d_8 , consistent with a transition from columnar assemblies to isolated H-bonded trimers based on comparisons with non-assembling model compounds. We are hopeful that this
- ⁵⁰ approach can serve as a complementary way to explore dynamic supramolecular assembly in solution and facilitate drawing relationships between structure and function in applicationoriented organic systems.

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60 Notes and references

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110