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**Title**

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**Journal**

Macromolecules, 50(6)

**ISSN**

0024-9297

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**Publication Date**

2017-03-28

**DOI**

10.1021/acs.macromol.6b02410

Peer reviewed

# Identifying Atomic Scale Structure in Undoped/Doped Semicrystalline P3HT Using Inelastic Neutron Scattering

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## Abstract

2  
3 The greatest advantage of organic materials is the ability to synthetically tune  
4 desired properties. However, structural heterogeneity often obfuscates the relation-  
5 ship between chemical structure and functional properties. Inelastic neutron scattering  
6 (INS) is sensitive to both local structure and chemical environment and provides atomic  
7 level details that cannot be obtained through other spectroscopic or diffraction meth-  
8 ods. INS data is composed of a density of vibrational states with no selection rules,  
9 which means that every structural configuration is equally weighted in the spectrum.  
10 This allows the INS spectrum to be quantitatively decomposed into different structural  
11 motifs. We present INS measurements of the semiconducting polymer P3HT doped  
12 with F4TCNQ supported by density functional theory calculations to identify two dom-  
13 inant families of undoped crystalline structures, and one dominant doped structural

14 motif, in spite of considerable heterogeneity. The differences between the undoped and  
15 doped structures indicate that P3HT side chains flatten upon doping.

## 16 **Introduction**

17 Recent progress in polymer based organic devices has led to significant milestones in perfor-  
18 mance, such as organic photovoltaic (OPV) power efficiencies of  $>10\%$ <sup>1,2</sup> and electron and  
19 hole mobilities  $>1 \text{ cm}^2\text{V}^{-1}\text{s}^{-1}$ .<sup>3,4</sup> However, despite the successes of these materials, it is dif-  
20 ficult to precisely identify the microstructural details that lead to desired performance. The  
21 structural and electronic details of organic electronic materials in particular are challenging  
22 to characterize and describe due to high levels of structural and electronic heterogeneity at  
23 length scales from Angstroms to micrometers.<sup>5-8</sup> In contrast to inorganic semiconductors,  
24 polymeric semiconductors do not follow the paradigm that increased crystalline order and re-  
25 peating texture necessarily leads to improved conductivity.<sup>9</sup> For polymeric semiconductors,  
26 a low degree of dihedral variation can lead to highly delocalized and anisotropic charged  
27 states in materials with little or no long range crystalline order.<sup>10</sup>

28 The determination of a detailed atomic structure is challenging because the weak bonding  
29 between polymer chains in the film does not encourage long range order, allowing for many  
30 structural configurations with similar energetic minima. The lack of long range order limits  
31 the information that can be obtained from traditional structural characterization techniques  
32 such as XRD, which probes ordered domains.<sup>11-18</sup> Scattering techniques such as small angle  
33 neutron or X-ray scattering (SANS or SAXS) return average distance correlations but fail  
34 to determine specific geometries.<sup>19-21</sup> Spectroscopic techniques, such as pulsed laser experi-  
35 ments, can provide information about the energetic distribution of heterogeneous structures,  
36 but cannot link this distribution to any specific molecular geometry.<sup>22-26</sup> Even electron mi-  
37 croscopy techniques, which can determine the location of a specific atom within a lattice to  
38 within  $<0.1 \text{ \AA}$ , cannot distinguish between adjacent carbon atoms (due to limited contrast),

39 and hydrogen atoms are invisible to most microscopy techniques.<sup>27–29</sup>

40 Analogous to inorganic semiconductors, polymeric semiconductors can be doped for a  
41 conductivity increase of up to 11 orders of magnitude.<sup>30</sup> However, unlike inorganic semi-  
42 conductors, many polymer–dopant mixtures have low doping efficiencies, which requires in-  
43 creased doping ratios to achieve desired conductivity increases. Such high doping ratios cause  
44 structural rearrangements that affect the alignment of energy levels and transfer integrals,  
45 which may negatively impact the macroscopic electrical properties. For example, films of  
46 poly(3–hexylthiophene) (P3HT) and 2,3,5,6-Tetrafluoro-7,7,8,8-tetracyanoquinodimethane  
47 (F4TCNQ) have different morphologies at different doping ratios, and the conductivity does  
48 not increase predictably with dopant concentration<sup>31–33</sup> demonstrating the complex effect  
49 that structural rearrangement has on doping efficiency and conductivity. At high doping  
50 concentrations, F4TCNQ intercalates between P3HT chains in crystalline regions creating  
51 a new phase, which coincides with changes in the dependence of F4TCNQ concentration  
52 on conductivity.<sup>31</sup> Clearly, the local structure of the P3HT–F4TCNQ mixture has a large  
53 effect on electrical properties, but very little is known about the unit cell structure at atomic  
54 length scales.

55 In this context, we highlight inelastic neutron spectroscopy (INS) as an appropriate mea-  
56 surement to infer structural motifs within films of doped or undoped P3HT. Unlike electro-  
57 magnetic radiation, neutrons do not interact with mobile charges present in doped materials.  
58 For this reason, the observed INS spectra only contains peaks from nuclear vibrations, and  
59 not the dielectric response of the P3HT polaron, which simplifies the comparison between  
60 doped and undoped P3HT. The INS technique hinges on resolving the time–of–flight of  
61 scattered neutrons to calculate the energy loss of the neutron after the transfer of neutron  
62 energy to a vibrational mode of the sample. The measurement provides the vibrational den-  
63 sity of states weighted by the incoherent neutron cross section and vibrational amplitude.  
64 INS provides different information than FTIR and Raman spectroscopy because of the dif-  
65 ferent atomic cross sections for neutrons and electromagnetic radiation. Scattering from all



66 hydrogen vibrations is visible using neutrons, but only vibrations that cause a change in  
67 dipole moment (such as C–H stretch vibrations) are also detectable in FTIR/Raman spec-  
68 troscopy, as shown in supplemental Figures S2a,b. Those selection rules are not present in  
69 inelastic neutron scattering processes.<sup>34</sup> FTIR has further restrictions as the only allowed  
70 transitions must be fundamental (0–1) within the harmonic approximation. This restriction  
71 leads to sharper, better defined peaks than INS because INS allows simultaneous excitation  
72 of multiple vibrational modes, which broadens observed peaks.

73 To illustrate this principle, consider a system with a single high energy mode (e.g.  
74 C–H stretch), and a distribution of low energy modes (e.g. torsional vibrations between  
75 monomers). An incoming neutron may excite the high energy mode and a random low  
76 energy mode simultaneously, causing the neutron to lose energy proportional to the sum  
77 of the frequencies of the excited modes, which leads to a distribution of observable energy  
78 transfers that broadens observed peaks. An incoming photon is allowed to excite one of the  
79 modes (assuming there is a dipole), leading to a sharp, well-defined peak that presents an  
80 advantage of FTIR over INS: the peaks that are observable for FTIR are better defined than  
81 INS peaks. This concept is illustrated when considering the C–H stretch peaks observed in  
82 FTIR (SI Figure S2) versus INS (SI Figure 1). A major disadvantage of FTIR is the lack  
83 of peaks relative to INS spectra that come from bend, wag, and other complex motions, as  
84 can be seen in supplemental Figures S2b,c. Finally, electromagnetic radiation interacts with  
85 free electrons, yielding a large background signal in highly conductive samples like doped  
86 P3HT (see supplemental figure S2c). The information in an INS scattering profile can be  
87 decomposed into a linear combination of contributions from dominant structural themes.  
88 In comparison, FTIR provides structural information of the sample, but the selection rules  
89 could provide different weightings on that information from different structural configura-  
90 tions. As a result, an average structure can be computed from FTIR, but it is not possible  
91 to decompose the spectrum into different populations.<sup>35</sup> In contrast, INS provides informa-  
92 tion about the full distribution of structural configurations weighted by their density in the

93 sample.<sup>34</sup> Thus, vibrational neutron spectroscopy is complementary to FTIR/Raman, but  
94 also yields significant new information and is much better suited to investigate conductive  
95 and heterogeneous samples.

96 Because the INS spectrum has no selection rules and is a quantitative measurement of  
97 the vibrational density of states, it can be decomposed into multiple populations. Each  
98 population can then be modeled separately with high accuracy using density functional  
99 theory (DFT).<sup>36,37</sup> Thus, the experimental spectra can be matched to a combination of  
100 proposed structures whose spectral contributions are determined from DFT. In addition,  
101 the charge distribution calculated using the DFT model can be constrained by fitting to the  
102 measured data, which has numerous modes. In this way, the structure of the configuration  
103 in the sample is inferred from the dynamics of the atoms in the film, which is dependent on  
104 the charge distribution. Configurations that do not adequately represent the experimental  
105 INS spectra can be summarily removed.

106 Previously, INS has been used to study the vibrational dynamics and structure of poly-  
107 mers. Early studies used INS to understand the low frequency dynamics of poly-acetylene,<sup>38</sup>  
108 polyaniline,<sup>39</sup> and PPV<sup>40</sup> demonstrating that INS provides valuable low-energy dynamic in-  
109 formation for conjugated polymers that could be important for understanding charge trans-  
110 port. However, the short range dynamic information was missing from the spectra due to the  
111 low neutron flux and small energy range of early INS instruments; instrument resolution de-  
112 creased with increasing energy transfers, meaning that short range dynamic information was  
113 missing from the spectra. Current INS spectrometers, including VISION at the Spallation  
114 Neutron Source,<sup>41,42</sup> TOSCA at the ISIS laboratory,<sup>43</sup> and LAGRANGE at the Institute  
115 Laue-Langevin,<sup>44</sup> are capable of providing high-resolution spectra across the entire vibra-  
116 tional energy range (5 meV – 1 eV). Thus, these spectrometers can observe long-range,  
117 low energy lattice dynamics, and high-energy, short range bond vibrations simultaneously.  
118 With these high-resolution instruments, the local structure of polymers has been inferred  
119 from INS spectra because the vibrational modes sensitively depend on the spatial arrange-

120 ment of neighboring atoms/molecules. For instance, the TOSCA spectrometer was used to  
 121 identify the zig-zag structure of PEO under confinement,<sup>45</sup> as well as the local structure of  
 122 adsorbed hydrogen on a nickel catalyst.<sup>46</sup> On the other hand, the VISION spectrometer was  
 123 recently used to identify the complex structure of highly disordered kerogen<sup>47</sup> demonstrating  
 124 that high-resolution INS is capable of providing structural as well as dynamic information.  
 125 Hermet et al. demonstrated that INS can also be used to identify the presence of different  
 126 crystalline polymorphs of quaterthiophene and sexithiophene.<sup>48</sup>

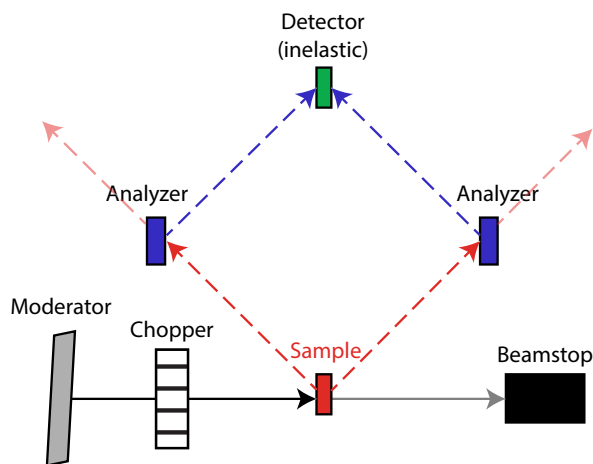


Figure 1: *Schematic representation of the VISION spectrometer. The analyzers determine the final neutron energies, which are used to find the amount of energy transferred to the sample by recording and analyzing the time of flight of the neutrons from each pulse.*

127 Our study builds on prior studies by demonstrating that INS can be used to study the  
 128 structure of both doped and undoped disordered organic electronic materials. We demon-  
 129 strate here that the distribution of configurations contained in heterogeneous films of doped  
 130 and undoped P3HT can be resolved into contributions from dominant structural motifs  
 131 that we can identify using the VISION spectrometer and DFT. We decompose the undoped  
 132 spectrum into contributions from the crystalline and amorphous regions to find three key  
 133 structural motifs in the film, one in the amorphous region, two in the crystalline region.  
 134 We also find a dominant structural motif of a P3HT film that has been heavily doped by  
 135 F4TCNQ, producing a crystal structure of P3HT with intercalated F4TCNQ that agrees

136 with both theory and experiment.

## 137 **Materials and Methods**

### 138 **Materials**

139 Regioregular P3HT ( $>98\%$ ,  $M_n = 54 - 75$  kDa), regiorandom P3HT and F4TCNQ ( $\geq 97\%$ )  
140 were purchased from Sigma–Aldrich. Undoped samples were used as received by the manu-  
141 facturer in INS experiments unless otherwise indicated. For the doped samples, P3HT and  
142 F4TCNQ were first dissolved in chloroform. Solutions were then mixed in appropriate ratios  
143 to achieve the desired doping levels, and the solvent was evaporated inside a nitrogen glove-  
144 box at low temperature over several days. The solids were chopped into a powder (500 mg  
145 for each sample) using a razor blade and loaded into rectangular aluminum sample holders  
146 and sealed for measurement.

### 147 **Inelastic Neutron Scattering**

148 We used the VISION spectrometer at the Spallation Neutron Source at Oak Ridge National  
149 Laboratory (Figure 1) to characterize the ensemble of structural configurations present in our  
150 doped and undoped P3HT samples. The VISION spectrometer is ideally suited for inferring  
151 structural information from polymer films. Figure 1 shows the experimental setup of the  
152 VISION spectrometer. The analyzers and the Be filters select the final neutron energy loss.  
153 Time–of–flight analysis allows for computation of the energy spectrum transferred from the  
154 incident white beam of neutrons to the polymer film. The experimental set up improves  
155 the count rate and signal to noise ratio over other INS instruments by over two orders of  
156 magnitude, allowing rapid acquisition of full spectrum data.<sup>41</sup> To minimize the Debye–Waller  
157 effect, which causes broadening of the spectral information due to random fluctuations of  
158 the atomic position, the scans were performed at 5 K. This low temperature provides the  
159 level of detail required to probe sensitive structural changes during doping. It is possible

160 that the low temperature of the measurement could induce crystalline phase transformation.  
161 However, we do not expect major morphological changes in P3HT, because main chain and  
162 side chain melting occurs at  $\sim 200^\circ\text{C}$  and  $\sim 50^\circ\text{C}$ , respectively.<sup>49,50</sup> In addition, Poelking  
163 et al used molecular dynamics to show that polymorph I' does not reform from polymorph  
164 I after cooling.<sup>51</sup> Large scale reorganization would require mobile main chains and/or side  
165 chains, which is unlikely as the temperature is lowered. Nevertheless, it remains possible  
166 that temperature-induced structural reorganization is taking place.

## 167 Computational Methods

168 We used plane-wave density functional theory with norm-conserving pseudopotentials and  
169 the PBE functional<sup>52,53</sup> to calculate the optimized structures within the CASTEP frame-  
170 work.<sup>54</sup> PBE is a common choice for understanding the structure of molecular crystals,  
171 including quaterthiophene and sexithiophene, which are chemically similar to P3HT.<sup>48,55,56</sup>  
172 Density functional perturbation theory (DFPT) was used to calculate the phonon spectrum  
173 of the optimized structures instead of finite difference methods because DFPT allows com-  
174 putation of vibrational modes at multiple  $k$ -points whereas the finite difference method is  
175 restricted to the gamma point only. The initial structures for the simulations were created  
176 within the Materials Studio<sup>®</sup> suite. The lattice parameters were determined from published  
177 X-ray diffraction,<sup>31</sup> and molecular dynamics studies.<sup>51</sup> The atomic coordinates were relaxed  
178 so that the maximum interatomic force was below  $0.01 \text{ eV}/\text{\AA}$ . We used a  $2 \times 3 \times 3$  and a  $1 \times$   
179  $2 \times 2$   $k$ -point sampling grid to simulate the undoped and doped configurations, respectively.  
180 More  $k$ -points were used to sample longer dimensions in the Brillouin zone (smaller real space  
181 lattice dimensions). We used aCLIMAX 6.0<sup>57</sup> to convert the finished phonon calculations to  
182 INS spectra that can be compared to experiments. The details of the conversion from vibra-  
183 tional modes to an INS spectrum are found elsewhere.<sup>34,57</sup> All simulated frequencies were  
184 scaled by a constant factor (1.015 in this study) to better agree with the experiments. The  
185 use of a scaling factor has been used in many prior DFT studies to replicate experimental

186 vibrational modes.<sup>46,58,59</sup>

187 Since the initial geometries were relaxed prior to calculating vibrational frequencies, the  
188 relaxed configuration commonly differed greatly from the initial geometry, which led to  
189 problems with using experimental unit cells as starting points. We altered the dihedral  
190 angles of the first bond of the side chain, and the displacement of thiophene rings in the  
191 direction of the backbone until we achieved a relaxed geometry that matched configurations  
192 determined from X-ray data.

193 We iteratively adjusted parameters such as the dihedral angles between thiophene rings,  
194 the dihedral angle of the first bond of the side chain and the displacement of one chain  
195 relative to another chain to generate a set of configurations likely to represent the data  
196 and consistent with previously published crystallographic data. Likely configurations were  
197 chosen because the associated simulated spectra showed peaks at the correct frequencies  
198 compared to experimental data. This means that only configurations that were matched to  
199 both published crystallographic and our INS data are presented.

## 200 **Results and Discussion**

### 201 **INS Experimental Data**

202 Regio-regular P3HT is a semicrystalline polymer that forms films with 20–50% crystalline  
203 and 50–80% amorphous phase fractions.<sup>9,50,60</sup> The molecular structures of P3HT and F4TCNQ  
204 are shown in Figure 2b,e. When F4TCNQ comes into contact with an electron donor, like  
205 P3HT, charge transfer to F4TCNQ results in a positive charge that is delocalized along  
206 the polymer backbone which changes a large number of molecular vibrations visible to high  
207 resolution INS. In addition, the crystalline portion of P3HT has been reported to show sig-  
208 nificant paracrystallinity, which means that the individual monomers experience significant  
209 structural heterogeneity even within crystalline domains.<sup>9,51</sup> The total structural heterogene-  
210 ity is expressed in the INS spectrum as a weighted sum of all different contributions, which

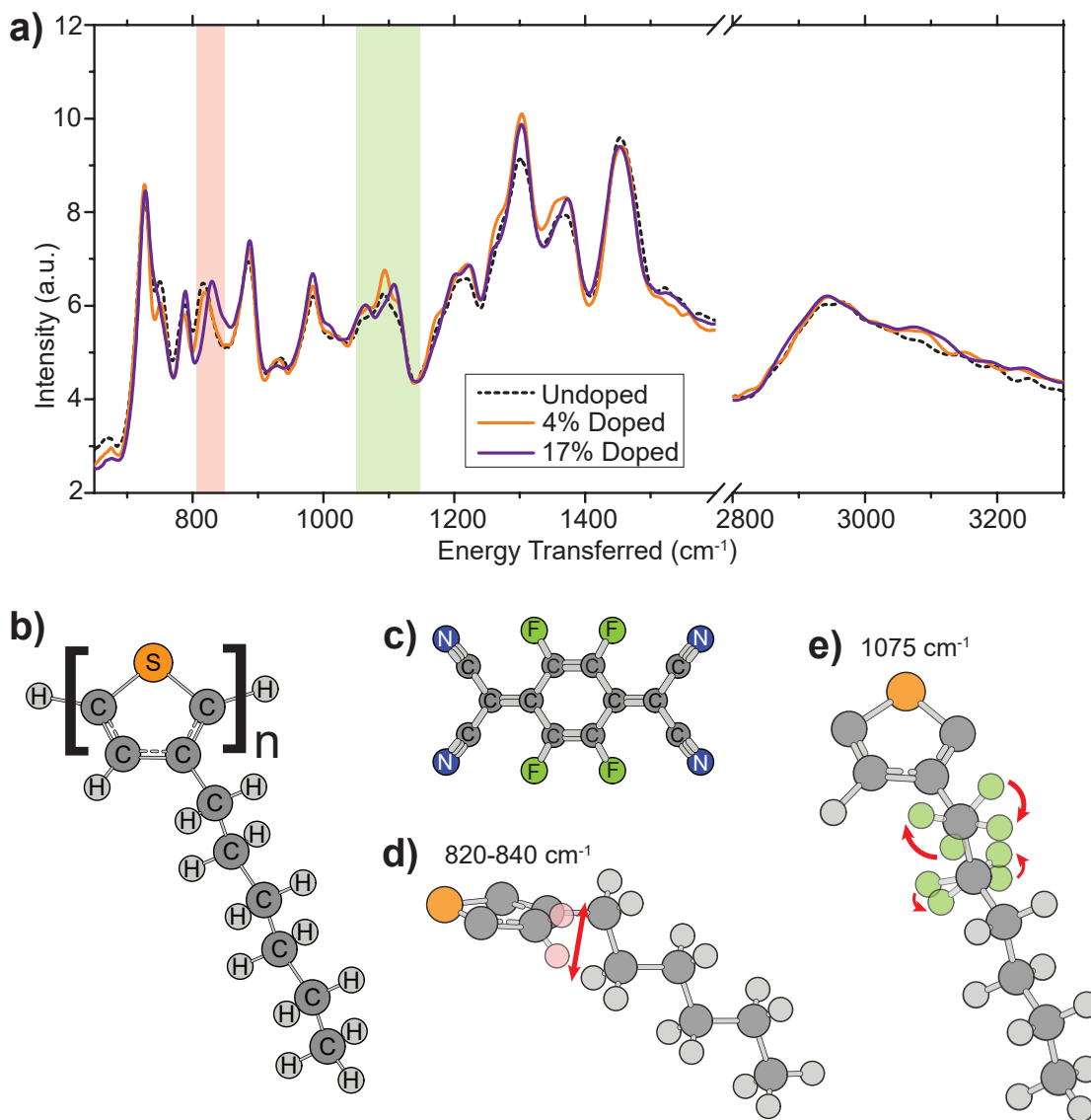


Figure 2: a) Experimental INS spectrum from 650 – 3300  $\text{cm}^{-1}$  of undoped P3HT, P3HT doped with 4 mol% F4TCNQ, and P3HT doped with 17 mol% F4TCNQ. The highlighted regions correspond to vibrational modes depicted in d,e. b,c) Molecular structures of P3HT and F4TCNQ. d,e) Vibrational modes that are most sensitive to doping from F4TCNQ. We show an out of plane bending mode of the backbone hydrogen at 820 – 840  $\text{cm}^{-1}$  (d), and a torsion between the first and second side chain carbons at 1075  $\text{cm}^{-1}$  (e).

211 leads to peak width (along with dispersion and phonon wings) and even the presence of sev-  
 212 eral different peaks originating from identical motions in differing molecular environments  
 213 (for example in crystalline and amorphous domains).

214 The vibrational spectra in Figure 2a represent samples of undoped, 4 mol% doped, and

215 17 mol% doped regio-regular P3HT with F4TCNQ. The spectra were normalized to equal  
216 background intensity. These doping levels were chosen because 17 mol% doping was reported  
217 to be the highest possible doping level (without formation of a pure F4TCNQ phase) for  
218 P3HT by F4TCNQ.<sup>31</sup> Doping at 4 mol% represents a lightly doped material in which most  
219 of the F4TCNQ is expected to reside in the amorphous domains of the polymer.<sup>31,33</sup>

220 Each peak corresponds to a vibrational transition that occurs when an incident neutron  
221 interacts with the nuclei in the sample. The peak height is proportional to the amplitude  
222 of the vibrational motion, and the energy of each transition reflects the frequency of the  
223 vibrational mode. Changes in energy of the same peak in different samples stem from  
224 changes in inter-atomic forces causing a particular mode to either stiffen (shift to higher  
225 energy) or relax (shift to lower energy). The changes in the height and frequency of a given  
226 peak may come from compositional changes to the ensemble of configurations present, as  
227 well as the reasons provided above. We model the data using DFT well enough to assign  
228 these peaks to particular vibrational modes, which agrees with prior studies.<sup>49</sup> We highlight  
229 two peaks, denoted by shaded regions in Figure 2a, most affected by doping, corresponding  
230 to the two vibrational modes in Figure 2d,e. The peaks are associated with the out of plane  
231 (OOP) bending of the backbone hydrogen, and the torsional bending of the first/second  
232 carbons in the side chain occur at  $\sim 820\text{--}840\text{ cm}^{-1}$ , and  $\sim 1075\text{ cm}^{-1}$ , respectively.

233 The OOP bending mode of the backbone hydrogen stiffens when heavily doped. The  
234 17 mol% doped sample displays a blue shift in this peak compared to the 4 mol% doped and  
235 undoped P3HT spectra. This blue shift is likely due to the intercalation of F4TCNQ into the  
236 crystalline regions at high doping ratios, in which the reduced distance between F4TCNQ  
237 and P3HT increases the intermolecular forces acting on the OOP mode. At 4 mol% doping  
238 ratio, F4TCNQ primarily remains in the amorphous regions of the polymer. Two prior  
239 studies and the lack of blue shift in the OOP bending peak corroborate this assignment.<sup>31,33</sup>

240 The torsional vibrations of the first and second carbons of the side chain are very sensitive  
241 to the torsional angle between the side chain and backbone of P3HT. Using DFT (discussed



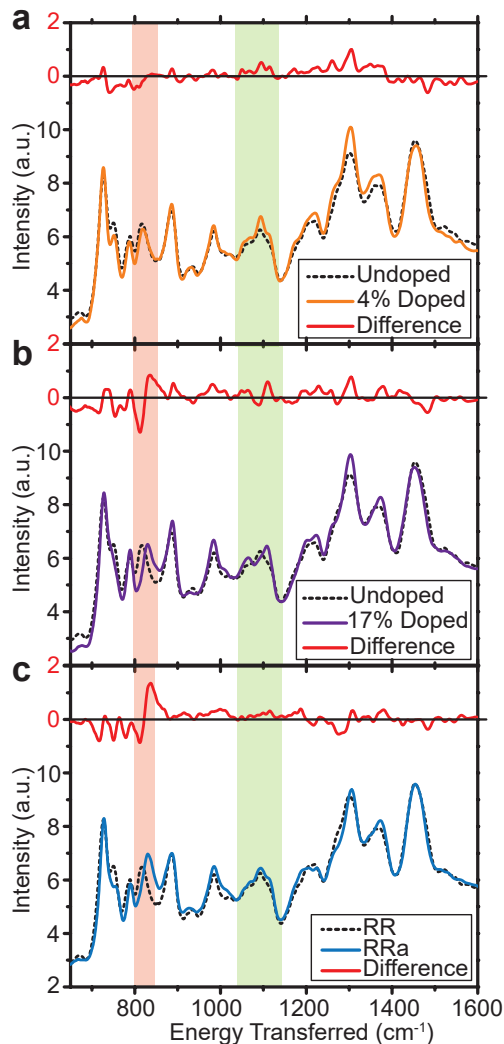


Figure 3: a) The regio-regular (RR) undoped and 4 mol% RR-doped spectra are compared between 650–1600  $\text{cm}^{-1}$ . The red curve corresponds to the difference between the 4 mol% doped curve and the undoped curve. b) The RR-undoped and 17 mol% RR-doped spectra are compared between 650–1600  $\text{cm}^{-1}$ . The red curve is the difference between the 17 mol% doped curve and the undoped curve. c) The RR-undoped and regiorandom (RRa) undoped spectrum are compared between 650–1600  $\text{cm}^{-1}$ . The highlighted regions correspond to vibrational modes depicted in Figure 2. The red curves above the spectra in a–c are the differences between the spectra.

242 later), we see that the change in torsional angle causes two overlapping peaks at 1075  $\text{cm}^{-1}$   
 243 to shift to higher and lower energies at high doping levels. The shift occurs as a larger  
 244 proportion of P3HT configurations contain side chains in the plane of the backbone as we  
 245 will show below. The flattening of side chains agrees with X-ray diffraction showing an

246 increase in the long axis of the unit cell upon doping.<sup>31</sup> Our DFT results also show that  
247 when a P3HT side chain is flat with respect to its backbone, the torsional mode blue shifts,  
248 and a coupled carbon-carbon stretching mode along the length of the side chain slightly  
249 increases in amplitude causing an increase in intensity (see the Supplementary Information  
250 for a visualization of the modes).

251 The spectrum also shows that the peak at  $1300\text{ cm}^{-1}$  increases in amplitude when compar-  
252 ing undoped and 4% doped. Using our DFT results, we assigned this peak as a combination  
253 of many twisting modes of the side chain hydrogens. The increase in peak height could be due  
254 to a combination of many factors. In general, changes in peak height are due to changes in  
255 vibrational amplitude and/or compositional changes in the polymer. Compositional changes  
256 include local configurational changes, and changes in the amorphous/crystalline volume  
257 fractions. Doping P3HT results in a charged backbone, which could cause configurational  
258 changes, and changes in intramolecular forces, both of which can affect the vibrational mode  
259 energies and amplitudes. It seems that this peak is unaffected by changes in amorphous  
260 volume fraction because this peak does not show a change in height when comparing un-  
261 doped RRa and RR P3HT. In addition, prior X-ray studies show only a small amount of  
262 structural reorganization when comparing undoped and 4% doped, so it would make sense  
263 that charging the backbone is responsible for the increase in peak height (either through  
264 changes in local configuration or increase in vibrational amplitude). However, we cannot  
265 make any conclusions about the cause of the increase in this peak's height at this time.

266 Figure 3 shows the compositional dependence of the INS spectra. The spectral differences  
267 between undoped and 17% doped are clearly shown in Figure 3b, where the OOP bending  
268 peak shifts to higher energy, and the overlapping modes in the green region shift into two  
269 distinct peaks. In contrast, Figure 3a shows almost no spectral differences between undoped  
270 and 4% doped samples. The difference curves in Figure 3a,b reinforce the contrast between  
271 the two doping levels. A doping ratio of 4 mol% does not affect the vibrational structure as  
272 much as a doping ratio of 17 mol%, demonstrating that the local structure is changed most

273 in a heavily doped sample.

274 The spectra in Figure 3 represent samples that have both amorphous and crystalline  
275 phases, which convolute our interpretation of the spectra because we can't explicitly attribute  
276 a spectral change between samples to a change in intermolecular forces (small scale) versus a  
277 change in phase volume fractions (large scale) because doping likely changes the crystalline  
278 phase fraction. Thus, we need a sample with a single phase to distinguish between crystalline  
279 and amorphous configurations. Regio-random (RRa) P3HT has an identical chemical struc-  
280 ture to regio-regular (RR) P3HT, however the hexyl side chain is randomly attached to  
281 either the second or third carbon of the thiophene ring, which generates films that do not  
282 show an X-ray crystal structure and therefore are uniformly amorphous.<sup>61-63</sup> Thus, we can  
283 study the vibrational structure of the amorphous region independently, assuming that the  
284 random change in side chain attachment does not strongly affect the vibrational modes of  
285 the polymer.

286 Figure 3c shows a comparison of the INS spectra of undoped RR-P3HT and RRa-P3HT,  
287 with the difference between the spectra plotted above. This comparison demonstrates the  
288 vibrational difference between a composite amorphous/crystalline system (RR-P3HT), and  
289 a uniformly amorphous system (RRa-P3HT). The OOP bending region at 820-840  $\text{cm}^{-1}$   
290 is the only peak with a vibrational frequency that is strongly affected by the change in  
291 phase composition, indicating that the OOP bending mode is relaxed by more delocalized  
292 electronic states present in crystalline regions of polymer chains. Electronic delocalization is  
293 a result of decreased dihedral angle variation between backbone monomers.

## 294 **Simulation of Crystalline Domains**

295 Plane wave density functional theory is the best approach to model the INS data as it al-  
296 lows the accurate prediction of vibrational modes in crystalline samples. The periodicity  
297 implicit in using a plane wave basis set allows the simulations to model extended solids,  
298 which provides the intermolecular interactions absent in typical gas phase DFT simulations.

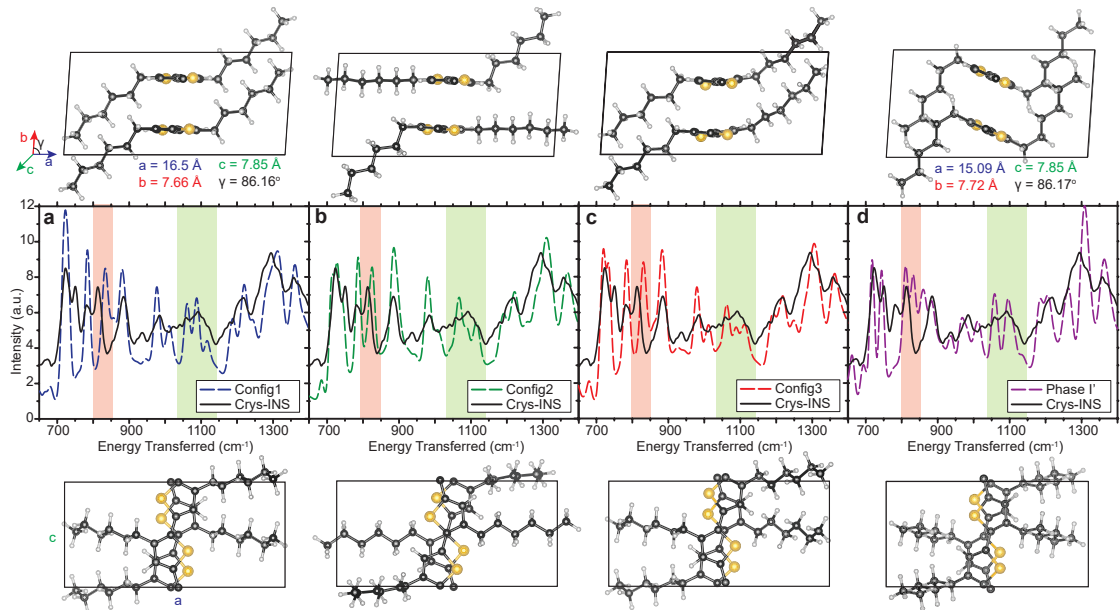


Figure 4: Simulations of three different potential P3HT crystalline configurations (Config1(a), Config2(b), Config3(c), and Phase I'(d)). In all cases, the simulated spectrum is plotted against the experimental INS spectrum of undoped P3HT between 650–1400  $\text{cm}^{-1}$ . The side view (top row) and top view (bottom row) of each configuration is shown.

299 Semicrystalline P3HT has  $\sim 40\text{--}50\%$  crystalline domains by volume,<sup>9</sup> indicating that a sig-  
 300 nificant portion of P3HT can be described by plane-wave DFT, but long-range structural  
 301 correlations will be overestimated. To correct these errors, we would need to simulate large  
 302 unit cells to account for the proper decay in structural order. However, the computational  
 303 expense of DFT limits the size of the unit cell, meaning long range curvature present in  
 304 real polymer systems cannot be modeled. We are assuming that the curvature effects on  
 305 the spectrum are small compared to changes in local ( $\sim 10 \text{ \AA}$ ) structure, which is reasonable  
 306 provided that we only simulate modes that are sufficiently high in energy (i.e.  $> 700 \text{ cm}^{-1}$ ).  
 307 In this energy range, we expect that the local ordering of adjacent polymers will be the  
 308 dominant perturbation to the vibrational modes.

309 Semicrystalline P3HT contains several different crystalline polymorphs which have been  
 310 identified from X-ray measurements.<sup>15,17,18,51,64</sup> The most thermodynamically favorable poly-  
 311 morph is Phase I as determined by Poelking et al using a molecular dynamics approach. We

312 also consider the Phase I' polymorph proposed by Poelking et. al.,<sup>51</sup> but do not consider  
313 other polymorphs, such as Phase II because Phase II is irreversibly transformed into Phase I  
314 at room temperature and requires specific sample preparation that we did not apply.<sup>65,66</sup> The  
315 assumed structure of Phase I P3HT has the backbone plane parallel to the top/bottom face  
316 of the unit cell and side chains are tilted at an angle of approximately 47° with respect to the  
317 backbone plane.<sup>17,51,64,67</sup> Phase I' P3HT is energetically favored at low temperatures.<sup>51</sup> In  
318 the structure of Phase I', the backbone plane is tilted with respected to the top/bottom face  
319 of the unit cell, and the angle between the side chains and the backbone is approximately  
320 79° relative to the plane of the backbones.

321 RR-P3HT films are composed of amorphous and crystalline regions. We can generate  
322 an approximation for the crystalline region spectrum if we assume that the RRa-P3HT  
323 spectrum describes the amorphous regions of P3HT perfectly. Here we assume that the  
324 experimental spectrum is comprised of a combination of independent amorphous and crys-  
325 talline spectra, and we can assume that the amorphous phase comprises 55 vol% based on  
326 the work of Noriega et al.<sup>9</sup> The crystal INS spectrum is calculated as the total RR-P3HT  
327 spectrum minus the RRa-P3HT spectrum multiplied by its volume fraction. Here, we must  
328 assume that RRa-P3HT has the same INS spectrum as 100% amorphous RR-P3HT. To sup-  
329 port this assumption, we highlight the recent work of Shen et al, in which they demonstrated  
330 that the amorphous phase of RR-P3HT contains locally ordered sections of polymer likely  
331 due to interfacial regions between crystalline and amorphous domains, and that RRa-P3HT  
332 contains locally ordered regions as well due to assembly of local regio-regular segments.<sup>68</sup>  
333 Due to the structural similarities between RR and RRa P3HT, Shen et al subtracted GI-  
334 WAXS spectra of RRa-P3HT from RR-P3HT to find the crystalline volume fraction of  
335 RR-P3HT, which agrees with the results by Noriega et al.<sup>9</sup> Based on these results, we can  
336 directly compare the results of plane-wave DFT simulations to the crystalline INS spectrum  
337 to isolate structural motifs likely present in semicrystalline P3HT.

338 Simulations of an infinite crystal of Phase I P3HT (referred to as Config1) with identical

339 side chain configurations show a poor fit to INS data (Figure 4a). We therefore introduce  
340 heterogeneity in the crystalline phase of P3HT by simulating multiple side chain angles (Con-  
341 fig2) and dihedral disorder (Config3), while maintaining the same unit cell, which improved  
342 the fits (Figure 4b and 4c). Differences between measured and modeled data are primarily  
343 visible in the wag and bend vibrations between 650–1400  $\text{cm}^{-1}$ . The entire simulated spec-  
344 trum can be found in Supplementary Information Figure S3 and S4. The simulated spectrum  
345 for Config1 shows the largest error (from a least squares regression) with the measured INS  
346 data in the OOP bending region (red shaded region in Fig 4), indicating that this stacking  
347 motif likely represents only a small fraction of undoped P3HT. The best fit (largest fraction)  
348 for the backbone hydrogen OOP bending mode is displayed by Config2 (Figure 4b). This  
349 indicates that the OOP bending mode of the backbone hydrogen is coupled to the dihedral  
350 angle of the first carbon on the side chain, as seen in the INS spectrum. In Config2, each  
351 side chain alternates between a bent and flat dihedral angle with respect to the backbone  
352 plane. This side chain alternation leads to a 0.502 Å offset between P3HT backbones along  
353 the a-direction of the unit cell. For Config3 (Figure 4c), the dihedral angle between adja-  
354 cent backbone monomers was adjusted to induce a slight twist along the backbone, which  
355 captures the uncertainty in the monomer-monomer dihedral angle seen in prior molecular  
356 dynamics studies.<sup>51,69</sup> This twist creates a simulated spectrum that is between Config1 and  
357 Config2 in quality of fit. Config2 and Config3 both show a slight splitting/broadening of  
358 the peak at 750  $\text{cm}^{-1}$  (representative of torsional vibrations of side chain hydrogens), which  
359 is completely absent from the spectrum of Config1, but consistent with the measured INS  
360 data.

361 The choice of alternating flat and angled side chains within the unit cell of Config2 may  
362 seem arbitrary, but was chosen to favorably pack flat side chains into an experimentally  
363 verified unit cell. A configuration with completely flat side chains require a longer unit cell  
364 along the a axis. Extending one side chain (out of four) to the flat configuration results  
365 in unstable steric interactions between the flat side chain and the angled side chain in the

366 adjacent unit cell. Incorporating two flat side chains allows the polymers to nicely pack into  
367 an appropriately sized unit cell. Certainly one could imagine additional configurations that  
368 fit into the unit cell (by twisting dihedral number 2–5 along the side chain), but properly  
369 sampling of these configurations is computationally infeasible. In addition, we do not expect  
370 that configurational variations to the side chain will have a strong effect on the OOP and  
371 torsional vibrations. Nevertheless, it is important to stress that each of the configurations  
372 presented in Figure 4 are representative of different structural families/motifs.

373 The flat side chain motif in Config2 shows two distinct peaks in the torsional region of  
374 the spectrum (green shading). The presence of these two peaks does not agree with the  
375 measured INS data. The highest peak in the highlighted green region is due to a coupled  
376 carbon–carbon stretching mode that is amplified due to the completely flat side chain. Peak  
377 intensities in the region between 1050–1200  $\text{cm}^{-1}$  are better described by Config1. The fact  
378 that Config1 fits better in the torsional (green shading) region of the spectrum and Config2  
379 fits better in the OOP bending (red shading) shows that the experimental spectrum can  
380 not be explained using a single molecular configuration. Instead, this analysis indicates that  
381 there are multiple different configurations present in the sample.

382 We also tested whether the proposed Phase I' crystalline structure of P3HT was consis-  
383 tent with the measured INS spectrum (Figure 4d). The Phase I' structure has been measured  
384 and discussed as an energetically but not entropically favorable structure, meaning that it  
385 could be dominant with some thermal processing and would be stable at low temperatures  
386 like those used for the INS measurements.<sup>51</sup> Recent studies (both experimental and compu-  
387 tational<sup>15,18,51,69</sup>) predict the presence of P3HT configurations with tilted backbone planes  
388 with respect to the top/bottom faces of the unit cell, demonstrating that the configuration  
389 of P3HT inside the unit cell is still disputed. The simulated spectrum of Phase I' P3HT  
390 fits to the crystalline spectrum better (lower sum of squared error) than Config1, but less  
391 well than Config2 and Config3. However, the spectrum for Phase I' does have a prominent  
392 double peak near 700  $\text{cm}^{-1}$ , which is mostly absent from the configurations present in Fig-

393 ure 4, indicating that Phase I' may be present as a fraction of the total volume. Since INS  
 394 allows the superposition of spectral contributions of different configurations that sum to the  
 395 total observed spectrum, a linear superposition of the four crystalline, simulated spectra is  
 396 appropriate to determine phase fractions in crystalline domains.

## 397 Linear Combinations of Spectra

398 The semicrystalline nature of RR-P3HT requires a linear combination of spectral contri-  
 399 butions from different configurations and/or phases. Conveniently, the INS data can be  
 400 decomposed into contributions from different configurations whose weights represent specific  
 401 volume fractions of the particular configurations and/or phases because INS is a technique  
 402 that equally weights the contribution from every vibrational mode (no selection rules).

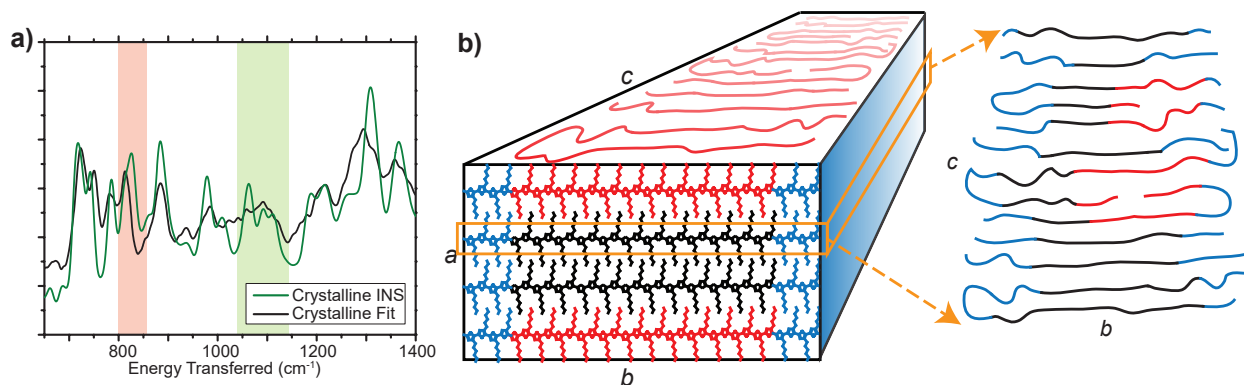


Figure 5: (a) Fitted linear combination of simulated spectra plotted against the experimental INS spectra (generated by subtracting the amorphous (RRa) fraction from the total undoped spectrum). (b) Cartoon of crystalline region of P3HT where black, red, and blue regions represent different configurations.

Table 1: Linear Combination Weights

System	Config1	Config2	Config3	Phase I'
Crystalline	0%	60%	2%	38%

403

*The weights of the optimal linear combination of effective crystalline spectrum (defined in text) to 4 crystalline configurations.*



404 Using the previously discussed crystalline INS spectrum, we combine the four crystalline  
405 DFT simulations to find the optimal linear combination of structural motifs. Figure 5a  
406 represents a least squares fit of the four crystalline simulations to the crystalline experi-  
407 mental spectrum. The contribution of each simulation is given by Table 1 indicating that  
408 the dominant structural motif is Config2 (60%), followed by Phase I' (38%) and Config3  
409 (2%). The error of the least-squares fit depends on the accuracy of the instrument and the  
410 computational method. As the experimental uncertainty within the energy transfer window  
411 discussed (600–1400  $\text{cm}^{-1}$ ) is quite low, the majority of the error lies with the assumptions  
412 and convergence criteria (very tight convergence criteria was used) of DFT. Thus, these pro-  
413 portions are the linear combination fit of the four crystalline structures is correct within the  
414 accuracy given by the PBE functional. However, as mentioned before, these structures are  
415 representative of different structural motifs. We expect more side chain heterogeneity in a  
416 real system, which would produce broader peaks in the non-highlighted regions of Figure 5a.  
417 Thus, the structural proportions presented in Table 1 only approximately correlate to the  
418 true proportions of structural families in the real material. Despite the approximate nature  
419 of our results, we can still draw conclusions from the fit between the four representative  
420 configurations.

421 There is no contribution from Config1, the most often cited crystalline motif from X-  
422 ray diffraction measurements. The high percentage contribution from Phase I' is due to  
423 the presence of a second peak around 750  $\text{cm}^{-1}$ , and absence of a peak at 775  $\text{cm}^{-1}$ . The  
424 other configurations do not display a peak at 750  $\text{cm}^{-1}$ , and overestimate the height of the  
425 peak at 775  $\text{cm}^{-1}$ . Thus the combination of Phase I' and Config2 synergistically describes  
426 the experimental data. The ratio of Config2 and Phase I' contributions to the fit of the  
427 INS data is fairly consistent in both the crystalline only fit and to the fit of the unaltered  
428 RR-P3HT spectrum (see Supplementary Information). This indicates in particular that  
429 there are a variety of side chain angles with respect to the backbone in crystalline P3HT  
430 domains and that the angle of the backbone with respect to the unit cell is not constant. We

431 again point out that P3HT forms crystalline ribbons and the crystalline chains in the center  
432 of the ribbon may have significantly different geometry due to the presence of an ordered  
433 crystal field compared to P3HT that forms the top, bottom, internal domain boundaries,  
434 and ribbon edges (Figure 5b). In addition, real morphologies likely contain continuous  
435 transitions between crystalline and amorphous regions, which should increase the structural  
436 heterogeneity, overall peak width, and background of our simulated spectrum.

437 The question of whether or not the backbone is tilted with respect to the top/bottom face  
438 of the unit cell has long been disputed.<sup>18,51,67</sup> Our results demonstrate that both stacking  
439 motifs are present in the crystalline region, and can be elucidated with INS, emphasizing  
440 the sensitivity of this measurement technique. Poelking et al used molecular dynamics  
441 to show that the crystalline phase had thiophene rings that were flat with respect to the  
442 top/bottom of the unit cell,<sup>51</sup> whereas a similar study using a different force field showed  
443 that similarly sized polymer chains (20-mers) generate a unit cell with a tilted backbone.<sup>69</sup>  
444 This discrepancy highlights the theoretical difficulty in identifying the local atomic-level  
445 structure. In the latter study, the backbones changed from a tilted configuration to a flat  
446 configuration when the chain length was increased to 40-mers, suggesting that the crystalline  
447 phase may contain regions with both tilted and flat backbones.

448 Our results help rationalize the differences observed in prior molecular dynamics<sup>51,69</sup> and  
449 experimental studies<sup>15,18,64,67</sup> considering the picture of the crystalline domain in Figure 5a.  
450 The INS data suggests that the crystalline domain is comprised of at least three distinct  
451 structural motifs, which can be explained by three different local field regimes within the  
452 crystalline domain that cause changes in local structure of the polymer. The inner region of  
453 the crystalline domain is different from the regions at the top/bottom and left/right of the  
454 domain (Figure 5a), which may cause distortions in the side chain tilt and/or dihedral angle  
455 between monomers. We can postulate that the blue, red, and black segments of Figure 5a  
456 correspond to dihedral distortions (Config3), side chain tilt variation (Config2), and a highly  
457 crystalline morphology (Phase I'), respectively. While this postulate intuitively makes sense,

458 we cannot assign the colored segments of Figure 5a to specific configurations presented in  
 459 this study without a larger length scale model. However, we can conclude that the crystalline  
 460 domain of P3HT is structurally heterogeneous at the atomic level with three local structural  
 461 motifs dominating the distribution of configurations.

## 462 Doped Simulations

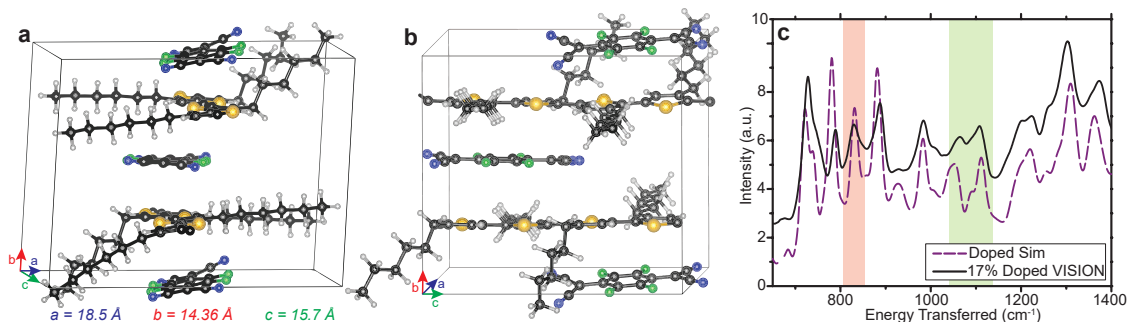


Figure 6: (a) Front view, and side view (b) of structure of doped P3HT in the crystalline phase used for simulation. (c) The simulated spectrum is plotted against the 17 mol% doped INS spectrum.

463 Recent publications about P3HT doped with F4TCNQ<sup>31,33</sup> clearly show that at low  
 464 doping ratios ( $\leq 3 \text{ mol\%}$ ), F4TCNQ is mostly found in amorphous domains. At higher  
 465 doping ratios, F4TCNQ inserts between P3HT backbone planes. Although at 4% doping  
 466 some F4TCNQ likely does intercalate, this does not have a significant effect on the vibrational  
 467 spectrum, as seen in Figure 3. The maximum doping ratio is approximately 17 mol% and  
 468 represents full intercalation of F4TCNQ into crystalline domains in addition to doping of  
 469 the amorphous sections of the polymer.<sup>31,70</sup> The 17 mol% doped sample represents a much  
 470 more uniform sample than lower doping ratios because doping likely causes planarization of  
 471 the P3HT backbone, which removes local structural heterogeneity.

472 Figure 6 shows a fit to the INS spectrum of P3HT doped with 17 mol% F4TCNQ that  
 473 represents a dominant configuration. Since we do not know the volume of amorphous do-  
 474 mains in the doped polymer, the peak heights are difficult to quantitatively explain, so we

475 will focus on the peak energies. The simulated spectrum shows good agreement with the  
476 doped experimental spectrum for all vibrational peak frequencies. This indicates that the  
477 structure presented in Figure 6a is dominant in the doped ensemble. Both highlighted re-  
478 gions are well explained by the simulation showing that the stacking motifs present in the  
479 structure are widespread throughout the system. F4TCNQ is intercalated between P3HT  
480 chains, but not adjacent to the same monomers; one is displaced along the  $c$  axis relative  
481 to the other. The lattice constants were taken from a combination of X-ray diffraction<sup>31</sup>  
482 and molecular dynamics studies.<sup>51</sup> The plane of the backbone (created by the flat face of  
483 the 5-membered thiophene ring) is parallel to the plane of the closest dopant. The two  
484 chains are not vertically aligned with one another, indicating that F4TCNQ prefers to in-  
485 teract with a single P3HT chain. The molecular planes of the dopants are not parallel,  
486 causing both P3HT chains to distort and twist along the backbone indicating that charging  
487 the backbone/F4TCNQ combination may cause some puckering/distortions in the backbone  
488 to minimize the electrostatic potential. The average charge transfer between P3HT and  
489 F4TCNQ in this configuration is 0.92 electrons from the P3HT to F4TCNQ using Mulliken  
490 analysis.<sup>71,72</sup> This near integer charge transfer from P3HT to F4TCNQ is consistent with  
491 spectroscopic analysis.<sup>73</sup>

492 The INS data shows strong interactions in the OOP bending and torsional modes, which  
493 involves hydrogens on the backbone and the first and second carbon of the side chain. In our  
494 simulated structure, we see that some of the nitrogen/fluorine groups are nearest to either  
495 the backbone hydrogen or the hydrogens attached to the first carbon of the side chain.  
496 The electronegative atoms are close enough to interact with the hydrogens responsible for  
497 the OOP bending and torsional bending modes but not close enough to call these hydrogen  
498 bonds. These dipolar interactions, in addition to the electrostatic attraction of the negatively  
499 charged dopant to the positively charged polymer, have a strong effect on the determination  
500 of the dopant location with respect to the polymer. The fact that the simulated spectrum  
501 matches at every vibrational peak in these regions is a strong indication that the proposed

502 structure is valid. In addition, each peak in the displayed energy range is well explained by  
 503 a single configuration. Since we required two major configurations to explain the undoped  
 504 spectrum, we can infer that the distribution of local configurations shrinks when heavily  
 505 doped. In other words, there appears to be more local order when P3HT is heavily doped.

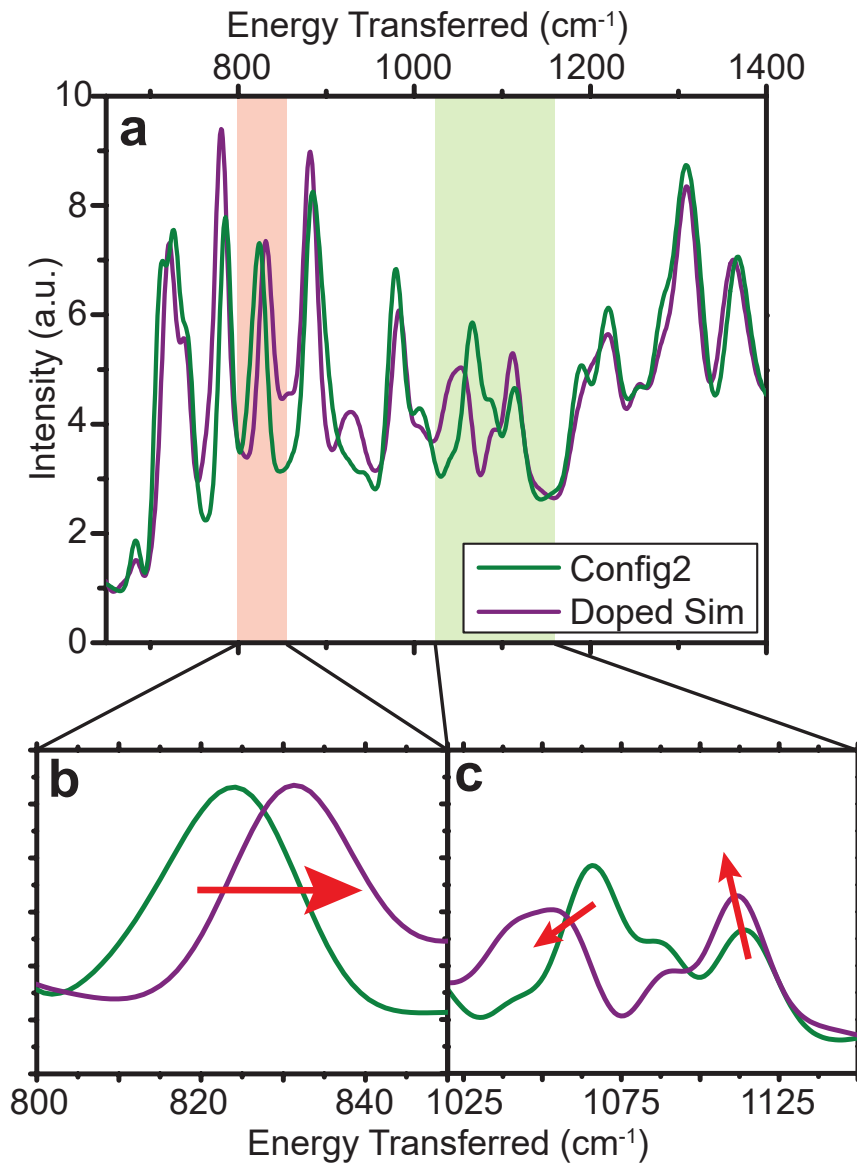


Figure 7: *The simulated spectra of Config2 and the doped configuration are compared. The two shaded regions are expanded to demonstrate the specific effect of doping.*

506 The simulated spectra of undoped Config2 and the simulated doped configuration are  
 507 compared to demonstrate the internal consistency of the modeling approach. Config2 was

508 chosen because it is the undoped configuration most similar to the doped configuration,  
509 and the most prevalent configuration in the undoped crystal. The OOP bending peak at  
510  $\sim 820 \text{ cm}^{-1}$  blue shifts upon doping as seen in the experimental spectra (Figure 7b). This  
511 blue shift is likely due to a combination of increased electrostatic interactions between dopant  
512 and backbone hydrogen and/or charging of the backbone (through doping) leading to a stiffer  
513 mode. F4TCNQ is  $3.59 \text{ \AA}$  away from the P3HT backbone, whereas the P3HT chains are  
514  $3.83 \text{ \AA}$  away from each other when undoped, which reduces the average interaction distance  
515 of the backbone hydrogen. As a result, the backbone hydrogen experiences greater forces  
516 when moving perpendicular to the plane of the backbone.

517 The doped simulation displays a greater splitting between peaks in the torsional region  
518 of the spectrum (Figure 7). The peak at  $\sim 1120 \text{ cm}^{-1}$  increases in amplitude, while the peak  
519 at  $\sim 1065 \text{ cm}^{-1}$  shifts to lower energy upon doping. The former is due to a red shift (lower  
520 energy) in an in-phase, coupled rowing vibration of hydrogens along a flat side chain (see  
521 Supplemental Information Figure S9d). When P3HT is doped, this motion has a similar  
522 energy to the torsional vibration as shown in Figure 2c, causing the increase in peak height.  
523 The shift in the peak at  $1065 \text{ cm}^{-1}$  to lower energy is the result of a relaxation of coupled  
524 carbon-carbon stretching modes along the length of the side chains. The motion of the side  
525 chain carbons and hydrogens close to the backbone are less stiff when doped. The positive  
526 charge on the backbone, coupled with the electron donating nature of the hexyl side chain,  
527 make the side chain bonds less stiff. All of these simulated vibrational changes due to doping  
528 are consistent with the trends observed in the experimental spectra.

## 529 **Conclusions**

530 The INS instrument VISION records a spectrum that is similar to the pure density of vibra-  
531 tional states, which means that the data is a superposition of the amplitude and frequency  
532 for each vibrational mode for each configuration of P3HT. By investigating the major trends

533 in amplitude and frequency for several indicator modes, we are able to ascertain configura-  
534 tions of P3HT that are the most prevalent structural themes. Specifically, we demonstrate  
535 that the crystalline configurational distribution contains at least three dominant structural  
536 motifs represented by individual configurations: one with flat side chains and flat backbones  
537 (Config2  $\sim 60\%$  present), one with tilted side chains and a slightly twisted backbones along  
538 the chain (Config3  $\sim 2\%$  present), and one with tilted side chains and a tilted backbone  
539 (Phase I'  $\sim 38\%$  present). Backbone "flatness" is defined relative to the top and bottom  
540 face of the unit cell. The observed mixture of backbone/side chain tilt angles highlights the  
541 large amount of structural heterogeneity in undoped P3HT that makes the determination  
542 of atomic scale structure using traditional experiments such as XRD more difficult. The  
543 simultaneous occurrence of tilted and flat backbone unit cells reconciles differences in prior  
544 experimental studies reporting crystal structures,<sup>15,18,64,67</sup> and makes sense in the context of  
545 the formation of nanocrystallites that are known to form in thin films of solution-processed  
546 P3HT.<sup>9,33</sup>

547 P3HT doped with F4TCNQ is dominated by structures with F4TCNQ intercalated  
548 between P3HT chains. Here we report a unit cell structure for P3HT with intercalated  
549 F4TCNQ that is consistent with INS data, which provides atomic-scale structural informa-  
550 tion, providing more detail than previous reports which only reported unit cell dimensions.  
551 The intercalation of F4TCNQ increases the inter-atomic forces acting on the backbone hy-  
552 drogen, which causes a blue shift in the OOP bending mode in the INS spectrum. The  
553 splitting in the torsional region of the INS spectrum is similar to the splitting displayed by  
554 Config2 (Figure 4b), indicating that the presence of F4TCNQ causes the side chains to flat-  
555 ten. The intercalation of F4TCNQ molecules into the crystalline region of P3HT causes the  
556 polymer chain to become flatter to best accommodate the packing of a flat dopant molecule.  
557 Since each peak is captured by a single configuration, we expect that high doping ratios  
558 produce a smaller distribution of local configurations and a more homogeneous film than  
559 undoped P3HT, which could impact the electrical properties.

560 We must emphasize that high-resolution INS contains the information to infer the detailed  
561 structural characteristics of disordered systems. However, our ability to model semicrys-  
562 talline and doped semiconductors polymers is limited by methodology and computational  
563 expense. Thus, improved modeling techniques are needed to fully model longer range (lower  
564 energy) vibrations over a much larger volume. In particular, modeling techniques that incor-  
565 porate longer length scales but maintain electronic accuracy are lacking. We anticipate that  
566 INS spectroscopy will become a valuable tool in the study of polycrystalline and disordered  
567 materials; the structures of countless other disordered materials can be optimized using INS  
568 spectroscopy at the VISION instrument.

## 569 **Supporting Information**

570 Video files of all relevant vibrational modes are present in GIF format.

## 571 **Acknowledgements**

572 This research project was supported by the U.S. Department of Energy, Office of Basic  
573 Energy Sciences, Division of Materials Sciences and Engineering, under Award No. DE-  
574 SC0010419. This research was made possible by the use of the VISION beamline at ORNLs  
575 Spallation Neutron Source, which is supported by the Scientific User Facilities Division, Office  
576 of Basic Energy Sciences (BES), U.S. Department of Energy (DOE), under Contract No.  
577 DE-AC0500OR22725 with UT Battelle, LLC. Computational work was conducted on the  
578 Oak Ridge Leadership Computing Facility (OLCF) supported by the DOE. The computing  
579 resources were made available through the VirtuES (Virtual Experiments in Spectroscopy)  
580 project, funded by Laboratory Directed Research and Development program. No. 7739).



## 581 Additional Information

582 **Competing Financial Interests:** The Authors declare no competing financial interests.

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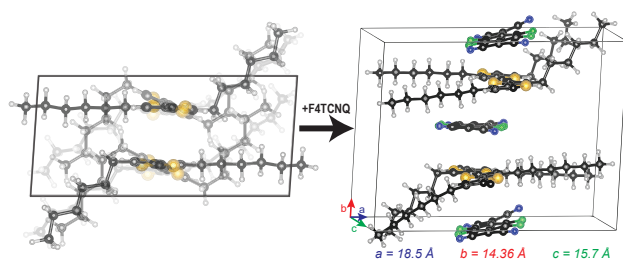
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