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► **To cite this version:**

Yu Ogawa. Electron microdiffraction reveals the nanoscale twist geometry of cellulose nanocrystals. *Nanoscale*, 2019, 10.1039/C9NR06044H . hal-02360161

HAL Id: hal-02360161

<https://hal.science/hal-02360161>

Submitted on 14 Nov 2020

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1 **Electron microdiffraction reveals the nanoscale twist geometry of cellulose nanocrystals**

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6

7 **Abstract**

8 Nanocellulose consisting of crystalline cellulose nanoparticles has a high potential to
9 serve as a building block for bio-based functional materials. The intrinsic chirality of
10 cellulose crystals provides them with high added values such as optical properties and chiral
11 induction ability. At the nanoscale, this chirality is connected to the right-handed longitudinal
12 twisting of these fibrous crystallites. However, this nanoscale fibrillar twist has been a matter
13 of debate due to contradictory data between ultrastructural observations and molecular
14 simulations and so far, the exact twist geometry has not been elucidated. Here, an electron
15 microdiffraction (μ ED) analysis under cryogenic condition reveals the continuous twisting of
16 cellulose nanocrystals (CNCs) in aqueous suspension. This intrinsic regular twist is
17 drastically modified to discontinuous sharp twists when the CNCs are dried on flat surface.
18 The present μ ED-based analysis at the single nanoparticle level allows establishing
19 quantitative structure-properties relationship of various solid and colloidal nanocellulose
20 systems.

21

22 Keywords: Cellulose. Electron diffraction. Cryogenic transmission electron microscopy.

23 Twist. Chirality. Nanoparticle.

24

25 **Introduction**

26 Cellulose is attracting renewed scientific and industrial interests thanks to its high
27 potential as nanosized building blocks for functional bio-based materials.¹⁻⁴ In this context,
28 crystalline cellulose fibrils, commonly called nanocellulose, obtained from native cellulosic
29 tissues, are categorized as short cellulose nanocrystals (CNCs) or long cellulose nanofibers
30 (CNF). Together with their outstanding mechanical properties,⁵⁻⁹ one of the most fascinating
31 aspects of nanocelluloses is their intrinsic chirality, which gives rise to various useful
32 properties such as chiral optical activity¹⁰ and chiral induction ability.¹¹ The chirality of
33 nanocelluloses is present at various length scales, from the optically active glucosyl residues,
34 *i.e.* the molecular building block of cellulose, to that of their suspensions, which are
35 susceptible to spontaneously self-organize into chiral-nematic structures that can be
36 preserved in a solid state after drying.^{10, 12, 13} At the nanometric scale, the chirality of
37 cellulose can be observed as unidirectional twists along the fibrillar axes of nanocelluloses.¹⁴⁻
38 ¹⁶ Such twists are of great importance since they are considered to govern the chiral-related
39 properties of nanocellulose materials.^{1, 11, 17} Despite this, the ultrastructural details of the
40 fibrillar twists of nanocelluloses have not been elucidated and have been in under debate due
41 to contradicting experimental and computational observations as described below.

42 The twist of nanocelluloses has been observed using conventional imaging techniques
43 such as atomic force microscopy (AFM), scanning and transmission electron microscopy
44 (SEM and TEM, Fig. 1a-d) together with cryogenic TEM and electron tomography.^{14-16, 18}
45 Hanley and co-workers¹⁴ have reported ultrastructural details of the twisted morphology of
46 large isolated nanofibrils extracted from the cell wall of the alga *Micrasterias denticule* dried
47 on a flat substrate. In these, 180° sharp right-handed twist regions occurred regularly spaced
48 with flat and untwisted regions.¹⁴ Such sharp discontinuous right-handed twists, as illustrated
49 in Fig. 1e, were also reported for nanocelluloses extracted from other sources including wood

50 and bacterial celluloses under similar observation conditions.¹⁵ These experimental
51 observations have been confronted with atomistic molecular dynamics (MD) simulations that
52 also proposed right-handed twist occurrence for finite-size cellulose crystals.¹⁹⁻²³ However,
53 these MD simulations predict continuous smooth twists along the fiber axes (Fig. 1f) unlike
54 the alternate occurrence of the sharp twists and flat untwisted regions observed
55 experimentally. It remains to be determined whether the observed discontinuous twists in
56 dried nanocellulose correspond to artifacts resulting from sample preparation or to intrinsic
57 features of the samples.

58 To address the question of fundamental understanding of the twist morphology, one
59 requires devising a methodology that allows quantitative description of local geometry of the
60 cellulose crystal at the single nano-object level. In this work, a combination of cryogenic
61 transmission electron microscopy (cryoEM) and electron microdiffraction (μ ED) was used to
62 follow the exact geometry of the twist of nanocelluloses as they occur in aqueous
63 environment. Typically, a μ ED analysis can provide crystallographic information on
64 individual nano-sized crystalline domains, thus leading to their local crystallographic
65 orientation from two-dimensional diffraction patterns. By recording sequential μ ED patterns
66 along the axis of cellulose nanocrystals embedded into vitreous ice, the structural details of
67 the twist of CNCs, such as twist angle and pitch could be described. CNCs under dry and
68 frozen conditions were compared, which allowed elucidating the effect of their drying on a
69 supporting substrate.

70

71 **Results**

72 **Sample**

73 In this study, tunicate CNCs (tCNCs) extracted from the mantle of *Halocynthia*
74 *roretzi* were used for their high crystallinity and large crystalline domains, hence their

75 relatively high resistance toward electron radiation dose. The cross-section of a given tCNC
76 consists in an elongated octagon where cellulose molecular sheets organized by hydrophobic
77 interactions are stacked along its wider side (Supplementary Information, Fig. S1c).²⁴ The
78 basic structural characterization of the tCNCs used in this study is summarized in Fig. S2 in
79 Supplementary Information file. Briefly, the tCNCs correspond to highly crystalline cellulose
80 I β crystals as described in literature. The average width such nanocrystals is estimated as 14.4
81 \pm 3.3 nm based on the cryoTEM observations. The acid hydrolysis with sulfuric acid induced
82 sulfate half-ester groups on the surface of the tCNCs, giving a rise to negative surface charge
83 of tCNCs. Such sulfate ester groups are detectable in FT-IR spectrum at 807 cm⁻¹ (Fig. S2b).
84 The zeta potential of tCNC was estimated as -35.1 mV.

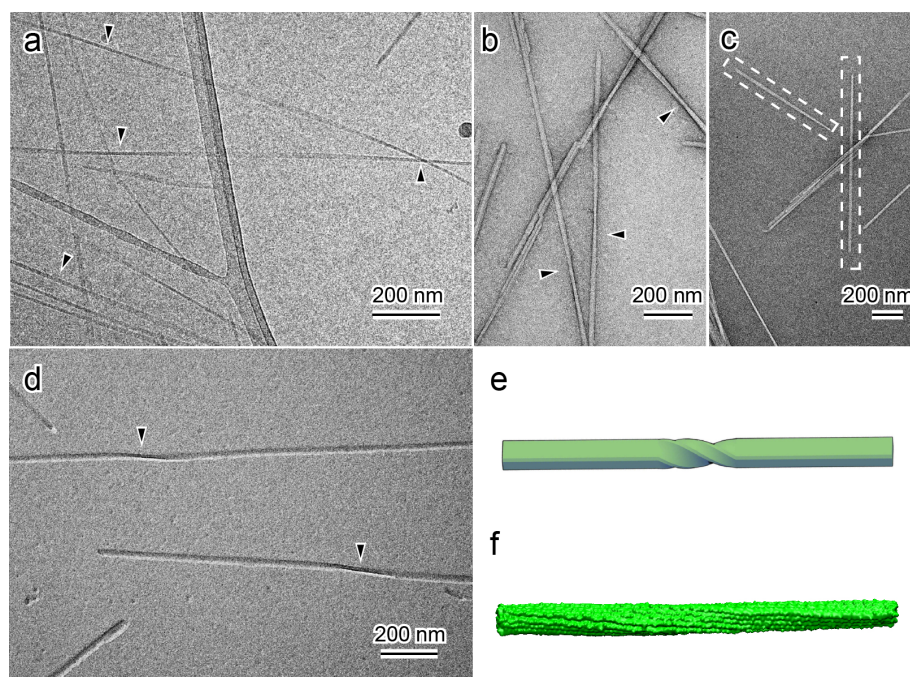
85

86 **Apparent fibrillar twist of cellulose nanocrystals visualized by TEM**

87 Figure 1a shows a cryoEM image of tCNCs in aqueous suspension where the twists are
88 revealed as a gradual change in their contrast and their width along their length. They were
89 also visible in tCNCs dried on a thin carbon film after negative staining (Fig. 1b) or metal
90 shadowing (Fig. 1d). The twists revealed in the images of shadowed tCNC are also right-
91 handed and discontinuous as in the literature. The twist was virtually invisible in some short
92 CNCs (Fig. 1c), being consistent with the low observation frequency of apparent twist, less
93 than 10 % in the report on twist morphology of wood and bacterial CNCs by Usov and co-
94 workers.¹⁵

95 Although the twist is visible in these micrographs, the elucidation of the exact geometry of
96 these twists is extremely challenging from the sole analysis of these images. This is partly
97 because the cellulose crystals have an intrinsic size distribution in lateral dimensions,²⁵⁻²⁷
98 which precludes a direct correlation between their dimensions and their local internal
99 structure.

100



101

102 **Figure 1. Twisted morphology of cellulose nanocrystals.**

103 (a) CryoEM image of tCNCs; (b, c) conventional TEM image of negatively stained tCNCs;

104 (d) images of tCNCs shadowed with W/Ta alloy. In Fig. 1a-1d, the arrowheads point toward

105 the apparent twisted regions, and dashed squares indicate apparently untwisted tCNCs. (e)

106 schematized discontinuous twist morphology observed on CNCs in dry condition. (f) MD-

107 simulated cellulose nanocrystal showing a right-handed twist.

108

109 **Intrinsic regular twist of tCNCs in aqueous suspension**

110 Fig. 2 summarizes the cryoEM/ μ ED analysis of a single tCNC which was embedded in

111 vitreous ice, in such a way that its nanoscale morphology was maintained as in the aqueous

112 suspension state. These sets of two-dimensional μ ED patterns were recorded with an electron

113 probe of a diameter of approximately 70 nm. The reflections in these patterns were indexed

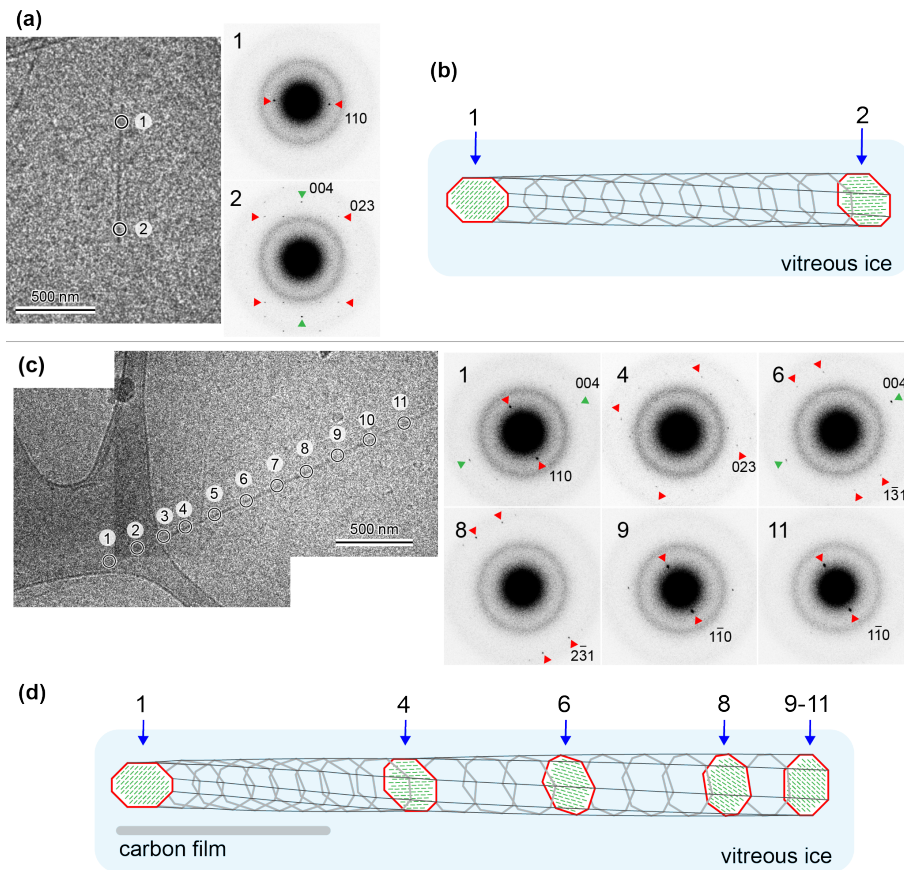
114 according to the unit cell of cellulose I β ,^{28,29} (Supplementary Information Figs. S1a and S1b),

115 allowing the determination of the exact crystallographic orientation of the tCNC at each

116 acquisition position with respect to the incident electron beam, as schematized in Figs S3 and
117 S4.

118 As can be seen in Figure 2a, the first tCNC is relatively short with a length of
119 approximately 800 nm and isolated from other tCNCs and the supporting lacey carbon
120 network. Two diffraction patterns recorded at both ends show different projections, one
121 containing the $1\ 1\ 0^*$ vector perpendicular to the crystal direction and the other containing b^*
122 and c^* indicating that this crystal is rotated along its longitudinal axis (Supporting
123 Information, Figure S3). Considering the right-handedness of the twist, comparison of the
124 two diffraction pattern is equivalent to the recording of another diffraction pattern with 45°
125 clock-wise rotation at point 1. Thus, as previously demonstrated by diffraction experiment of
126 a series of rotation,³⁰ the polarity of this tCNC was determined as having the c-axis pointing
127 upward in this image. Therefore, the reducing end of this crystal is located at its top-end since
128 the cellulose molecules are packed into a parallel-up manner in their unit cell.^{29, 31}

129



130

131 **Figure 2. Longitudinal twist of tCNCs embedded in vitreous ice studied using μ ED.**

132 (a, c) Diffraction contrast images of tCNCs under cryo-frozen condition, together with μ ED

133 patterns recorded from acquisition areas indicated by the circles. (b, d) Schematic

134 illustrations of the cross-sectional orientation of tCNC along their longitudinal axis at each

135 acquisition point.

136

137 The second tCNC is longer and one of its ends is in contact with the carbon of the

138 supporting lacey film (Figure 2c). For this tCNC, a series of μ ED patterns were taken along

139 the tCNC longitudinal axis over a 2.2- μ m distance at intervals of approximately 200 nm.

140 Selected μ ED patterns are shown in Fig 2c 1-11 and the full diffraction data set is given in

141 Figure S3 together with schematic presentations of corresponding projections of reciprocal

142 lattice. The diffraction patterns at acquisition points 1-9 correspond to different projections of

143 the reciprocal lattice, indicating a continuous rotation of the crystal (Figs. 2d) with a rotation

144 angle of 90° from position 1 to 9 separated by $1.6 \mu\text{m}$. On the other hand, the patterns from
145 positions 9-11 are identical, showing a projection containing the $1 -1 0^*$ vector. As in the first
146 example, the polarity of this tCNC can be determined considering the right handedness of
147 twist, indicating that the reducing end is located toward the left-hand end of the tCNC in the
148 figure.

149 As demonstrated by these μED analyses in Fig 2 together with other examples in Fig. S5,
150 the tCNCs clearly possess the slow continuous twist along their longitudinal axes in the
151 aqueous suspension state. Their twist rates along the longitudinal axis of the crystal are not
152 constant not only between these two crystals but also within a single crystal itself as in the
153 second tCNC. The twist rate of the first crystal between points 1 and 2 is of $5.8^\circ/100 \text{ nm}$
154 (Figure 2b), while the one of the second one is about $9^\circ/100 \text{ nm}$ between positions 1 and 4,
155 and $4^\circ/100 \text{ nm}$ between positions 4 and 9, respectively (Figure 2d). The nanocrystals in Fig.
156 S5 gave twist pitches of about $10^\circ/100 \text{ nm}$ (Fig. S5a) and $13^\circ/100 \text{ nm}$ (Fig. S5b), respectively.
157 The difference between the twist geometries of these crystals is reasonable considering that
158 the tCNCs have a broad distribution in width and cross-sectional shape.²⁵⁻²⁷ MD simulations
159 predict that the difference in cross-sectional shape of a cellulose crystal greatly affects its
160 twist rate, where the crystal with a larger diameter should be less twisted.³² This effect is due
161 to the contradictory contribution of the twisting tendency of individual chains and the elastic
162 constraint against axial deformation that increases with distance to the center of the
163 nanocrystal in the lateral plane. The change of twist rate in the second tCNC is presumably
164 due to the fact that this tCNC is apparently in contact with the carbon film between points 1
165 and 3, which may cause a distortion of the twist geometry as a consequence of the interaction
166 between cellulose and the carbon film.

167

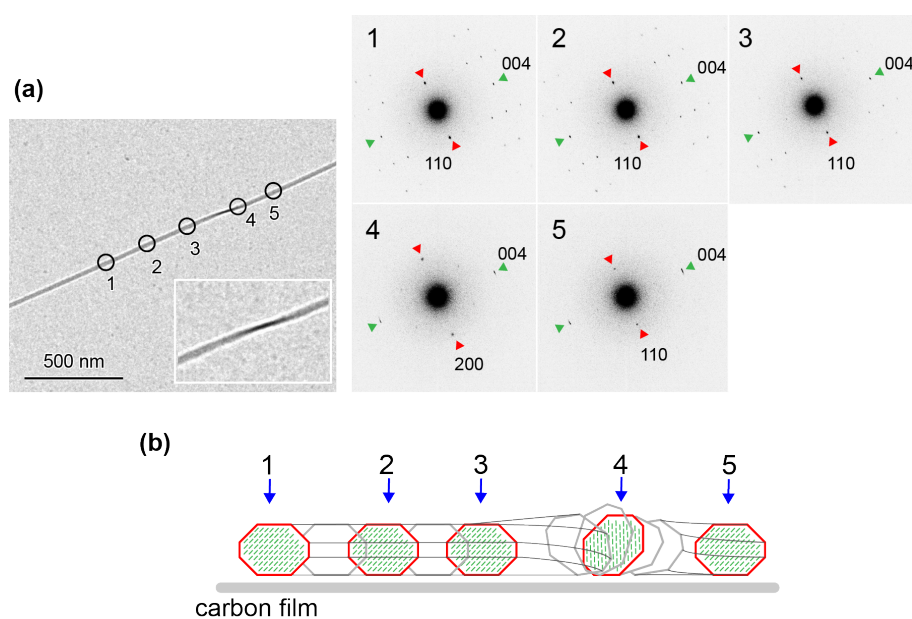
168 **Alteration of twist geometry of tCNCs upon drying on flat surface**

169 The μ ED experiments under cryogenic condition reveals the presence of intrinsic regular
170 twist of tCNCs in the aqueous suspension state as predicted in MD simulations (Fig. 1f). It is,
171 however, incompatible with the apparent twist geometry observed in dry conditions (Fig. 1d
172 and 1e). The similar μ ED analysis was thus performed on a tCNC dried on a flat substrate,
173 under a condition equivalent to those in the literature.¹⁴⁻¹⁶ Figure 3a summarizes a sequential
174 μ ED series obtained along a tCNC dried on an amorphous carbon film. The indexation of
175 diffraction spots in Fig. 3a is given in Figure S6. The part of the tCNC subjected to this
176 diffraction analysis shows a twist region separated by two flat segments. The flat segments
177 give identical diffraction patterns, a projection of the reciprocal lattice containing c^* and $1\ 1\ 0^*$
178 vectors (Fig. 3a, positions 1-3, and 5), so that the tCNC lays flat with its wider side flat on
179 the carbon supporting film (Fig. 3b). The diffraction pattern obtained from the apparent twist
180 region is a projection of planes containing c^* and a^* (Fig. 3a, position 4), indicating that the
181 longer side of the octagonal cross-section was oriented diagonally with respect to the carbon
182 film plane. This is consistent with the fact that the twisted region appeared to be thinner,
183 compared to the flat segment regions. Thus, the tCNC is twisted by 180° along its axis over
184 the apparent twist position (Fig. 3b), the twist localized over a distance of less than 400 nm
185 along the axis of the tCNC, between positions 3 and 5.

186 In this study, 28 other tCNCs dried on carbon films were analyzed using the same
187 sequential μ ED measurements. Two examples of the analysis are given in Figs. S7 and S8 in
188 the Supplementary Information file. All the nanocrystals possessed flat segments extending
189 over more than 500 nm. Among 42 observed flat segments, the majority (74%) of them gave
190 ED patterns of the projection of $1\ 1\ 0^*-c^*$ plane, while the other 26% were of the $1\ 2\ 0^*-c^*$
191 projection. 18 apparent sharp twist regions were examined. They provided the nanocrystals
192 with either a 180° ($1\ 1\ 0^*-c^*$ to $1\ 1\ 0^*-c^*$, Figs. 3 and S7) or 194° flip ($1\ 1\ 0^*-c^*$ to $1\ 2\ 0^*-c^*$,
193 Fig. S8). No smaller twist angle was observed in this study.

194 The difference of the twist geometries of the tCNCs under dry and cryo-frozen conditions
 195 is striking, as the tCNCs show a sharp localized twist when dried on a carbon film as opposed
 196 to a continuous extended twist in vitreous ice. The sharp twists observed under dry condition,
 197 not only in this study, but also in many images found in literature¹⁴⁻¹⁶ are thus a consequence
 198 of the drying process of CNCs on flat substrates, as suggested earlier by Hanley and co-
 199 workers.¹⁴ Considering that the surface tension between water and solid are significant at
 200 nanoscale, it is likely that the intrinsic twisting habit of cellulose nanocrystals is deformed by
 201 the large capillary force resulting from drying, with the result of the local de-twisting
 202 observed over the flat areas.

203



204

205 **Figure 3. Longitudinal twist of a dry tCNC followed by μED.**

206 (a) Diffraction contrast image of a tCNC dried on a carbon film, together with μED patterns
 207 recorded from the acquisition points indicated by circles. The diameter of the circle is equal
 208 to that of the electron probe used to record the patterns. Inset: enlarged image of the twisted
 209 region. (b) Schematic illustration of the cross-sectional orientation of the tCNC along
 210 longitudinal axis at each acquisition point.

211

212 **μ ED analysis of parallel assembly of tCNCs.**

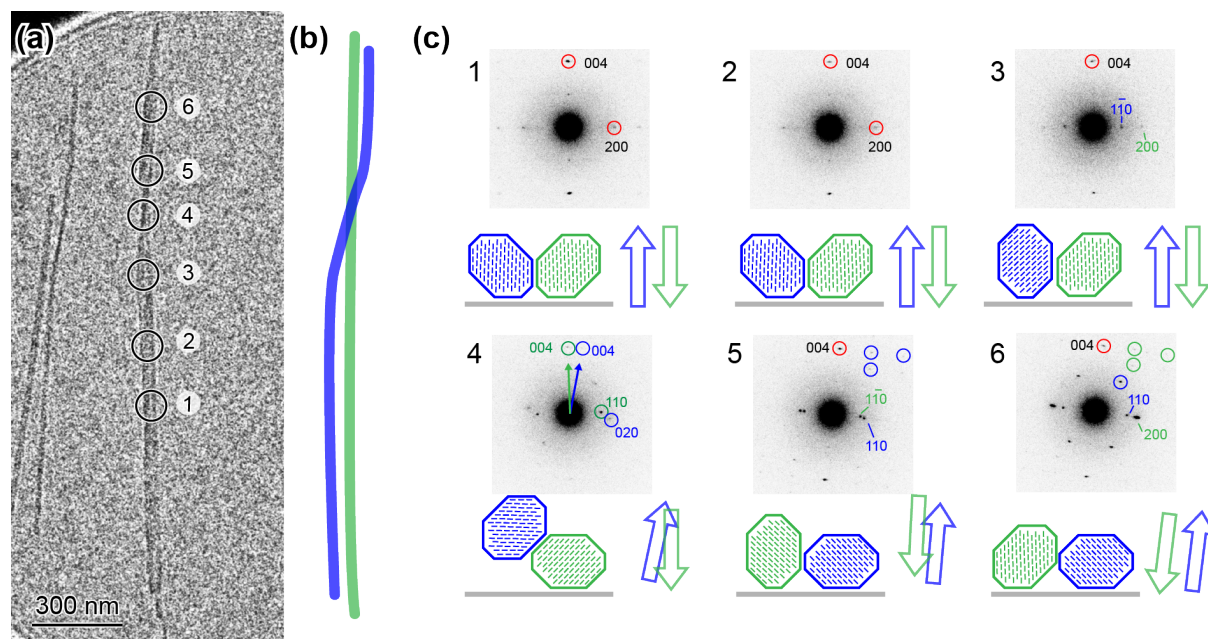
213 The use of μ ED analysis is not only meaningful for single tCNCs, but is also extendable to
214 fibrillar bundles of nanocelluloses, commonly observed with various imaging techniques.¹⁵
215 Ultrastructural details of such assemblies are important as bundles of CNCs behave as single
216 particles in suspensions and thus have an impact on the suspension properties. In this line, it
217 is important to reveal (i) whether in a given bundle of parallel CNCs, the crystals have all the
218 same polarity with respect to their reducing direction, (ii) whether the twisted regions may
219 lead to the intertwinement of several CNCs, and (iii) whether the twists of individual
220 crystallites are in sync.

221 Fig. 4a shows an assembly composed of two adjacent parallel tCNCs dried on a carbon
222 film. As schematized in Fig. 4b, there is an apparent intertwinement of both tCNCs at their
223 mid-lengths. μ ED patterns taken from a series of acquisition points along the tCNCs are
224 shown in Fig 4c. The diffraction patterns obtained from the bottom half (Fig.4c, 1 and 2)
225 correspond only to one projection containing a^* and c^* vectors, indicating that two crystals
226 have the same cross-sectional orientation with respect to the incident electron beam. On the
227 other hand, the patterns taken from positions 3 to 6 contain two different projections. The
228 evolution of the cross-sectional orientation along the axis indicates that these two crystals
229 have different rotation directions. Assuming that both crystals maintain their right
230 handedness, these two crystals are therefore associated in an antiparallel manner with their
231 crystal polarities being opposite to one another. In Figure 4a and its model in 4b, the “blue”
232 nanocrystal has its reducing end toward the top of the pictures whereas in the “green” one, it
233 is toward the bottom.

234 Based on this series of μ ED patterns, one can also deduce which of the surfaces of both
235 crystals are in contact as illustrated in Fig. 4c. It is noticeable that the two crystals do not
236 twist at the same location, but successively one after the other. Also, the contact planes

237 between the two crystals are modified during the twisting, both crystals being jointed by their
 238 hydrophobic (200) planes in the locations 1 and 2, while at the end of the twist, in position 6
 239 it is the (110) planes of one crystal, which is adjacent to the (200) plane of the other.

240



241

242 **Figure 4. Joint assembly of two intertwined tCNCs.** (a) Diffraction contrast image of the
 243 joint assembly of two tCNCs. (b) Schematic illustration of the assembly. (c) Structural details
 244 of the assembly at each acquisition point indicated in (a). Each subpanel contains a μ ED
 245 pattern, a cross-sectional organization of both tCNCs, together with hollowed arrows
 246 indicating the polarity of the tCNCs pointing toward their reducing ends. In c4, the solid
 247 arrows indicate that the directions of fiber *c* axes are different for the two crystals.

248

249 Discussion

250 The comparison between the twist geometry of tCNCs under dry and cryo-frozen
 251 conditions highlights the importance of using cryoEM techniques to preserve the intact
 252 morphology of nanocelluloses in aqueous suspension. The effect of drying on the nanoscale
 253 morphology of tCNCs is significant, since it converts their initial regular longitudinal twist
 254 into long flat untwisted segments alternating with sharp right-handed flips with a twist angle

255 of about 180° . Although the twist morphology observed in the dried condition is ultimately
256 derived from that in the suspension state, it remains to be investigated if and how these two
257 morphologies are quantitatively correlated. Based on the amyloid fibril systems, it has been
258 suggested that the adsorption of twisting nanofibers on a flat substrate would not alter the
259 twist pitch when a full periodicity is observable even when the exact twist geometry was
260 altered by the surface interactions with supporting substrates.³³ However, this is often not the
261 case for cellulose nanocrystal systems where the crystallite length can be shorter than its
262 periodicity. In such a case the fibrillar twists are apparently suppressed due to the drying
263 artifact as shown in Figure 1c. A systematic μ ED analysis with different surface properties of
264 CNC and supporting substrates would be necessary to fully understand the influence of
265 drying on the nanoscale morphology of such nanocrystals.

266 The drying on flat surface alters not only the twist geometry but also other nanoscale
267 morphological features of nanocelluloses, such as the flexibility of these nanoelements. The
268 occasional absence of the meridional $0\ 0\ 4$ reflections was observed in these μ ED patterns
269 recorded under cryo condition (Figure 2d - 4, 8, 9, 11) unlike those recorded under dry
270 condition where the $0\ 0\ 4$ reflection along the tCNC axis was always present (Figure 3a). The
271 absence of this reflection indicates that the c^* axis of the crystal is locally tilted with respect
272 to a nearly flat Ewald sphere surface, hence, in the real space, the tCNC is not a straight rigid
273 rod but rather consists in a somewhat wavy structure no longer flat in the vitrified ice plane.
274 This is surprising as especially short CNCs are often considered as rigid and straight bodies
275 thanks to their high axial elastic modulus. However, as recently reported, based on
276 computational simulations,^{34, 35} cellulose crystals can be considered as flexible in their
277 transverse and diagonal directions, which may lead to crystals deformation in such a waving
278 manner along their axes.

279 Another aspect of using sequential μ ED acquisition on right-handed twisted cellulose
280 nanocrystals or nanofibrils consists in allowing the identification of the crystal polarity of
281 these elements without resorting to the reducing-end silver-labeling of the cellulose
282 molecules, as described in previous studies.³⁶ Determining the crystal polarity of cellulose
283 crystals is important not only to characterize the assembled fibrillar structures composed of
284 multiple cellulose crystallites, but also to understand the organization of cellulose in plant
285 tissues. Indeed, during biosynthesis, the cellulose synthesizing complexes polymerize the
286 nascent cellulose molecules from their non-reducing end and thus give its unique polarity to
287 each nanofibril.³⁰ Despite such unidirectional organization, adjacent cellulose nanofibrils
288 have been characterized as statistically antiparallel in the cell wall of green algae, such as
289 those of *Valonia*.³⁶ This organization is likely due to the biogenesis of these nanofibrils,
290 which are generated by slender synthesizing complexes of several hundred of nm in length,
291 which themselves can align in a parallel or antiparallel manner in the plasma membrane of
292 the corresponding alga.³⁷ To date, one does not know whether such statistical distribution is
293 common to the cellulose microfibrillar assembly in all plant cell walls, in particular to the
294 more common ones where cellulose is being synthesized by the much smaller rosettes. The
295 use of sequential μ ED on bundle of nanofibrils pulled out from the corresponding plant cell
296 wall could answer such questions.

297 The twisted structures described in this report correspond to tCNCs isolated from their
298 environment. It is not clear whether these cellulose elements are also twisted in the native
299 tunicate mantle from which they have been extracted. In this line, it is interesting to mention
300 the case of *M. denticulata* cellulose as the observed twisting reported by Hanley et al.¹⁴
301 occurs only on isolated nanofibrils. In the cell wall of the corresponding alga, electron and X-
302 ray diffraction experiments have demonstrated that all nanofibrils are strictly laid flat in the
303 algal cell wall with no hint of departure from the uniplanar orientation.³⁸ This observation

304 indicates that the inherent forces leading to the twist deformation of the nanofibrils can be
305 easily counterbalanced by the surfaces forces resulting from the close contact with the
306 neighboring nanofibrils as observed in the parallel assembly of tCNCs in this study (Fig. 4).
307 While it has been suggested that the twist morphology of cellulose nanofibrils in plant cell
308 wall may hinder a coalescence of individual nanofibrils into merged fibrils,^{22,39} the
309 microdiffraction study shown here for the model samples suggests that the twisting do not
310 avoid tight adhesion of adjacent microfibrils on non-matching crystallographic surfaces. The
311 persistence of the twisted structures *in-planta* remains to be investigated.

312

313 **Conclusions**

314 This report demonstrates that the μ ED analysis allows quantitatively describing the
315 nanoscale structural features of nanocelluloses. The exact geometry of intrinsic continuous
316 twist of tCNC in aqueous suspension state was revealed for the first time under cryogenic
317 condition. With dried samples, this regular twist was significantly altered into discontinuous
318 sharp twist separated by long flat segments, giving rise to the discrepancy between the
319 apparent morphology obtained from conventional TEM and AFM imaging and molecular
320 simulations of CNCs. The similar μ ED analysis applied to a fibillar assembly of tCNCs
321 provided the structural details of this common structural unit of nanocellulose systems,
322 namely the molecular polarities and twist rates of constituent nanocrystals as well as
323 interacting surfaces between crystals. The quantitative structural description at single
324 nanoparticle level leads to establish the structure-properties relationship of various solid and
325 colloidal nanocellulose systems. Such knowledge will pave a way to use the renewable
326 cellulose for rationally designing specific biobased functional materials.

327

328 **Experimental**

329 **Sample preparation.** Cellulose nanocrystals (CNCs) were prepared according to the method
330 described by Elazzouzi-Hafraoui et al.²⁶ Briefly, *Halocynthia roretzi* tunicin was purified by
331 alternating aqueous treatments with 1 N KOH and 0.3% NaClO₂ at room temperature
332 followed by hydrolysis with 48% sulfuric acid at 55 °C for 15 h. Acid-free CNCs aqueous
333 suspensions were obtained after repeated centrifugations/washing cycles followed by
334 ultrasonication.

335

336 **Structural characterization of tCNC.** X-ray diffraction measurement was carried out on a
337 dried film of tCNC with Ni-filtered CuK α radiation ($\lambda = 0.15418 \text{ \AA}$) using a Philips PW3830
338 generator operated at 30 kV and 20 mA. FTIR measurement was performed with the same
339 dried tCNC film using a Spectrum Two spectrometer (Perkin Elmer) operated under
340 attenuated total reflection (ATR) mode. The zeta potential was measured by electrophoresis
341 coupled with laser Doppler velocimetry using a Malvern NanoZS instrument. The 0.1 wt%
342 tCNC suspension was measured at pH 7 and with 0.01 M NaCl. Data were averaged over
343 three measurements.

344

345 **Transmission electron microscopy (TEM).** Drops of about 0.001 wt% CNC aqueous
346 suspensions were deposited on glow-discharged carbon-coated copper grids. Negative
347 staining was achieved using 2 wt% uranyl acetate aqueous solution. Metal shadowing was
348 performed with a BAL-TEC MED 020 coating system (Leica microsystems, Germany), using
349 tungsten/tantalum alloy sputtered from an electron gun, with a shadowing angle of 30°.
350 Specimens for observations under cryogenic conditions were prepared by quench-freezing in
351 liquid ethane by using an automatic plunge freezer (EM-GP, Leica Microsystems, Germany).
352 Transmission electron microscopy was performed using a JEM-2100Plus (JEOL Ltd., Japan)
353 operated at an accelerating voltage at 200 kV. All electron micrographs and electron

354 diffraction patterns were recorded on a Gatan Rio 16 camera (Gatan Inc., USA). An Elsa
355 cryo-transfer holder (Gatan Inc., USA) was used for cryoEM observations. Low-dose bright-
356 field imaging and microED (μ ED) measurements were achieved using the SerialEM
357 program.⁴⁰ Micrographs and μ ED patterns were analyzed using the Fiji program.⁴¹

358

359 **Molecular dynamics (MD) simulation.** MD simulation of cellulose nanocrystals was
360 performed using Gromacs version 5⁴² and Gromos 56Acarbo force field.⁴³ The nanocrystals
361 were composed of 18 molecular chains having 80 glucose residues. The details of the
362 simulation are described elsewhere.³⁴

363

364 **Conflicts of interest**

365 There are no conflicts to declare.

366 **Acknowledgements**

367 I would like to thank Drs. H. Chanzy, Y. Nishiyama, K. Mazeau, J-L Putaux, and S.
368 Vignolini for suggestions during the writing of this work. I also would like to thank Drs.
369 Bruno Jean, Christine Lancelon-Pin and Pierre Saillar for their help in zeta-potential
370 measurement, metal shadowing, and drawing of schematic illustration. The NanoBio-ICMG
371 platform (FR 2607) is acknowledged for granting access to the electron microscopy facility.

372

373 **Supplementary Information**

374 Supplementary information file is available online.

375

376 **References**

377 1. Y. Habibi, L. A. Lucia, and O. J. Rojas, *Chem. Rev.* **110**, 3479–3500 (2010).

- 378 2. D. Klemm, B. Heublein, H-P. Fink, and A. Bohn, *Angew. Chem. Int. Edit.* **44**, 3358–
379 3393 (2005).
- 380 3. R. J. Moon, A. Martini, J. Nairn, J. Simonsen, and J. Youngblood, *Chem. Soc. Rev.*
381 **40**, 3941–3994 (2011).
- 382 4. A. Isogai, *J. Wood Sci.* **59**, 449–459 (2013).
- 383 5. I. Sakurada, Y. Nukushina, and T. Ito, *J. Polym. Sci.* **37**, 651–660 (1962).
- 384 6. T. Nishino, K. Takano, and K. Nakamae, *J. Polym. Sci. Part B Polym. Phys.* **33**,
385 1647–1651 (1995).
- 386 7. I. Diddens, B. Murphy, M. Krisch, and M. Müller, *Macromolecules* **41**, 9755–9759
387 (2008).
- 388 8. Y. Nishiyama, *J. Wood Sci.* **55**, 241–249 (2009).
- 389 9. T. Saito, R. Kuramae, J. Wohler, L. A. Berglund, and A. Isogai, *Biomacromolecules*
390 **14**, 248–253 (2013).
- 391 10. B. D. Wilts, A. G. Dumanli, R. Middleton, P. Vukusic, and S. Vignolini, *APL*
392 *Photonics* **2**, 040801 (2017).
- 393 11. M. Kaushik, K. Basu, C. Benoit, C. M. Cirtiu, H. Vali, and A. Moores, *J. Am. Chem.*
394 *Soc.* **137**, 6124–6127 (2015).
- 395 12. J-F. Revol, H. Bradford, J. Giasson, R. H. Marchessault, and D. G. Gray, *Int. J. Biol.*
396 *Macromol.* **14**, 170–172 (1990).
- 397 13. R. M. Parker, G. Guidetti, C. A. Williams, T. Zhao, A. Narkevicius, S. Vignolini, and
398 B. Frka-Petesic, *Adv. Mater.* **30**, 1704477 (2018).
- 399 14. S. J. Hanley, J-F. Revol, L. Godbout, and D. G. Gray, *Cellulose* **4**, 209–220 (1997).
- 400 15. I. Usov, G. Nyström, J. Adamcik, S. Handschin, C. Schütz, A. Fall, L. Bergström, and
401 R. Mezzenga, *Nat. Commun.* **6**, 7564-1–7564-11 (2014).

- 402 16. M. Arcari, E. Zuccarella, R. Axelrod, J. Adamcik, A. Sánchez-Ferrer, R. Mezzenga,
403 and G. Nyström, *Biomacromolecules* (2019).
- 404 17. W. J. Orts, L. Godbout, R. H. Marchessault, and J-F. Revol, *Macromolecules* **31**,
405 5717–5725 (1998).
- 406 18. J. Majoinen, J. Haataja. D. Appelhans. A. Lederer, A. Olszewska, J. Seitsonen, V.
407 Aseyev, E. Kontturi, H. Rosilo, M. Österberg. N. Houbenov, and O. Ikkala. *J. Am. Chem. Soc.*
408 **136**, 865–869 (2013).
- 409 19. J. F. Matthews, C. E. Skopec, P. E. Mason, P. Zuccato, R. W. Torget, J. Sugiyama, M.
410 W. Himmel, and J. W. Brady, *Carbohydr. Res.* **341**, 138–152 (2006).
- 411 20. S. Paavilainen, T. Rog, and I. Vattulainen, *J. Phys. Chem. B* **115**, 3747–3755 (2011).
- 412 21. J. A. Hadden, A. D. French, and R. J. Woods, *Biopolymers* **99**,746–756 (2013).
- 413 22. L. Bu, M. E. Himmel, and M. F. Crowley, *Carbohydr. Polym.* **125**, 146–152 (2015).
- 414 23. S. K. Kannan, D. P. Oehme, M. S. Doblin, M. S. Gidley, A. Bacik, and M. T. Dowton,
415 *Carbohydr. Polym.* **175**, 433–439 (2017).
- 416 24. W. Helbert, Y. Nishiyama, T. Okano, and J. Sugiyama, *J. Struct. Biol.* **124**, 42–50
417 (1998).
- 418 25. Y. Van Daele, J-F. Revol, F. Gaill, and G. Goffinet, *Biol. Cell.* **76**, 87–96 (1992).
- 419 26. S. Elazzouzi-Hafraoui, Y. Nishiyama, J-L. Putaux, L. Heux, F. Dubreuil, and C.
420 Rochas, *Biomacromolecules* **9**, 57–65 (2008).
- 421 27. B. Frka-Petesic, J. Sugiyama, S. Kimura, H. Chanzy, and G. Maret, *Macromolecules*
422 **48**, 8844–8857 (2015).
- 423 28. J. Sugiyama, T. Okano, H. Yamamoto, and F. Horii, *Macromolecules* **23**, 3196-3198
424 (1990).
- 425 29. Y. Nishiyama, P. Langan, and H. Chanzy, *J. Am. Chem. Soc.* **124**, 9074–9082 (2002).

- 426 30. M. Koyama, W. Helbert, T. Imai, J. Sugiyama, and B. Henrissat, *Proc. Natl. Acad.*
427 *Sci. U.S.A.* **94**, 9091–9095 (1997).
- 428 31. A. Sarko, and R. Muggli, *Macromolecules* **7**, 486–494 (1974).
- 429 32. Z. Zhao, O. E. Shklyae, A. Nili, M. N. A. Mohamed, J. D. Kubicki, V. H. Crespi,
430 and L. Zhong, *J. Phys. Chem. A* **117**, 2580–2589 (2012).
- 431 33. S. Assenza, J. Adamcik, R. Mezzenga, and P. De Los Rios, *Phys. Rev. Lett.* **113**,
432 268103 (2014).
- 433 34. P. Chen, Y. Ogawa, Y. Nishiyama, A. E. Ismail, and K. Mazeau, *Phys. Chem. Chem.*
434 *Phys.* **18**, 19880–19887 (2016).
- 435 35. G. Molnar, D. Rodney, F. Martoia, P. J. J. Dumont, Y. Nishiyama, K. Mazeau, and L.
436 Orgeas, *Proc. Natl. Acad. Sci. U.S.A.* **115**, 7260–7265 (2018).
- 437 36. N-H. Kim, T. Imai, M. Wada, and J. Sugiyama, *Biomacromolecules* **7**, 274–280
438 (2006).
- 439 37. T. Itoh, and R. M. Brown, *Planta* **160**, 372-381 (1984).
- 440 38. N-H. Kim, W. Herth, R. Vuong, and H. Chanzy, *J. Struct. Biol.* **117**, 195-203 (1996).
- 441 39. A. N. Fernandes, L. H. Thomas, C. M. Altaner, P. Callow, V. T. Forsyth, D. C.
442 Apperley, C. J. Kennedy, and M. C. Jarvis, *Proc. Natl. Acad. Sci. U.S.A.* **108**, 1195-1203
443 (2011).
- 444 40. D. N. Mastronarde, *Microsc. Microanal.*, **9**, 1182–1183 (2003).
- 445 41. J. Schindelin, I. Arganda-Carreras, I. E. Frise, et al. *Nat. Methods* **9**, 676-682 (2012).
- 446 42. M. J. Abraham, T. Murtola, R. Schulz, S. Páll, J. C. Smith, B. Hess, and E. Lindahl,
447 *SoftwareX*, **1**, 19–25 (2015).
- 448 43. H. S. Hansen, and P. H. Hünenberger, *J. Comp. Chem.* **32**, 998–1032 (2011).