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ARTICLE

Amyloid-like fibrils formed from intrinsically disordered caseins: physicochemical and nanomechanical properties

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Amyloid-like fibrils are studied because of their significance in understanding pathogenesis and creating functional materials. Amyloid-like fibrils have been studied by heating globular proteins at acidic conditions. In the present study, intrinsically disordered α -, β -, and κ -caseins were studied to form amyloid-like fibrils at pH 2.0 and 90 °C. No fibrils were observed for α -caseins, and acid hydrolysis was found to be the rate-limiting step of fibrillation of β - and κ -caseins. An increase of β -sheet structure was observed after fibrillation. Nanomechanical analysis of long amyloid-like fibrils using peak-force quantitative nanomechanical atomic force microscopy showed the lowest and highest Young's modulus for β -casein (2.35 ± 0.29 GPa) and κ -casein (4.14 ± 0.66 GPa), respectively. The dispersion with β -casein fibrils had a viscosity more than 10 and 5 times higher than those of κ -casein and β -lactoglobulin, respectively, at 0.1 s^{-1} at comparable concentrations. The current findings may assist not only the understanding of amyloid fibril formation but also the development of novel functional materials from disordered proteins.

Keywords: amyloid-like fibrils, intrinsically disordered protein, casein, nanomechanical properties, viscosity

1 Introduction

2 Amyloid fibrils formed in human tissues such as brain are
3 significant to the understanding of pathogenesis of diseases
4 related to neurodegenerative disorders and amyloidosis like
5 Alzheimer's and Parkinson's diseases.¹ The formation
6 mechanism of amyloid fibrils has been investigated
7 extensively *in vivo* and *in vitro* with the hope to find a
8 preventive therapy or revert fibril structures. Although it
9 has not been identified conclusively, there are also some
10 amyloid-like fibrils found in the mammary tissue of
11 animals such as bovine, which cause the reduction of milk
12 secretion.² It is generally believed that the formation of
13 amyloid or amyloid-like fibrils involves the nucleation of
14 partially unfolded proteins or peptides, followed by
15 extension reactions to form beta-sheets through anti-parallel
16 stacking of beta-strands by inter-strand hydrogen bonds.^{3,4}
17 Besides proteins related to pathogenesis, several globular
18 proteins with a significant portion of β -sheet structures

19 have been observed to form amyloid-like fibrils *in vitro*,
20 such as β -lactoglobulin, yeast prions, α -lactalbumin,
21 ovalbumin, and lysozyme.⁵⁻¹¹ Instead of applying
22 physiological conditions, heating these globular proteins at
23 acidic conditions is commonly used to form amyloid-like
24 fibrils that can be used to unveil the fibrillation process.
25 Studies have shown that the *in vitro* fibrillation involves
26 steps of acid hydrolysis, nucleation and growth of acid-
27 hydrolyzed peptides by β -sheet alignment.^{6, 12} β -
28 lactoglobulin is among the most extensively studied
29 globular proteins and can form amyloid-like fibrils at a
30 wide pH range below its isoelectric point (pI) of around pH
31 5.3 by heating.¹³ A recent analysis using atomic force
32 microscopy (AFM) demonstrated a comparable
33 amyloidosis process between heat-denatured β -
34 lactoglobulin and amyloid fibrils.¹⁴ Several parameters
35 affect the rate and type of fibril formation. Longer fibrils
36 are formed at a higher rate when heated at a lower ionic
37 strength, a higher temperature, and without co-solutes such

38 as polysaccharides and polyols.^{12, 15-18} These amyloid-like
39 fibrils also have potential for use as functional materials to
40 provide unique rheology, encapsulate compounds, and
41 serve as scaffolds in tissue engineering.^{9, 19, 20}

42 *In vitro* fibril formation from amorphous proteins is
43 relatively scarce.¹ Caseins abundant in mammalian milk
44 lack of defined three-dimensional structures and are
45 classified as intrinsically disordered proteins.²¹ Several
46 studies have characterized the properties of bovine caseins
47 in forming fibrils. In bovine milk, caseins consist of about
48 80% of proteins²² and four types of caseins (α_{s1} , α_{s2} , β , and
49 κ) assemble as mostly spherical colloidal particles with a
50 diameter of 50-250 nm.²³ Caseins have less secondary or
51 tertiary structures than globular proteins under
52 physiological conditions and can function as molecular
53 chaperones to stabilize proteins through electrostatic and
54 hydrophobic interactions.^{24, 25} Caseins also have numerous
55 applications based on their balanced
56 hydrophilicity/hydrophobicity and self-assembling
57 properties.²⁶⁻²⁹ κ -casein can form amyloid-like fibrils under
58 physiological conditions with or without a reducing agent.³⁰
59 The fibrillation involves the formation of multimers from
60 individual κ -casein molecules and conformational
61 rearrangement to form more organized species, which
62 subsequently undergo micellar assembling and further
63 growing to form amyloid-like fibrils.^{30, 31} The fibrillation of
64 κ -casein can be inhibited by α_s - and β -caseins, which
65 results in the absence of amyloid-like fibrils when different
66 types of caseins are present under physiological
67 conditions.³⁰ The fibrillation of α_{s2} -casein at physiological
68 conditions was also observed but was inhibited by β - or
69 α_{s1} -casein.^{30, 32} These findings collectively suggest the
70 prevention of amyloidosis accumulation in mammary
71 tissues and the development of corpora amylacea.^{30, 32} It
72 was also observed that α_{s1} -casein inhibits fibrillogenesis of
73 A β 1-40, a peptide involved in Alzheimer's disease, by
74 increasing the nucleation lag-time and slowing down the
75 overall fibrillation.³³ Currently, formation of amyloid-like
76 fibrils by heating caseins at acidic conditions has not been
77 studied.

78 In the present study, the objective was to study amyloid-
79 like fibrils produced from bovine caseins by heating at 90
80 °C and pH 2.0, and compare their physicochemical and
81 nanomechanical properties with fibrils formed from β -
82 lactoglobulin. The fibrillation kinetics was monitored using
83 Thioflavin T (Th-T) fluorescence spectroscopy for up to 48
84 h. The morphology of fibrils was characterized using
85 transmission electron microscopy (TEM) and AFM). The
86 hydrolysis kinetics was quantified using sodium dodecyl
87 sulfate-polyacrylamide gel electrophoresis (SDS-PAGE),
88 and changes of secondary structures were studied using
89 Fourier transform infrared (FTIR) spectroscopy and circular
90 dichroism (CD) spectroscopy. Finally, rheological
91 properties of fibril dispersions formed from different
92 proteins were characterized.

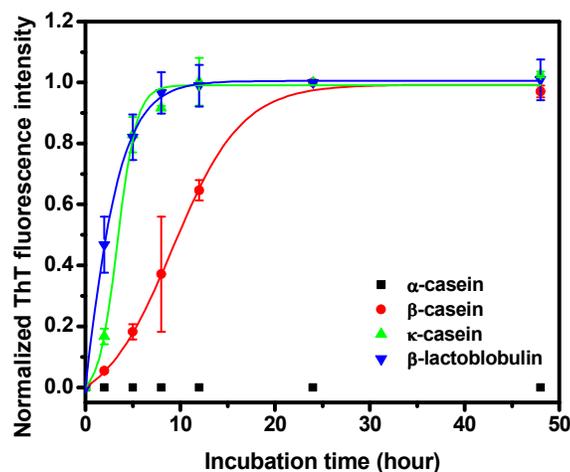
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94 Results and discussions

95 Amyloid-like fibril formation kinetics

96 Amyloid-like fibrils are formed by sequentially stacking of
97 beta-strands to beta-sheets perpendicularly to the main axil
98 of the fibril.^{3, 4} Because the selective binding between ThT
99 and fibrils strengthens the emitted fluorescence intensity,
100 the quantified fluorescence intensity can be used to monitor
101 the growth of amyloid-like fibrils.^{34, 35} Figure 1 shows the
102 normalized ThT fluorescence intensity of samples after heating
103 α -, β -, and κ -caseins and β -lactoglobulin for different
104 durations. Heating α -casein at 90 °C and pH 2.0
105 did not change ThT fluorescence, indicating the absence of
106 fibrils. The α -casein in the current study is a mixture of α_{s1} -
107 and α_{s2} -caseins. The mixed α -casein does not form fibrils at
108 physiological conditions, but the purified α_{s2} -casein does,
109 which indicates the suppression of fibril formation from
110 α_{s2} -casein by α_{s1} -casein.^{30, 32} Unfortunately, we were not
111 able to find purified α_{s1} - and α_{s2} -caseins to verify similar
112 inhibition occurred during heating at pH 2.0.

113 κ -Casein and β -lactoglobulin showed similar and typical
114 sigmoidal increase in ThT fluorescence intensity at
115 extended heating durations (Figure 1). The kinetic
116 parameters after fitting data to Eqs. 2-4 are summarized in
117 Table 1. Both proteins reached a plateau after heating at 90
118 °C for 12 h, which is similar to previous studies.^{10, 36}
119 Different from κ -casein and β -lactoglobulin, β -casein
120 showed a delayed increase in ThT fluorescence in the first
121 12 h and did not reach an equilibrium after 24 h. Therefore,
122 each casein showed distinctly different properties in
123 forming amyloid-like fibrils during heating at pH 2.0.



124 **Figure 1.** Normalized ThT fluorescence of α -casein, β -casein, κ -
125 casein, and β -lactoglobulin solutions after heating at 90 °C and pH
126 2.0 for 0-48 h at a protein concentration of 10 mg/mL. Error bars
127 are standard deviations from triplicates. Solid lines indicate the
128 fitting of data to Eq. 2.

131 Table 1. Fibrillation parameters estimated from ThT
132 fluorescence based on Eqs. 2-4.

Protein	t_{lag} (h)	$t_{1/2max}$ (h)	Adjusted r^2
β -casein	2.96	9.74	0.9983
κ -casein	1.51	3.45	0.9924

β -lactoglobulin	0.05	2.21	0.9994
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133 Acid hydrolysis kinetics of proteins

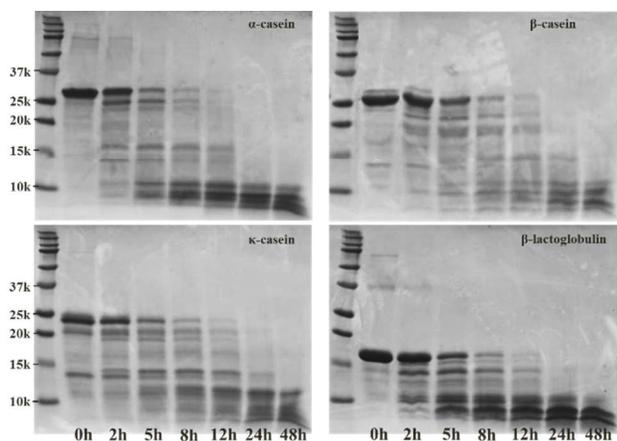
134 The fibril formation kinetics can be affected by several
 135 parameters such as pH and temperature that determine the
 136 kinetics of acid hydrolysis, as well as nucleation and
 137 growth which are a function of both hydrolysate structures
 138 and environment conditions.^{15, 37} To understand the rate-
 139 determining step during fibril formation, SDS-PAGE was
 140 used to monitor acid hydrolysis during heating (Figure 2).
 141 Hydrolysis of proteins to peptides smaller than 10 kDa
 142 mostly completed in the first 12 h, with β -casein being the
 143 slowest. The hydrolysis kinetics quantified from SDS-
 144 PAGE after fitting the model in Eq. 5 is shown in Figure 3,
 145 and the estimated rate constants are presented in Table 2. α -
 146 Casein was hydrolyzed the fastest, with a k_h of $0.332 \pm$
 147 0.008 h^{-1} . κ -Casein and β -lactoglobulin showed similar
 148 hydrolysis kinetics with k_h of $0.301 \pm 0.022 \text{ h}^{-1}$ and $0.310 \pm$
 149 0.015 h^{-1} , respectively. The k_h of β -casein hydrolysis (0.228
 150 $\pm 0.018 \text{ h}^{-1}$) was significantly smaller than the other three
 151 proteins. The calculated hydrophobicity of proteins follows
 152 the decreasing order of β -casein > κ -casein = β -
 153 lactoglobulin > α_{s1} -casein > α_{s2} -casein.³⁸ Therefore, a more
 154 hydrophobic protein is hydrolyzed more slowly during
 155 heating at acidic conditions. This is expected because a
 156 more hydrophobic protein reduces the exposure of amide
 157 bonds for acid hydrolysis, which also is in agreement with
 158 the ThT fluorescence intensity changes (Figure 1). With the
 159 exception of α -casein (no fibril formation), acid hydrolysis
 160 is thus the rate-limiting step in fibrillation of these proteins
 161 during heating at acidic conditions.

162
 163 Table 2. Protein hydrolysis rate constant (k_h) estimated from SDS-
 164 PAGE (Figure 2).

Protein	$k_h \text{ (h}^{-1}\text{)}^*$	Adjusted r^2
α -Casein	0.332 ± 0.008	0.9970
β -casein	0.228 ± 0.018	0.9683
κ -Casein	0.301 ± 0.022	0.9748
β -lactoglobulin	0.310 ± 0.015	0.9889

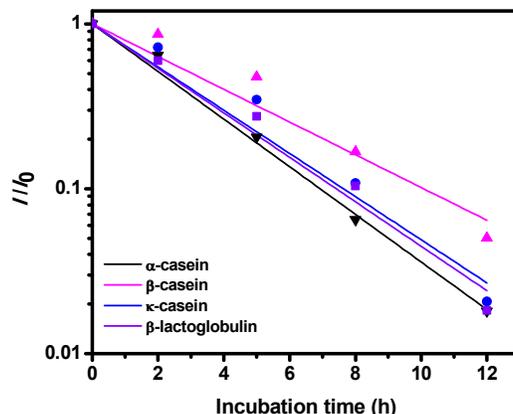
*Data are mean \pm standard error from fitting.

165
 166
 167



168

169 **Figure 2.** SDS-PAGE analysis of α -casein, β -casein, κ -casein, and
 170 β -lactoglobulin after heating at 90 °C and pH 2.0 for 0-48 h.



171
 172 **Figure 3.** Normalized SDS-PAGE band intensity showing
 173 hydrolysis kinetics of α -casein, β -casein, κ -casein, and β -
 174 lactoglobulin after heating at 90 °C and pH 2.0 for up to 12 h.

175

176 Morphology and nano-mechanical properties of fibrils

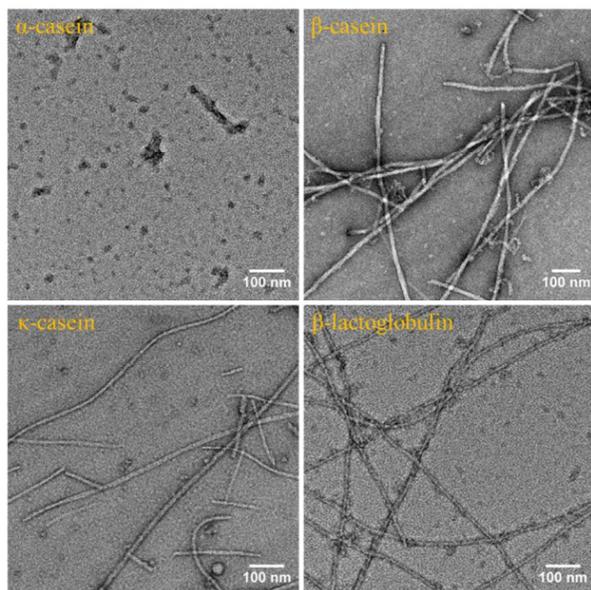
177 The morphology of proteins after heating at 90 °C and pH
 178 2.0 for 24 h was studied using TEM, shown in Figure 4. As
 179 indicated for absence of fibrils based on ThT fluorescence
 180 (Figure 1), individual particles and aggregates without
 181 amyloid-like fibrils were observed for α -casein (Figure 4).
 182 In contrast, amyloid-like fibrils were evident for β -casein,
 183 κ -casein, and β -lactoglobulin treatments.

184 AFM is also commonly used to characterize the structure
 185 of amyloid-like fibrils, and the nanoscale structures
 186 facilitate the understanding of fibrillation mechanism.^{13, 39}
 187 Characterization of nanoscale mechanical properties of
 188 materials is a unique feature of AFM, and the peak force
 189 value in the force-distance curves can be used to calculate
 190 Young's modulus in real time by fitting data in the
 191 Derjaguin–Mueller–Toporov model.^{39, 40} Figures 5-7 show
 192 morphological and mechanical properties of β -casein, κ -
 193 casein, and β -lactoglobulin before and after incubation at
 194 pH 2.0 and 90 °C for 24 h. All three proteins showed
 195 spherical structures before heating. After heating, amyloid-
 196 like fibrils with a length greater than 1 μm were observed.
 197 General observations showed that fibrils of β -lactoglobulin
 198 were overall longer than those of β -casein and κ -casein.

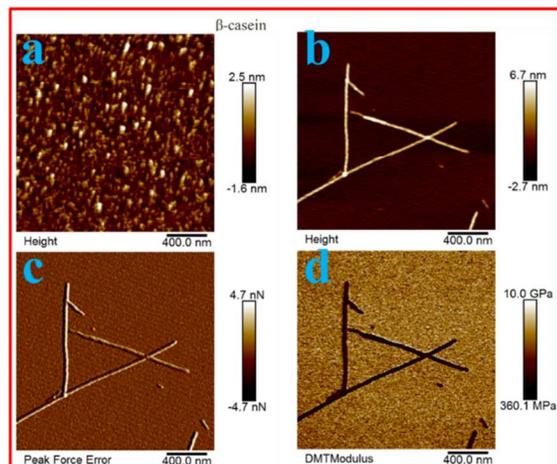
199 The average Young's moduli measured from different
 200 fibrils are summarized in Table 3. The Young's modulus of
 201 β -lactoglobulin fibrils (3.45 GPa) was close to the
 202 previously reported value of 3.7 GPa.⁴¹ The Young's
 203 modulus of β -casein fibrils (2.35 GPa) was lower than that
 204 of β -lactoglobulin but was close to that of α -synuclein and
 205 end-capped heptapeptide $\text{CH}_3\text{CONH-}\beta\text{A}\beta\text{AKLVFF-}$
 206 CONH_2 fibrils.^{41, 42} κ -Casein fibrils had the highest
 207 modulus (4.14 GPa) among the three protein fibrils. The
 208 difference in fibril modulus can be caused by the strength
 209 of intermolecular forces and can be correlated to material
 210 stiffness.⁴³ These fibrils may be used to design functional
 211 materials with different mechanical properties.

212

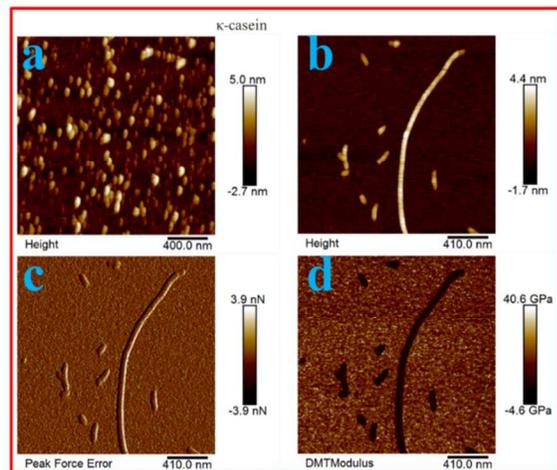
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215 **Figure 4.** TEM images showing structures of α -casein, β -casein, κ -
216 casein, and β -lactoglobulin after heating at pH 2.0 and 90 °C for 24
217 h.

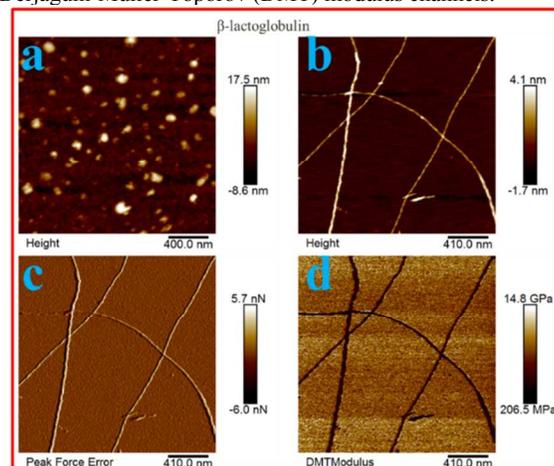


218
219 **Figure 5.** AFM images of β -casein (a) before treatment in height
220 channel and after treatment in (b) height, (c) peak force error, and
221 (d) Derjaguin-Muller-Toporov (DMT) modulus channels.
222



223

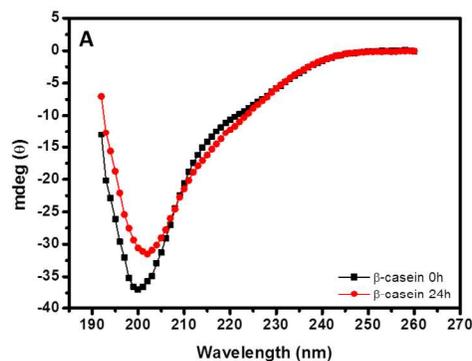
224 **Figure 6.** AFM images of κ -casein (a) before treatment in height
225 channel and after treatment in (b) height, (c) peak force error, and
226 (d) Derjaguin-Muller-Toporov (DMT) modulus channels.



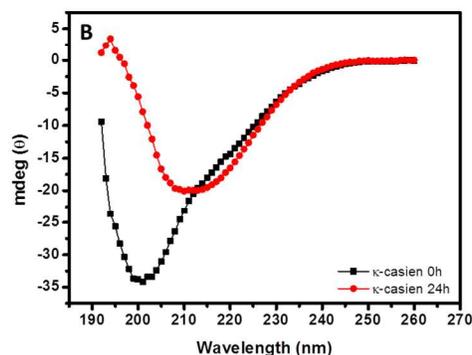
227
228 **Figure 7.** AFM images of β -lactoglobulin (a) before treatment in
229 height channel and after treatment in (b) height, (c) peak force
230 error, and (d) Derjaguin-Muller-Toporov (DMT) modulus
231 channels.
232

233 Structural changes of proteins analyzed using FTIR and 234 CD spectroscopy

235 The secondary structural changes of caseins before and
236 after heating for 24 h were analyzed using ATR-FTIR, with
237 the spectra presented in supplementary Figure S1.
238 Contrasting with no changes of α -casein, IR peaks of β -
239 κ -caseins shifted to a lower wavenumber after heating
240 (supplementary Figure S1), indicating the increased
241 contents of ordered structures.^{31, 44}



242



243

244 **Figure 8.** CD spectra of (A) β -casein and (B) κ -casein before and
 245 after heating at 90 °C and pH 2.0 for 24 h.

246
 247 To confirm with ATR-FTIR results, Far-UV CD spectra
 248 of β -casein and κ -casein were measured before and after
 249 heating for 24 h (Figure 8). The large negative ellipticity
 250 centered on 200 nm and 201 nm for β -casein and κ -casein,
 251 respectively, before heating and the low ellipticity at 220
 252 nm are characteristics of disordered structures.²⁴ After
 253 heating, there were redshifts of the minimum in ellipticity
 254 for β -casein and κ -casein to 202 nm and 210 nm,
 255 respectively. Coupled by the increase of ellipticity at 192-
 256 200 nm region and decrease at 210-200 nm region, the
 257 spectra indicated an increase in β -sheet and a reduced
 258 content of irregular structure after heating.³² The greater
 259 extent of redshift observed for the CD spectrum of κ -casein
 260 after heating indicates the greater increase of contents of
 261 regular structures than β -casein.

262
 263 **Table 3.** Average Young's modulus of β -casein, κ -casein, and β -
 264 lactoglobulin fibrils produced by heating at 90 °C and pH 2.0 for
 265 24 h measured using atomic force microscopy operated in
 266 quantitative nanomechanics mode.

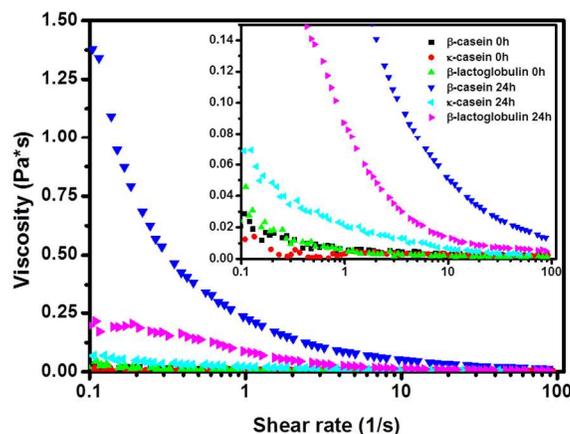
Protein	Young's Modulus (GPa)*
β -casein	2.35 ± 0.29^c
κ -casein	4.14 ± 0.66^a
β -lactoglobulin	3.45 ± 0.67^b

267 *Data are mean \pm standard deviation from 100 fibrils.

268 269 Rheological property of fibril dispersions

270 Viscosity is an important property to assess the potential of
 271 materials in applications such as texture modifiers or
 272 adhesives.⁴⁵ The viscosity of all protein samples increased
 273 after heating for 24 h, which has also been observed in
 274 other fibril systems.^{10, 18} The increased viscosity can be
 275 explained by the increased mass and hydrodynamic
 276 diameter of amyloid-like fibrils after acid hydrolysis and
 277 structure assembly.^{10, 46, 47} The apparent viscosity of κ -
 278 casein fibril dispersion at 0.1 s^{-1} ($\sim 0.1 \text{ Pa}\cdot\text{s}$) was about one-
 279 half of the β -lactoglobulin fibril sample. The β -casein fibril
 280 dispersion showed the highest viscosity and was more than
 281 one decade higher than the viscosity of κ -casein fibril
 282 dispersion at 0.1 s^{-1} . There was a negative correlation
 283 between Young's modulus and viscosity among β -casein,
 284 κ -casein, and β -lactoglobulin ($p < 0.05$). For a model
 285 system studied with actin, the maximum shear strain that a
 286 network can withstand was found to decrease with the
 287 increase of modulus or chain stiffness, and vice versa.⁴⁸
 288 Therefore, this correlation may be interpreted by the
 289 flexibility of fibrils that is negatively correlated to Young's
 290 modulus (Table 1).

291
292
293
294



295
296 **Figure 9.** Viscosity of dispersions with 10 mg/mL β -casein, κ -
 297 casein, and β -lactoglobulin before and after heating at 90 °C and
 298 pH 2.0 for 24 h. The inset shows the partial data up to 0.15 Pa·s.
 299

300 Conclusions

301 In conclusion, the present study suggests for the first time
 302 that amyloid-like fibrils can be formed from β - and κ -
 303 caseins in a controlled manner during heating at acidic pH,
 304 especially β -casein that has long been used as a molecular
 305 chaperone to inhibit the formation of amyloid-like fibrils *in*
 306 *vitro*. The fibrillation of caseins was limited by the acid
 307 hydrolysis rate and resulted in more organized structures
 308 with increased contents of β -sheet. Fibrils of β - and κ -
 309 caseins differed in nanomechanical properties and bulk
 310 viscosity, suggesting the potential of κ -casein fibrils as
 311 novel biomaterials to provide mechanical strength and β -
 312 casein fibrils as an effective texture modifier or adhesive.
 313 The current findings may be important to not only the
 314 enhanced understanding of amyloid formation but also the
 315 exploration of novel biomaterials.

316

317 Experimental methods

318 Chemicals

319 α -, β -, and κ -caseins and β -lactoglobulin were purchased
 320 from Sigma Aldrich Corp. (St. Louis, MO) and were used
 321 without further purification. Polyacrylamide gels and
 322 protein markers were purchased from Bio-Rad Laboratories
 323 Inc. (Hercules, CA). Dialysis tubing with a nominal
 324 molecular weight cutoff (MWCO) of 6000-8000 Da and β -
 325 mercaptoethanol were purchased from Sigma-Aldrich Corp.
 326 (St. Louis, MO). Other chemicals were of analytical grade
 327 and were purchased from Fisher Scientific, Inc. (Pittsburgh,
 328 PA).

329 Preparation of fibrils

330 Each protein was dissolved at 10 mg/mL in deionized
 331 water, and the pH was adjusted to 2.0 using 1.0 M HCl.
 332 Samples were heated at 90 °C for up to 48 h by incubating
 333 in the glycerol bath heated on a hot magnetic stir pate
 334

operating at 300 rpm. Samples were transferred and quenched in an ice bath at pre-determined time intervals.

Thioflavin T (ThT) fluorescence spectroscopy

The ThT fluorescent properties of samples were characterized using spectrofluorometry (model RF-1501, Shimadzu Corp., Tokyo, Japan) as an indicator of fibril formation.^{34,35} 50 μ L of a protein sample was mixed with 5 mL of the ThT working solution (0.8 mg/mL). The emission spectra at a range of 460–700 nm were acquired using an excitation wavelength of 440 nm. The fluorescence of protein samples was calibrated by subtracting the background of the ThT working solution without protein. The maximum fluorescence intensity of a sample (F_m) was used to calculate the normalized intensity by that of the same protein sample produced after 24-h heating (F_{24}) that showed the highest intensity among the time points:

$$\text{Normalized intensity} = \frac{F_m - F_0}{F_{24} - F_0} \quad (1)$$

where F_0 is the maximum fluorescence intensity of a sample before heating.

The ThT fluorescence data were fitted with the following model:^{12,49}

$$f_t = \alpha - \frac{\frac{\beta}{\gamma} + \alpha}{1 + \frac{\beta}{\alpha\gamma} \exp[t(\beta + \alpha\gamma)]} \quad (2)$$

where f_t is the normalized fluorescence intensity at time t , and α , β , and γ are arbitrary constants. The lag phase length (t_{lag}) and the time required to reach one-half of the maximum fluorescence ($t_{1/2\text{max}}$) were calculated using Eq. (3) and (4), respectively.

$$t_{\text{lag}} = \frac{1}{\beta + \alpha\gamma} \left(\ln\left(\frac{\alpha\gamma}{\beta}\right) - 4 \frac{\alpha\gamma}{\beta + \alpha\gamma} + 2 \right) \quad (3)$$

$$t_{1/2\text{max}} = \frac{\ln\left(2 + \frac{\alpha\gamma}{\beta}\right)}{\beta + \alpha\gamma} \quad (4)$$

Transmission electron microscopy (TEM)

The morphology of fibrils produced after 24-h heating was studied using TEM. The freshly prepared sample was diluted 100 times with deionized water. One drop of a diluted sample was placed on a freshly glow-discharged carbon film on a 400-mesh copper grid and then stained with 1% uranyl acetate. Samples were imaged with a Zeiss Libra TEM operating at 200 kV (Carl Zeiss Microscopy, Oberkochen, Germany).

Atomic force microscopy (AFM)

The topography and nano-mechanical properties of fibrils were characterized using AFM. 10 μ L of a 200-fold diluted sample was spread evenly onto a freshly cleaved mica sheet mounted on a sample disk (Bruker Corp., Santa Barbara, CA), and was rinsed with 1 mL deionized water previously adjusted to pH 2.0 after incubation at room temperature for

1 minute. A rectangular cantilever having an aluminum reflective coating on the backside and a quoted force constant of 40 N/m (RTESPA, Bruker Corp.) was used on a Multimode VIII microscope (Bruker AXS, Billerica, MA). The peak-force quantitative nanomechanical mode was used to scan the sample,^{41,50} and topography images were generated with a preset scan area of 2.0 \times 2.0 μ m at a scanning speed of 1 Hz. A relative method was used to calibrate the cantilever on the polystyrene film with a 2.7 GPa nominal modulus.

Sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE)

A precast 15% gradient polyacrylamide gel from Bio-Rad Laboratories, Inc. (Hercules, CA) was used in SDS-PAGE. 40 μ L of a protein sample was mixed with 200 μ L of a SDS-PAGE sample buffer (catalog number MB01015, GenScript Corp., Piscataway, NJ). After heating at 95 $^{\circ}$ C for 5 min, 10 μ L of each sample was loaded to the gel well for electrophoresis at 200 V till the indicator dye reached the gel bottom. The intensity of the band corresponding to each protein after heating (I) for a duration of t was quantified using ImageJ software (NIH, Bethesda, MD) and was normalized by the intensity of unheated protein (I_0) run on the same gel. The data were analyzed using the first-order kinetics:¹²

$$I = I_0 * e^{-k_h * t} \quad (5)$$

where k_h is the hydrolysis rate constant.

Circular dichroism (CD) spectroscopy.

The instrument used in the far-UV CD spectroscopy was a Jasco model J-815 CD spectrometer (Jasco, Inc., Easton, MD). Fibrils were dialyzed using a membrane with a MWCO of 6000-8000 Da at 4 $^{\circ}$ C for 4 days with mild stirring against water acidified to pH 2.0. The concentrations of proteins and fibrils were measured using the bichinonic acid assay method after 10-fold dilution in deionized water previously adjusted to pH 2. The diluted sample was scanned 10 times at a rate of 50 nm/min from 192 to 260 nm with a bandwidth of 1 nm.

Fourier-transform infrared spectroscopy (FTIR)

The structural changes of caseins before and after heating at 90 $^{\circ}$ C for 24 h were studied for FTIR following previously reported methods with modifications.^{44,51} Briefly, the liquid samples after dialysis as above were dried on a mica surface and the spectra were acquired using an ATR-FTIR Nicolet Nexus 670 IR spectrometer (Thermo Nicolet Corp., Madison, MI) which was equipped with a Germanium attenuated total reflection (ATR) accessory. The IR transmittance was acquired from 1600 to 1700 cm^{-1} with a resolution of 2 cm^{-1} . 100 repeated scans were undertaken for each sample. All signals were collected against a background spectrum recorded from the mica substrate only.

- 441
442
443 **Rheology**
444 The viscosity of samples before and after heating was
445 measured using a stress-controlled rheometer (model
446 AR2000, TA Instruments Inc., New Castle, DE). The cone-
447 plate geometry with a cone angle of 1° and a cone diameter
448 of 40 mm was used in shear rate ramps from 0.1 to 100 s⁻¹
449 at 25 °C.
450
451 **Statistical analysis**
452 Statistical analyses were performed using the SAS program
453 (version 9.3, SAS Institute, Cary, NC). One-way analysis of
454 variance (ANOVA) was carried out. Differences between
455 pairs of means were compared using a Tukey's test. The
456 significance level was set at 0.05.
457
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463 **Notes and references**
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