



Distinguishing anti-PEG antibodies by specificity for the PEG terminus using nanoarchitectonics-based antibiofouling cello-oligosaccharide platforms

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1 Distinguishing anti-PEG antibodies by specificity
2 for the PEG terminus using nanoarchitectonics-
3 based antibiofouling cello-oligosaccharide
4 platforms†

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12

13 ABSTRACT

14 The conjugation of poly(ethylene glycol) (PEG) to therapeutic proteins or nanoparticles is a widely
15 used pharmaceutical strategy to improve their therapeutic efficacy. However, conjugation can
16 make PEG immunogenic and induce the production of anti-PEG antibodies, which decreases both
17 the therapeutic efficacy after repeated dosing and clinical safety. To address these concerns, it is

18 essential to analyze the binding characteristics of anti-PEG antibodies to PEG. However,
19 distinguishing anti-PEG antibodies is still a difficult task. Herein, we demonstrate the use of
20 antibiofouling cello-oligosaccharide assemblies tethering one-terminal methoxy oligo(ethylene
21 glycol) (OEG) ligands for distinguishing anti-PEG antibodies in a simple manner. The OEG
22 ligand-tethering two-dimensional crystalline cello-oligosaccharide assemblies were stably
23 dispersed in a buffer solution and had antibiofouling properties against nonspecific protein
24 adsorption. These characteristics allowed enzyme-linked immunosorbent assays (ELISAs) to be
25 simply performed by cycles of centrifugation/redispersion of aqueous dispersions of the
26 assemblies. The simple assays revealed that the specific OEG ligand-tethering assemblies could
27 distinguish anti-PEG antibodies to detect a specific antibody that preferentially binds to the
28 methoxy terminus of the PEG chain with 3 repeating ethylene glycol units. Furthermore,
29 quantitative detection of the antibodies was successfully performed with high sensitivity even in
30 the presence of serum. The detectable and quantifiable range of antibody concentrations covered
31 those required clinically. Our findings open a new avenue for analyzing the binding characteristics
32 of anti-PEG antibodies in biological samples.

33

34 **1. Introduction**

35 The conjugation of poly(ethylene glycol) (PEG) to therapeutic proteins or nanoparticles, such
36 as liposomes and lipid nanoparticles, which is termed PEGylation, prolongs the blood circulation
37 time of the therapeutic component and shields immunogenic epitopes on the protein.¹⁻³ More than
38 twenty PEGylated drugs and nanoparticles have been approved by the US Food and Drug
39 Administration (FDA) for clinical applications. Generally, PEG is a biocompatible water-soluble
40 polymer; however, once conjugated to drugs and nanoparticles, PEG has actually been

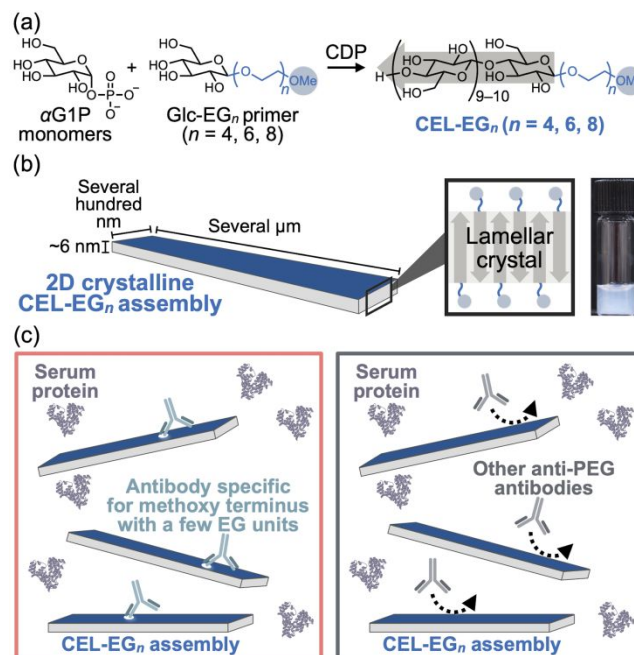
41 demonstrated to be weakly immunogenic and can induce the production of anti-PEG antibodies.^{4,5}
42 Anti-PEG antibodies are also found in healthy people who have never been exposed to PEGylated
43 therapeutics but to forms of PEG in daily products, such as cosmetics, shampoos, and
44 toothpastes.^{6,7} The retention of anti-PEG antibodies in patients potentially hinders the therapeutic
45 efficacy and safety of PEGylated therapeutics. Anti-PEG antibodies interact with PEGylated
46 therapeutics to form complexes that activate complement and cause an accelerated blood clearance
47 (ABC) phenomenon.^{8,9} Moreover, anti-PEG antibodies may cause severe allergic reactions in
48 some individuals upon the administration of PEGylated therapeutics, including PEGylated protein
49 drugs and COVID-19 mRNA vaccines.¹⁰⁻¹²

50 To develop technologies that reduce the adverse effects of PEGylated therapeutics, it is
51 essential to analyze the binding characteristics of anti-PEG antibodies to PEG. Anti-PEG
52 antibodies are a group of antibodies that recognize different parts of PEG chains and are broadly
53 classified into two groups: backbone-specific anti-PEG antibodies and methoxy-specific anti-PEG
54 antibodies. Backbone-specific anti-PEG antibodies bind to the backbone of the PEG chain with at
55 least 16 ethylene glycol (EG) units, whereas methoxy-specific anti-PEG antibodies bind to the
56 terminal methoxy group with a few to a few tens of EG units.¹³⁻¹⁵ This difference in specificity
57 has been revealed by a variety of methods, such as enzyme-linked immunosorbent assays
58 (ELISAs),^{13,16,17} surface plasmon resonance,¹⁸ hemagglutination,¹⁹ acoustic membrane microplate
59 technology,²⁰ and Western blot analysis.²¹ Nevertheless, distinguishing anti-PEG antibodies by
60 specificity is still a difficult task. For instance, in a typical ELISA, a series of PEGs with different
61 chain lengths and terminal groups are attached to proteins or lipids and immobilized onto surfaces
62 of microplates, after which the assay is performed with delicate procedures that include blocking
63 and washing steps for reducing the nonspecific adsorption of proteins (e.g., serum proteins or

64 enzyme-conjugated secondary antibodies for ELISAs) using appropriate surfactant
65 solutions.^{13,16,17} Overall, distinguishing anti-PEG antibodies by specificity in a simple manner
66 remains challenging.

67 Nanoarchitectonics is an arising concept to construct advanced materials via the fusion of
68 nanotechnology and other research disciplines, including supramolecular chemistry, synthetic
69 chemistry, bio-related technology, and others.²²⁻²⁵ Herein, we propose the use of
70 nanoarchitectonics-based two-dimensional (2D) cello-oligosaccharide assemblies²⁶⁻²⁸ with
71 structural designability and antibiofouling properties as platforms for ELISA. The assemblies have
72 nanoribbon morphologies and are composed of enzymatically synthesized cello-oligosaccharides
73 bearing one-terminal methoxy oligo(ethylene glycol) (OEG) chains with systematically varied
74 numbers of repeating units of 2, 4, 6, and 8; these oligosaccharides were named CEL-EG_{*n*} where
75 *n* corresponds to the number of repeating EG units.²⁹ The oligosaccharides form monolayer
76 lamellar crystals with an antiparallel molecular arrangement, in which the cello-oligosaccharides
77 adopt the cellulose II allomorph, thereby tethering the OEG ligands to both sides of the
78 nanoribbons in a 2D manner. Because of their 2D shape, the assemblies have a high specific
79 surface area with OEG ligands while remaining sufficiently large for centrifugation, allowing
80 ELISAs to be easily performed by cycles of centrifugation/redispersion of aqueous dispersions of
81 the assemblies. Moreover, these cello-oligosaccharide assemblies have antibiofouling properties
82 against proteins, which minimizes their nonspecific adsorption without the use of a surfactant. By
83 employing this nanoarchitectonics-based oligosaccharide platform that is attractive for use in
84 ELISAs, we successfully distinguished anti-PEG antibodies by specificity for the terminus of PEG
85 in a simple manner (Fig. 1).

86



87

88 **Fig. 1** (a) Synthetic scheme of CEL-EG_n (n = 4, 6, 8) and schematic illustrations for (b)

89 antibiofouling cello-oligosaccharide assemblies tethering one-terminal methoxy OEG ligands and

90 (c) distinguishing anti-PEG antibodies by the assemblies.

91

92 **2. Materials and methods**93 **2.1 Materials**

94 Bovine serum albumin (BSA, 98%, protease free) was purchased from Wako Pure Chemical

95 Industries. Anti-PEG immunoglobulin G (IgG) (clone PEG-B-47, produced in rabbit, monoclonal,

96 ab51257) and horseradish peroxidase (HRP)-conjugated anti-mouse IgG1 (produced in goats,

97 polyclonal, ab97240) were purchased from Abcam. HRP-conjugated anti-mouse IgG (produced in

98 rabbits, polyclonal, A9044), anti-PEG IgG (clone 15-2b, produced in mice, monoclonal,

99 MABS2002-25UG), and anti-PEG IgG (clone 6.3, produced in mice, monoclonal, MABS1966-

100 100UG) were purchased from Sigma–Aldrich Co. HRP-conjugated anti-rabbit IgG (produced in

101 goats, polyclonal, GTX213110-01) was purchased from GeneTex. A rabbit anti-PEG IgG ELISA

102 kit (PEGG-10) was purchased from Life Diagnostic, Inc. Fetal bovine serum was purchased from
103 BioWest and heated at 56 °C for 30 min to inactivate complement proteins. All other reagents were
104 purchased from Nacalai Tesque. The 96-well plates (clear, flat bottoms) were purchased from
105 Corning. A polypropylene tube (0.2 mL) was purchased from WATSON Bio Lab. Ultrapure water
106 (more than 18.2 MΩ·cm) was obtained from a Milli-Q Advantage A-10 system (Merck Millipore)
107 and used throughout the study.

108

109 **2.2 Enzyme-catalyzed production of cello-oligosaccharide assemblies**

110 CEL-EG_n (*n* = 2, 4, 6, and 8) assemblies, a plain cello-oligosaccharide assembly (named CEL-
111 OH), and a cello-oligosaccharide assembly that tethers loop-structured OEG chains with 5
112 repeating units to the surface (named CEL-EG₅-loop assembly) were prepared via cellodextrin
113 phosphorylase (CDP)-catalyzed reverse phosphorolysis reactions and characterized according to
114 our previously reported methods. In brief, for the preparation of the CEL-OH assembly,³⁰ α-D-
115 glucose 1-phosphate (αG1P) monomers (200 mM), D-glucose primers (50 mM), and CDP (1 U
116 mL⁻¹, derived from *Acetivibrio thermocellus* DSM 1313) were incubated in 500 mM 4-(2-
117 hydroxyethyl)-1-piperazineethanesulfonic acid buffer (pH 7.5) containing 50 μM
118 ethylenediaminetetraacetic acid at 60 °C for 3 d. For purification, the water-insoluble products
119 were washed with ultrapure water through centrifugation (20,400 *g*) and redispersion cycles to
120 remove at least 99.999% of the reaction solution. The obtained products dispersed in ultrapure
121 water were stored at 4 °C before use for matrix-assisted laser desorption ionization time-of-flight
122 (MALDI-TOF) mass spectrometry and atomic force microscopy (AFM). For ¹H nuclear magnetic
123 resonance (¹H NMR) spectroscopy, the product dispersions were lyophilized. The CEL-EG_n²⁹ and
124 CEL-EG₅-loop assemblies³¹ were similarly prepared using terminal methoxy OEG chain-bearing

125 β -D-glucose derivatives (Glc-EG_n, 50 mM) and penta(ethylene glycol) with cellobiose units at both
126 ends (Cello-BiP, 25 mM) as primers, respectively. Synthetic schemes of the primers are shown in
127 Scheme S1. The structural properties of cello-oligosaccharide assemblies are summarized in Table
128 1 and Fig. S1. The average degrees of polymerization (DP) of the cello-oligosaccharide moieties
129 ranged from approximately 9 to 11 (except for CEL-EG₅-loop), which were estimated based on
130 the integral ratios of internal (shown as 1' in Fig. S1b) and terminal (shown as 1 in Fig. S1b)
131 anomeric protons in the ¹H NMR spectra. The average DPs of cello-oligosaccharide chains in
132 CEL-EG₅-loop were estimated to be approximately 6 based on the integral ratios of anomeric
133 protons (shown as 1, 1' in Fig. S1b) and protons from the triazole rings (shown as c in Fig. S1b) in
134 the ¹H NMR spectrum. The thicknesses of the assemblies ranged from approximately 5 to 6 nm.
135 The widths of the assemblies were approximately 100 nm to several hundreds of nanometers. The
136 lengths of the assemblies were more than several hundred nanometers and showed large
137 distributions. ¹H NMR spectra were recorded at room temperature using an AVANCE III HD
138 spectrometer (500 MHz, Bruker Corporation) for cello-oligosaccharide solutions at concentrations
139 of more than 2% (w/v) in 4% sodium deuterioxide/deuterium oxide. The spectra were calibrated
140 using the water peak ($\delta = 4.79$) as an internal reference. An AXIMA-performance mass
141 spectrometer (Shimadzu Corporation) equipped with a nitrogen laser ($\lambda = 337$ nm) and pulsed ion
142 extraction using linear/positive mode was used to acquire the spectra under ambient conditions,
143 followed by calibration with Bradykinin (757.3997 Da), P₁₄R (1533.8582 Da), and ACTH
144 (2465.1989 Da). For sample preparation, cello-oligosaccharides were dispersed in
145 acetonitrile/water (1/1, v/v) containing 2,5-dihydroxybenzoic acid (2 mg mL⁻¹) and trifluoroacetic
146 acid (0.01% (v/v)), and the dispersions were deposited onto a sample target plate and dried under
147 ambient conditions. An FT/IR-4100 FTIR spectrometer (Jasco Corporation) was used to measure

148 attenuated total reflection-Fourier transform infrared (ATR-FTIR) absorption spectra of the
149 lyophilized products at a cumulative number of 100 and a resolution of 2.0 cm^{-1} under ambient
150 conditions. An SPM-9700HT (Shimadzu Corporation) was used to record AFM images in tapping
151 mode under ambient conditions. The samples for the cello-oligosaccharide assemblies were
152 prepared by spin casting the dispersions of the assemblies in water (0.001–0.005% (w/v)) on mica
153 at 600 rpm for 30 min. For the samples of the CEL-EG₄ assemblies after interaction with PEG-B-
154 47, aqueous dispersions of the assembly (0.5% (w/v) in PBS, 10 μL) were mixed with PEG-B-47
155 solutions (110 ng mL^{-1} in PBS, 100 μL) containing BSA (3.3% (w/v)). After incubation at 25 °C
156 for 1 h, the mixtures were centrifuged (20,400 g, 5 min), and 100 μL of the supernatants were
157 removed. Then, 100 μL of ultrapure water was added to redisperse the precipitates, and the
158 dispersions were centrifuged (20,400 g, 5 min), and 100 μL of the supernatants were removed.
159 This redispersion and centrifugation cycle was repeated three times to obtain the purified
160 assemblies, which were dispersed in 10 μL of ultrapure water. After adding 490 μL of ultrapure
161 water and mixing by gently pipetting, the dispersions were diluted one hundred-fold. The AFM
162 samples were prepared by spin casting the diluted dispersions on mica at 600 rpm for 30 min.

163

164 **2.3 Physical adsorption of BSA on the surfaces of the cello-oligosaccharide assemblies in PBS**

165 Aqueous dispersions of the cello-oligosaccharide assemblies in phosphate-buffered saline (PBS,
166 137 mM NaCl, pH 7.4) were mixed with PBS and BSA solutions in PBS. The final concentrations
167 of cello-oligosaccharide assemblies and BSA were 0.2% (w/v) and 20 μM , respectively. After
168 incubation at 25 °C for 1 h, the mixtures were centrifuged (20,400 g, 15 min) to precipitate the
169 assemblies. Ultraviolet–visible (UV–vis) absorption spectra for the supernatants were obtained by
170 using a V-670 spectrometer (JASCO) at room temperature. The BSA concentrations in the

171 supernatants were estimated based on the absorbance at 280 nm using an appropriate calibration
172 curve. Then, the amounts of protein adsorbed onto the assemblies were estimated by subtracting
173 the amounts in the supernatants from those in the initial solutions. To standardize the amount
174 adsorbed per the unit surface area of the assemblies, the areas for their base planes were estimated
175 from the unit cell area for the cellulose II allomorph (0.65 nm^2 per one cello-oligosaccharide chain)
176 as the apparent total surface area, similar to our previous reports.^{32–34} Notably, the surface areas of
177 the edges of the assemblies were ignored because the areas of the base planes appeared to be much
178 greater than those of the edges. The experiments were performed in triplicate.

179

180 **2.4 ELISAs of anti-PEG antibodies using cello-oligosaccharide assemblies**

181 Aqueous dispersions of cello-oligosaccharide assemblies (0.5% (w/v) in PBS, 10 μL) were mixed
182 with anti-PEG IgG solutions (11 ng mL^{-1} in PBS, 100 μL) containing BSA (3.3% (w/v)) as a
183 blocking reagent in polypropylene tubes. The final concentrations of the assemblies were 0.045%
184 (w/v) unless otherwise stated. After incubation at 25 °C for 1 h, the mixtures were centrifuged
185 (20,400 g, 5 min), and 100 μL of the supernatants were removed. Then, 100 μL of PBS was added
186 to redisperse the precipitates, and the dispersions were centrifuged (20,400 g, 5 min), and 100 μL
187 of the supernatants were removed. This washing cycle was repeated three times. Aqueous solutions
188 of HRP-conjugated anti-IgG (410 ng mL^{-1} and 440 ng mL^{-1} in PBS for PEG-B-47 and others,
189 respectively, 100 μL) containing BSA (3.3% (w/v)) were added to the resulting mixtures, and the
190 precipitates were dispersed by gentle pipetting. After incubation at 25 °C for 1 h, the mixtures
191 were centrifuged (20,400 g, 5 min), and 100 μL of the supernatants were removed. Then, 100 μL
192 of PBS was added to redisperse the precipitates, and the dispersions were centrifuged (20,400 g, 5
193 min), and 100 μL of the supernatants were removed. This washing cycle was repeated three times.

194 Then, *o*-phenylenediamine solutions (2 mg mL^{-1}) in a McIlvaine buffer solution (pH 4.0)
195 containing hydrogen peroxide (0.11% (w/v)) were added, and the enzymatic reactions were
196 performed at $25 \text{ }^{\circ}\text{C}$ for 30 min. After that, $100 \text{ }\mu\text{L}$ of sulfuric acid (1.5 M) aqueous solution was
197 added to stop the reactions. Then, $100 \text{ }\mu\text{L}$ of each reaction solution was added to a 96-well plate,
198 and the absorbance of each well was measured at 490 nm using a microplate reader (Model 680,
199 Bio-Rad Laboratories, Inc.) at room temperature. For ELISAs of PEG-B-47 at a concentration of
200 10 ng mL^{-1} , the absorbance values of the reaction solutions diluted twofold were measured. For
201 the other antibodies, the absorbance values of the reaction solutions were measured without
202 dilution. The measured absorbance values were subjected to the subtraction of the absorbance
203 obtained by performing with PBS instead of dispersions of cello-oligosaccharide assemblies. To
204 investigate the dependence of the absorbance value on PEG-B-47 concentration, ELISAs at
205 various final concentrations of PEG-B-47 ($1\text{--}100 \text{ ng mL}^{-1}$) were similarly performed in which
206 PEG-B-47 was incubated with the CEL-EG₄ assembly and BSA in the absence and presence of
207 10% fetal bovine serum. The absorbance values of the reaction solutions diluted fivefold were
208 measured and shown without the subtraction. The experiments were performed in triplicate.

209

210 **2.5 ELISAs of PEG-B-47 using microplates**

211 ELISAs of PEG-B-47 were performed using a commercially available microplate coated with
212 PEGylated BSA, in which the PEG ligand had a molecular weight of 20 kDa and a methoxy
213 terminus, according to the manufacturer's procedure. Aqueous solutions of PEG-B-47 ($1\text{--}100 \text{ ng}$
214 mL^{-1} in PBS, $100 \text{ }\mu\text{L}$) containing fetal bovine serum (10%) were added into the wells and
215 incubated using a plate shaker (150 rpm, 45 min, $25 \text{ }^{\circ}\text{C}$). After incubation, the solutions were
216 aspirated. The wells were washed with $1\times$ wash solutions (from the kit) five times and struck

217 sharply onto absorbent paper to remove all residues. HRP-conjugated IgG solutions (from the kit)
218 diluted appropriately were added into the wells and incubated using a plate shaker (150 rpm, 45
219 min, 25 °C). After incubation, washing processes were similarly performed. The 3,3',5,5'-
220 tetramethylbenzidine solutions (from the kit) were added into the wells to perform the enzymatic
221 reactions with incubation using a plate shaker (150 rpm, 20 min, 25 °C). After the reactions, 100
222 μL of the stop solutions (from the kit) were added with gentle mixing. The absorbance values of
223 the solutions diluted fivefold were measured within five minutes at 450 nm using a microplate
224 reader at room temperature. The experiments were performed in triplicate.

225

226 **3. Results and discussion**

227 **3.1 Protein adsorption properties of 2D cello-oligosaccharide assemblies**

228 Antibiofouling platforms contribute to the sensitive and selective detection of target
229 bio(macro)molecules under demanding conditions.^{35–37} Our previous studies revealed that the
230 CEL-OH assembly showed antibiofouling properties, such as protein nonadsorption.^{32–34} In this
231 study, the physical adsorption of BSA, which is the most abundant protein in serum, on the other
232 assemblies was similarly evaluated in PBS. Cello-oligosaccharide assemblies were stably
233 dispersed in PBS at an assembly concentration of 0.5% (w/v) for at least 1 h, which is a sufficient
234 amount of time for evaluating protein adsorption onto the assemblies in the dispersed state (Fig.
235 2). Fig. S2a shows the UV–vis absorption spectra of BSA solutions before and after incubation
236 with cello-oligosaccharide assemblies at a BSA concentration of 20 μM . The absorption curves
237 hardly changed with all cello-oligosaccharide assemblies used in this study, suggesting that BSA
238 was not adsorbed onto the assemblies. In fact, the amounts adsorbed were estimated to be
239 approximately 0 ng cm^{-2} (the respective estimated amounts adsorbed are shown in Fig. S2b). Thus,

240 OEG-tethering cello-oligosaccharide assemblies were expected to be excellent platforms for the
 241 sensitive and selective detection of anti-PEG antibodies.

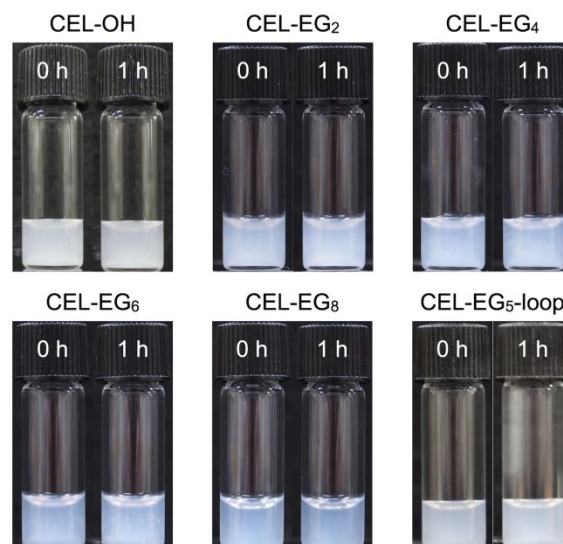
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243 **Table 1** Structural properties of the cello-oligosaccharide assemblies used in this study.

Assembly	Average degree of polymerization value of the cello-oligosaccharide moiety ^a	Thickness of the nanoribbons (nm)	Width of the nanoribbons (nm)
CEL-OH	~10	5.8 ± 0.4	Several hundreds
CEL-EG ₂	~9	5.4 ± 0.3	Several hundreds
CEL-EG ₄	~10	5.9 ± 0.5	Several hundreds
CEL-EG ₆	~10	5.9 ± 0.3	100~200
CEL-EG ₈	~11	6.3 ± 0.5	~100
CEL-EG ₅ -loop	~6 ^b	5.6 ± 0.2	Several hundreds

244 ^a Estimated from ¹H NMR spectra. ^b The value is for one of the two cello-oligosaccharide moieties
 245 in a molecular chain.

246



247

248 **Fig. 2** Photographs of dispersions of cello-oligosaccharide assemblies in PBS at an assembly
249 concentration of 0.5% (w/v) before and after incubation for 1 h.

250

251 **3.2 Distinguishing anti-PEG antibodies with the cello-oligosaccharide assemblies based on** 252 **ELISAs**

253 ELISAs of anti-PEG antibodies using microplates require delicate procedures that include
254 blocking and washing steps to decrease the nonspecific adsorption of anti-PEG antibodies, serum
255 proteins, and labeled antibodies (HRP-conjugated secondary antibodies in this assay). On the other
256 hand, in addition to the antibiofouling properties of cello-oligosaccharide assemblies as described
257 in the previous section (Fig. S2), the assemblies were readily precipitated by centrifugation and
258 redispersed in PBS for washing, which suggests the considerable potential of cello-oligosaccharide
259 platforms to perform ELISAs in a simple manner. The present ELISAs were performed in the
260 presence of 3% (w/v) BSA according to a previous study.³⁸

261 One-terminal methoxy PEG is the most frequently used for PEGylation because the chemical
262 reaction conditions are easily controlled.⁴ Consequently, anti-PEG antibodies with specificities for
263 not only the backbone PEG chain but also the terminal PEG chain containing a methoxy terminus

264 (namely, backbone-specific and methoxy-specific anti-PEG antibodies) are known to be
265 induced.^{39,40} Properties of anti-PEG antibodies used in this study are shown in Table 2. PEG-B-47
266 was used as a model target antibody, which requires a terminal methoxy PEG chain with at least
267 three repeating EG units for binding (i.e., methoxy-specific). We applied other anti-PEG
268 antibodies to the assays, 15-2b and 6.3, which require a terminal methoxy PEG chain with at least
269 twelve repeating EG units (i.e., methoxy-specific) and a PEG chain with at least sixteen repeating
270 EG units without terminal groups for binding (i.e., backbone-specific), respectively.

271 Fig. 3 shows the absorbance values at 490 nm for the products derived from HRP conjugated
272 to the secondary antibodies at an anti-PEG antibody concentration of 10 ng mL⁻¹ using cello-
273 oligosaccharide assemblies. The absorbance values correspond to the levels of anti-PEG antibodies
274 detected. In the case of the CEL-OH and CEL-EG₂ assemblies, the absorbance values were very
275 low regardless of the antibody clone, indicating that the cello-oligosaccharide assemblies hardly
276 adsorbed or detected the anti-PEG antibodies applied in the ELISAs (Fig. 3a–b). In the case of the
277 CEL-EG_{*n*} assemblies (*n* = 4, 6, and 8), absorbance values were clearly observed for PEG-B-47,
278 indicating the successful detection of PEG-B-47 (Fig. 3c–e). Considering the specificity of PEG-
279 B-47 for the terminal methoxy PEG chain with at least three repeating EG units, it was plausible
280 that the absorbance values drastically changed from the CEL-EG₂ to CEL-EG₄ assemblies. HRP-
281 conjugated secondary antibodies were hardly adsorbed onto the cello-oligosaccharide assemblies
282 used in this study (Fig. S3), supporting the conclusion that the assemblies detected PEG-B-47 with
283 high reliability based on ELISA. Moreover, the absorbance values were very low for 15-2b and
284 6.3, indicating that the CEL-EG_{*n*} assemblies (*n* = 4, 6, and 8) hardly detected the anti-PEG
285 antibodies (Fig. 3c–e), possibly due to the requirements of more repeating EG units for the binding
286 of 15-2b and 6.3. When the CEL-EG₅-loop assembly, in which the OEG ligands do not have a

287 methoxy terminus, was used, anti-PEG antibodies were not detected (Fig. 3f). These results
288 demonstrate that ELISAs using terminal methoxy OEG-tethering cello-oligosaccharide assemblies
289 allow for simply distinguishing anti-PEG antibodies by specifically identifying the PEG terminus.

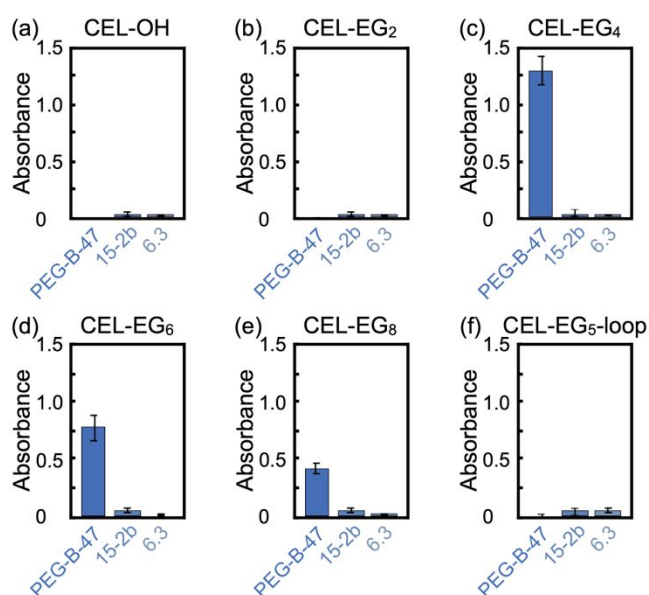
290 For PEG-B-47, the absorbance values for CEL-EG_n assemblies ($n = 4, 6, \text{ and } 8$) gradually
291 decreased with increasing number of EG units, suggesting that the binding of the anti-PEG
292 antibodies to the OEG ligands on the assemblies was suppressed with increasing OEG chain length.
293 This decrease was possibly derived from the steric hindrance and conformational restriction of the
294 surface OEG ligands. Considering that the assemblies are monolayer lamellar crystals with the
295 cellulose II allomorph, the area occupied by an OEG ligand on the assemblies can be estimated to
296 be 0.65 nm² (Fig. S4).⁴¹ On the other hand, the Flory radii for the EG₄, EG₆, and EG₈ chains were
297 estimated to be 0.76 nm, 0.97 nm, and 1.2 nm, respectively.⁴² These estimations suggest that the
298 OEG ligands interfere with each other to restrict their conformational freedoms as the number of
299 EG units increases. In other words, the steric hindrance of adjacent OEG ligands and the possible
300 conformational changes likely interfere with the binding of the anti-PEG antibodies to the longer
301 OEG ligand. It was therefore determined that the OEG ligand length on the surfaces of the cello-
302 oligosaccharide assemblies was an important factor for the effective detection of anti-PEG
303 antibodies.

304

305 **Table 2** Properties of the anti-PEG antibodies used in this study.

Antibody name	Preferred PEG structure for antibody binding	Preferred number of repeating EG units for antibody binding
PEG-B-47 ⁴³	Terminus PEG chain containing a methoxy terminus	≥ 3
15-2b ¹³	Terminus PEG chain containing a methoxy terminus	≥ 12
6.3 ¹³	Backbone PEG chain	≥ 16

306



307

308 **Fig. 3** ELISAs of anti-PEG antibodies using (a) CEL-OH, (b) CEL-EG₂, (c) CEL-EG₄, (d) CEL-
 309 EG₆, (e) CEL-EG₈, and (f) CEL-EG₅-loop at an anti-PEG antibody concentration of 10 ng mL⁻¹
 310 and an assembly concentration of 0.045% (w/v).

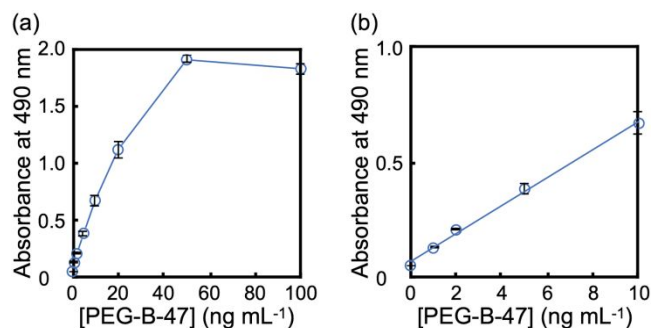
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312 The concentration of PEG-B-47 used for ELISAs with the CEL-EG₄ assemblies in PBS varied
 313 over the range of 0–100 ng mL⁻¹ (Fig. 4). The absorbance values increased with increasing
 314 antibody concentration in the range of 0–50 ng mL⁻¹, almost reaching a plateau above 50 ng mL⁻¹,
 315 resulting in a saturation profile (Fig. 4a). This observation possibly suggests that the anti-PEG

316 antibodies bind to the EG₄ ligand on the assembly surface in a monolayer. In fact, the atomic force
317 microscopy observations revealed nanometer-sized granules that could possibly be assigned to
318 PEG-B-47 on the assembly surface (Fig. S5). The absorbance value with 100 ng mL⁻¹ PEG-B-47
319 slightly decreased compared with that at 50 ng mL⁻¹. This might be attributed to a decrease in the
320 effective surface area of the assembly based on partial aggregation of the assembly with anti-PEG
321 antibodies (namely, crosslinking of the assemblies with anti-PEG antibodies).⁴⁴

322 Assuming that the dependence of the absorbance value on the PEG-B-47 concentration was
323 an adsorption isotherm, the profile in the range of 0–50 ng mL⁻¹ was fitted to the Langmuir
324 adsorption model with a coefficient of determination (R^2) value of 1.00 (Fig. S6). The apparent
325 dissociation constant (K_d) was estimated from the fitting to be 0.325 nM (this constant is discussed
326 below). In the range of 0–10 ng mL⁻¹, the plots were fitted to a linear equation with an R^2 value of
327 0.997 (Fig. 4b), suggesting that the present ELISAs can be used for quantitative anti-PEG antibody
328 detection in the concentration range. Using the slope of the linear plot and the standard deviation
329 (σ) of the absorbance value at an antibody concentration of 0 ng mL⁻¹, the limit of detection (LOD)
330 value ($= 3\sigma/\text{slope}$)⁴⁵ was estimated to be 0.065 ng mL⁻¹. Taken together, these results suggested
331 that these cello-oligosaccharide platforms have great potential to efficiently distinguish anti-PEG
332 antibodies and sensitively detect a specific anti-PEG antibody based on a simple ELISA method.

333



334

335 **Fig. 4** (a) ELISAs of PEG-B-47 using 0.045% (w/v) CEL-EG₄ assembly at different PEG-B-47
336 concentrations. (b) Linear fitting in the concentration range of 0–10 ng mL⁻¹.

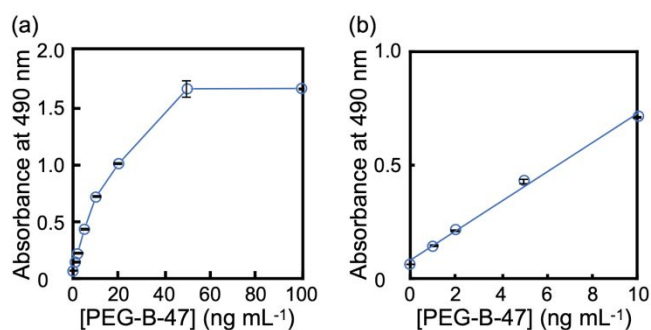
337

338 3.3 ELISAs of anti-PEG antibodies in the presence of serum

339 An essential question when analyzing the characteristics of anti-PEG antibodies is whether our
340 assays can quantify antibodies in serum samples for future practical applications. To evaluate the
341 practicability of the present ELISAs using antibiofouling cello-oligosaccharide platforms, the
342 concentration of PEG-B-47 in the ELISAs using the CEL-EG₄ assemblies was changed in the
343 range of 0–100 ng mL⁻¹ in the presence of 10% fetal bovine serum as a model serum (Fig. 5). The
344 absorbance values increased with increasing anti-PEG antibody concentration in the range of 0–
345 50 ng mL⁻¹ and then almost reached saturation above 50 ng mL⁻¹ (Fig. 5a), suggesting monolayer
346 adsorption of the anti-PEG antibodies on the assembly surface even in the presence of serum.
347 Monolayer adsorption was supported by the fact that the profile in the range of 0–50 ng mL⁻¹ fit
348 to the Langmuir adsorption model with an R^2 value of 0.998 (Fig. S7). The apparent K_d was
349 estimated from the fitting to be 0.219 nM, which was almost the same as that in the absence of
350 serum. Moreover, the plot was fitted to a linear equation with an R^2 value of 0.995 in the anti-PEG
351 antibody concentration range of 0–10 ng mL⁻¹ (Fig. 5b). The LOD was estimated to be 0.10 ng
352 mL⁻¹, which is comparable to that in the absence of serum (0.065 ng mL⁻¹). Therefore, serum
353 components such as proteins hardly affect the binding of anti-PEG antibodies to OEG ligands on

354 cello-oligosaccharide assemblies. Notably, the detectable and quantifiable range of anti-PEG
355 antibody concentrations was estimated to be 0.10–10 ng mL⁻¹, which covered the range of the
356 concentrations required for clinical analysis of the effects of anti-PEG antibodies on the
357 pharmacokinetics and/or safety of PEGylated therapeutics.⁶ It appears that the antibiofouling
358 properties of these cello-oligosaccharide platforms allow for the quantitative detection of specific
359 anti-PEG antibodies even in the presence of serum.

360



361

362 **Fig. 5** (a) ELISAs of PEG-B-47 using 0.045% (w/v) CEL-EG₄ assembly at different PEG-B-47
363 concentrations in the presence of 10% fetal bovine serum. (b) Linear fitting in the concentration
364 range of 0–10 ng mL⁻¹.

365

366 The LOD of PEG-B-47 by the cello-oligosaccharide platforms was almost the same as that
367 (0.050 ng mL⁻¹) for conventional ELISAs using commercially available microplates covered with
368 PEGylated BSA, of which PEG had a molecular weight of 20 kDa and a methoxy terminus (Fig.
369 S8). Nevertheless, the present ELISAs using the antibiofouling cello-oligosaccharide platforms
370 required simple procedures, including centrifugation/redispersion cycles, without the need for
371 delicate techniques. Overall, it is suggested that the simple ELISAs using nanoarchitectonics-based
372 antibiofouling cello-oligosaccharide platforms presented here have great potential for practical
373 applications.

374

375 4. Conclusions

376 In conclusion, distinguishing anti-PEG antibodies in a simple manner was demonstrated by
377 using antibiofouling 2D cello-oligosaccharide assemblies tethering one-terminal methoxy OEG
378 ligands. The assemblies were well dispersed in PBS, and the nonspecific adsorption of proteins on
379 the assemblies was very low level, making our assays not only reliable but also simple. The anti-
380 PEG antibodies were successfully distinguished based on our ELISAs using specific OEG ligand-
381 tethering crystalline assemblies. Notably, the quantitative detection of a particular methoxy-
382 specific anti-PEG antibody was successfully performed even under practical conditions with
383 serum, which hardly interfere with quantification. Considering that the ELISAs presented here can
384 be simply performed by centrifugation/redispersion cycles without the need for delicate techniques
385 and that mammalian body fluids do not normally contain enzymes that degrade cello-
386 oligosaccharides (e.g., cellulases), the present study opens a new avenue for the analysis of the
387 binding characteristics of anti-PEG antibodies in biological samples. Specifically, immunogenicity
388 of PEGylated therapeutics under use and development can be investigated reliably and simply by
389 using cello-oligosaccharide assemblies, which potentially contribute to designing the improved
390 therapeutics with excellent therapeutic efficacy and clinical safety.

391

392 Declaration of competing interests

393 The authors declare no competing financial interests.

394

395 Data availability

396 Data will be made available on request.

397

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