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by amylose inclusion complex

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1 **Lipophilization and molecular encapsulation of *p*-coumaric acid by amylose**  
2 **inclusion complex**

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## 39 **ABSTRACT**

40 *p*-Coumaric acid (CA) is a natural phenolic compound with a wide range of  
41 bioactivities, but its tendency to degrade during food processing and storage limits its  
42 application in functional foods. Forming amylose-guest inclusion complex is a  
43 technique to molecularly encapsulate guest molecules of interest and protect them  
44 from adverse environmental challenges. However, raw CA cannot be encapsulated  
45 into the helical cavity of amylose so this study used a strategy of lipophilization of CA  
46 to synthesize hexadecyl *p*-coumarate (HC), which might complex with amylose. The  
47 formation of the amylose inclusion complex was studied by complementary  
48 techniques: differential scanning calorimetry (DSC), X-ray diffraction (XRD), and  
49 Fourier transform infrared (FTIR) spectroscopy. To test the photo-stability of CA in  
50 the complex, a direct UV irradiation was performed. The *in vitro* release behavior of  
51 CA in the complex was also studied in both simulated gastric and intestinal fluids.  
52 The results showed that HC, but not CA, could form inclusion complex with amylose.  
53 Compared with raw CA, HC, and amylose-HC physical mixture, HC complexed with  
54 amylose showed a significant improvement in photo-stability ( $P < 0.05$ ), with 90%  
55 retained, double the 43% retention of raw CA after 72 h of direct UV irradiation, and  
56 exhibited higher release rates of CA during *in vitro* digestion. These findings suggest  
57 that the amylose inclusion complex is a promising technique to protect CA in foods  
58 during processing and storage and to increase its bioaccessibility.

59 *Keywords:* *p*-Coumaric acid; Amylose; Lipophilization; Molecular encapsulation.

## 60 **1. Introduction**

61 *p*-Coumaric acid (CA), also known as trans-4-hydroxycinnamic acid, is a  
62 phytochemical present in cereals, fruits and vegetables such as wheat, apples and  
63 potatoes, which form a vital part of our daily diet (King & Young, 1999). Recent  
64 studies have reported that CA possesses a wide range of bioactivities, such as  
65 antioxidant (Cinzia, et al., 1995), anti-inflammatory (Pragasam, Venkatesan, & Rasool,  
66 2013), anti-apoptotic (Peng, et al., 2018), antiplatelet (Luceri, et al., 2007),  
67 anti-nephrotoxic (Navaneethan & Rasool, 2014) and cardioprotective activities  
68 (Prasanna, Krishnan, & Rasool, 2013). Therefore, CA has the potential to be  
69 formulated into functional foods and nutritional supplements to improve health and  
70 prevent chronic diseases. However, CA is not stable and its properties can be  
71 compromised by environmental factors, such as light and oxygen, under certain  
72 processing and storage conditions.

73 To protect bioactive compounds from degradation and increase their stability and  
74 bioavailability, several encapsulation methods have been developed. For example,  
75 soluble starch has been used to formulate nanocomposite, which could increase the  
76 water solubility and stability of curcumin (Li, Shin, Lee, Chen, & Park, 2016) and a  
77 catechin- $\beta$ -cyclodextrin inclusion complex has been fabricated to increase the storage  
78 stability of catechin (Ho, Thoo, Young, & Siow, 2017). The amylose inclusion  
79 complex is another effective encapsulation method, but has not yet been extensively  
80 studied, especially its ability to encapsulate and protect phenolic compounds.

81 Amylose is a linear polymer of glucose units connected by  $\alpha$ -(1, 4)-glycosidic  
82 bonds. The polysaccharide can crystallize into A- and B-type allomorphs that both

83 consist of 6-fold left-handed double helices which are packed in parallel (Le,  
84 Choisnard, Wouessidjewe, & Putaux, 2018). Amylose also exists as single helix when  
85 crystallizing with some linear organic molecules, such as lauric acid, palmitic acid  
86 (C16), and stearic acid (Kong, Lee, Kim, & Ziegler, 2014; Lesmes, Cohen, Shener, &  
87 Shimoni, 2009; Zhang, Huang, Luo, & Fu, 2012). When forming inclusion complex,  
88 amylose adopts a left-handed single helical structure with a hydrophobic helical cavity,  
89 where the linear compound can be accommodated through the hydrophobic force  
90 between them (Putseys, Lamberts, & Delcour, 2010). It has been suggested that the  
91 amylose molecule can protect the entrapped guest molecules against adverse  
92 environmental challenges and regulate their release (Lesmes, Barchechath, & Shimoni,  
93 2008; Lesmes, et al., 2009).

94 CA is relatively hydrophilic and adopts a bulk phenyl group, which cannot be  
95 entrapped by amylose directly. Therefore, to enable the amylose to encapsulate CA, a  
96 strategy of lipophilization of CA was conducted in the present study. CA was  
97 lipophilized by grafting on a C16 alkyl chain (provided by 1-hexadecanol) to obtain  
98 hexadecyl *p*-coumarate (HC). The <sup>1</sup>H-NMR spectrum and mass spectrum were used  
99 to confirm the synthesis of the target compound. The resultant samples were then  
100 examined by complementary techniques: differential scanning calorimetry (DSC),  
101 X-ray diffraction (XRD), and Fourier transform infrared (FTIR) spectroscopy, to  
102 verify the successful formation of the inclusion complex. The photo-stability and  
103 release behavior of CA in the inclusion complex were investigated to evaluate the  
104 effectiveness of this encapsulation technique.

## 105 **2. Materials and methods**

### 106 *2.1. Materials*

107 Potato amylose (free from amylopectin), *p*-coumaric acid, 1-hexadecanol,  
108 pancreatin from porcine pancreas (8 × United States Pharmacopeial Convention  
109 specifications), and pepsin (501 U/mg) were purchased from Sigma-Aldrich (St.  
110 Louis, MO, USA), triphenyl phosphine and diethylazodicarboxylate from TCI Co.  
111 (Shanghai, China), acetonitrile from J&K Scientific (Beijing, China), and  
112 tetrahydrofuran (THF), dimethyl sulfoxide (DMSO) and all other reagents of  
113 analytical grade from Sinopharm Chemical Reagent Co Ltd. (Beijing, China).

### 114 *2.2. Lipophilization of CA*

115 CA was lipophilized by the Mitsunobu reaction with some modifications  
116 (Katsumi, et al., 2009). 1-Hexadecanol (242 mg, 1 mmol), CA (328 mg, 1 mmol), and  
117 triphenylphosphine (524 mg, 2 mmol) were dissolved in THF (7 mL) by stirring. A  
118 solution of diethylazodicarboxylate (0.38 mL, 2 mmol) in THF (3 mL) was then  
119 added at 0 °C for 15 min then the mixture was stirred for 1 h. The reaction mixture  
120 was concentrated *in vacuo*, diluted with 25 mL ethyl acetate, washed with brine, dried,  
121 filtered then concentrated into solid form. The residue was then purified by silica gel  
122 column chromatography (hexane/ethyl acetate = 9/1) and concentrated to form a  
123 white solid powder. The mass and <sup>1</sup>H-NMR spectra were recorded to confirm that the  
124 synthesized product was HC (Fig. 1).

### 125 *2.3. Preparation of amylose inclusion complex*

126 The DMSO method was used to prepare the amylose inclusion complex with

127 slight modifications (Kong & Ziegler, 2014b; Ma, Floros, & Ziegler, 2011). Amylose  
128 (500 mg) was dissolved in 10 mL of 95% (v/v) aqueous DMSO by stirring at 90 °C  
129 for 30 min. Then 1 mL of pre-dissolved CA or HC (50 mg) solution in 95% (v/v)  
130 aqueous DMSO was mixed into the amylose dispersion. The mixture was vigorously  
131 stirred for 2 min using a vortex stirrer. The mixture was held at 90 °C for 30 min,  
132 diluted with 25 mL of deionized water at 90 °C then held for another 15 min before  
133 being allowed to cool to room temperature (20 °C) over at least 24 h. The precipitate  
134 was collected by centrifugation at  $3000 \times g$  for 10 min and washed 3 times with 50%  
135 (v/v) aqueous ethanol. The resulting pellet was transferred into an aluminum dish with  
136 a small amount of ethanol then allowed to dry at room temperature (20 °C) in a  
137 desiccator. The dried samples were pulverized into a fine powder for further analysis.

#### 138 *2.4. Differential scanning calorimetry*

139 The thermal properties of the samples were studied using a Discovery DSC 250  
140 differential scanning calorimeter equipped with an autosampler (TA Instruments, New  
141 Castle, DE, USA). Approximately 5 mg of each sample was weighed into a 60  $\mu$ L  
142 stainless steel pan and deionized water added to obtain a 10% (w/v) suspension. For  
143 the raw CA and HC samples, deionized water was added to obtain a 5% (w/v)  
144 solution/dispersion. The pan was hermetically sealed then equilibrated at room  
145 temperature (20 °C) for at least 2 h. The samples were analyzed by heating from 20  
146 °C to 150 °C at a rate of 5 °C/min. The data were analyzed using TA Trios software  
147 (TA Instruments).

#### 148 *2.5. X-ray diffraction analysis*

149 Wide angle X-ray diffraction patterns were obtained using a Bruker D8 Discover  
150 X-ray diffractometer (Bruker Corporation, Billerica, MA, USA). Dried sample  
151 powders were exposed to Co K $\alpha$  radiation (0.179 nm) then the data were converted to  
152 correspond to 2 $\theta$  values based on Cu K $\alpha$  radiation (0.154 nm).

### 153 *2.6. Fourier transform infrared spectroscopy*

154 Fourier transform infrared spectra of the samples were obtained using a  
155 PerkinElmer Spectrum 100 FTIR spectrometer (PerkinElmer Inc., Waltham, MA,  
156 USA) with an attenuated total reflection accessory. Spectra were recorded in the  
157 wavelength region between 650 and 4000 cm<sup>-1</sup>. Each spectrum was obtained by  
158 averaging 32 scans at a resolution of 4 cm<sup>-1</sup>.

### 159 *2.7. Quantitative determination of p-coumaric acid and hexadecyl p-coumarate by* 160 *High Performance Liquid Chromatography (HPLC)*

161 All samples were analyzed as described in a previous study (Moron,  
162 Pozo-Morales, Benito Mora, Garvi, & Lebrato, 2018). The assay was carried out  
163 using a Shimadzu LC-20AT HPLC instrument (Kyoto, Japan) and an Agilent C-18  
164 reverse phase column (250 mm  $\times$  4.6 mm  $\times$  5  $\mu$ m) with UV detection at 280 nm for  
165 CA and 300 nm for HC. The injection volume was 10  $\mu$ L, and an isocratic system  
166 using acetonitrile was used as the mobile phase at a flow rate of 1 mL/min. The  
167 calibration curve with CA and HC standard solutions ranging from 0 to 200  $\mu$ M in  
168 concentration was constructed to quantify the amount of CA and HC.

### 169 *2.8. Loading capacity determination*

170 The total CA content encapsulated in the inclusion complex was determined  
171 using solvent extraction followed by HPLC measurement. About 10 mg of the  
172 inclusion complex powder was dissolved in 15 mL DMSO then diluted with 35 mL  
173 acetonitrile. The solution was treated in an ultrasonic cleaner for 15 min to extract the  
174 encapsulated HC. The sample was measured by the HPLC method described above  
175 (2.7). The loading capacity was calculated by the following formula:

$$176 \quad \text{Loading capacity (\%)} = \frac{\text{Weight of HC}}{\text{Weight of amylose}} \times 100\%$$

### 177 2.9. Photo-stability analysis

178 The photochemical stability of the amylose-HC inclusion complex was estimated  
179 in comparison with CA, HC and amylose-HC physical mixture. The test was based on  
180 a previous study with some modifications (Li, et al., 2016). Samples (about 50 mg)  
181 were weighed into a small glass vial then subjected to UV irradiation (254 nm) from a  
182 UV lamp (16 W) at room temperature for 72 h in the dark. The distance between the  
183 samples and the UV lamp was 5 cm. After irradiation for 6, 12, 24, 48 and 72 h, the  
184 amylose-HC physical mixture and inclusion complex samples were dissolved in  
185 DMSO while the CA and HC samples were dissolved in acetonitrile to determine the  
186 retention of CA or HC using the HPLC method described above (2.7). The CA or HC  
187 content in each sample without UV treatment was regarded as 100% CA retention.

### 188 2.10. *In vitro* release of CA

189 The *in vitro* digestion was conducted in simulated gastric and intestinal fluids to  
190 compare the release profile of the inclusion complex with references, using methods  
191 in previous studies with some modifications (Jain, Jain, Gupta, & Ahirwar, 2007;

192 Tagliacruzchi, Verzelloni, Bertolini, & Conte, 2010). Samples containing the  
193 equivalent amount of 10 mg of CA were first suspended in 50 mL of simulated gastric  
194 fluid (SGF, in 1 L distilled water: NaCl, 2.0 g; 36% HCl, 7.0 mL; Tween 80, 1.0 mL;  
195 pepsin, 0.6 g. pH = 1.2), then the suspensions were incubated at 37 °C in a shaker  
196 (150 rpm). The simulated intestinal digestion was conducted under the same  
197 conditions after 120 min of digestion in SGF: 10 mL SGF suspensions were collected  
198 then added to an equivalent amount of simulated intestinal fluid (SIF, in 1 L distilled  
199 water: KH<sub>2</sub>PO<sub>4</sub>, 6.8 g; NaOH, 0.616 g; Tween 80, 1.0 mL) The pH value was then  
200 adjusted to 7.4 with 1 M NaOH before adding 1.0 g/L pancreatin and 3 g/L porcine  
201 bile salts. During the *in vitro* release test, aliquots (1 ml) were obtained after 15, 30,  
202 60, and 120 min of both the SGF and SIF digestions, then immediately centrifuged at  
203 3000 × g for 10 min (Paramera, Konteles, & Karathanos, 2011). The supernatants  
204 were collected and their CA content was measured using the HPLC method described  
205 above (2.7).

#### 206 2.11. Statistical analysis

207 All tests were done in triplicate. The data from the photo-stability analysis and *in*  
208 *vitro* release were expressed as mean and standard deviation (mean ± SD) then  
209 analyzed by two-way analysis of variance (ANOVA) with Tukey *post hoc* tests using  
210 IBM SPSS Statistics from Windows (Version 20.0; IBM Corp., Armonk, NY, USA).  
211 A difference between mean values with  $P < 0.05$  was considered as statistically  
212 significant.

### 213 3. Results and discussion

### 214 3.1. Amylose inclusion complex with CA

#### 215 3.1.1. Thermal properties

216 The DSC thermograms recorded from all the samples are shown in Fig. 2. Heating  
217 aqueous solutions of CA with and without amylose at over 100 °C probably resulted  
218 in its degradation, suggested by the endotherms ranging from 100-120 °C (Fig. 2a &  
219 2c). The sample of amylose-HC inclusion complex exhibited an endotherm with a  
220 peak temperature at  $96.72 \pm 0.37$  °C and an enthalpy of  $19.91 \pm 1.56$  J/g (Fig. 2f).  
221 This endotherm of the amylose-HC inclusion complex was not caused by the melting  
222 of HC, which was around 80-82 °C as observed in the thermograms of both pure HC  
223 and the amylose-HC physical mixture (Fig. 2b & 2d). The dissociation temperature of  
224 the amylose-HC inclusion complex was very close to those found in amylose  
225 inclusion complexes with other C-16 lipid compounds, e.g. palmitic acid (Kong &  
226 Ziegler, 2014a) and ascorbyl palmitate (Kong & Ziegler, 2014b), indicating that  
227 amylose had complexed with the C-16 alkyl chain of hexadecyl *p*-coumarate.  
228 However, the dissociation enthalpy, proportional to the amount of guest compounds  
229 that have been complexed, was much higher than that found in inclusion complexes of  
230 amylose-palmitic acid and starch-ascorbyl palmitate (Kong & Ziegler, 2014a; Ma, et  
231 al., 2011) and was close to that of the starch-alkylresorcinol inclusion complex  
232 (Gunenc, Kong, Elias, & Ziegler, 2018). This result was actually in agreement with  
233 the loading capacity, 6.8%. In contrast, the tentative amylose-CA sample produced no  
234 noticeable endotherm (Fig. 2e), indicating that amylose had not formed inclusion  
235 complex with CA.

### 236 3.1.2. X-ray diffraction analysis

237 The XRD patterns of raw CA and HC, and their inclusion complex samples are  
238 shown in Fig. 3. Both the raw CA and HC displayed sharp diffraction peaks, an  
239 indication of their highly crystalline nature. The 23.7 ° peak seen in the XRD pattern  
240 of HC has been observed as a major peak in many fatty acids and lipid molecules  
241 (Lesmes, et al., 2009; Tang & Copeland, 2007), reconfirming the successful  
242 lipophilization of CA. The absence of any characteristic peaks for CA and HC in the  
243 inclusion complex samples, and the absence of their melting endotherms on DSC  
244 thermograms, showed the effectiveness of the washing steps during the preparation  
245 procedure: the uncomplexed CA and HC had been washed away, leaving only those  
246 in the amylose single helices, if any. The tentative amylose-CA sample showed a very  
247 diffuse pattern with hints of a B type, i.e. weak peaks at 17 °, 19.5 °, and 22 ° (Kong,  
248 et al., 2014). The sample of amylose-HC inclusion complex, exhibiting peaks at 13 °  
249 and 20 °, proved to be characteristic of the V<sub>6h</sub> amylose XRD pattern seen in  
250 amylose-lipid inclusion complexes (Kong, Bhosale, & Ziegler, 2018). This suggested  
251 that HC had been entrapped intra-helically, instead of being precipitated into its only  
252 phase.

### 253 3.1.3. Fourier transform infrared investigation

254 The FTIR spectra provide additional evidence for the presence of guest compounds  
255 in the samples (Fig. 4). The FTIR spectrum of CA showed the characteristic phenolic  
256 compound bands at around 1500, 1600, 1670, and 3375 cm<sup>-1</sup>. The bands centered at  
257 around 1500 and 1600 cm<sup>-1</sup> were assigned to vibrations of the benzene ring; 3375

258  $\text{cm}^{-1}$  and  $1670 \text{ cm}^{-1}$  were respectively attributed to O–H stretching and the vibrations  
259 including C=C and C=O on CA (Ravikumar, Gaddamanugu, & Anand Solomon,  
260 2013; Swisłocka, Kowczyk-Sadowy, Kalinowska, & Lewandowski, 2012). Compared  
261 with CA, new bands appeared on the HC spectrum: bands at  $2850$  and  $2920 \text{ cm}^{-1}$ ,  
262 both of which were attributed to the stretching vibrations on  $-\text{CH}_2$  of the alkyl chains;  
263 bands at  $1710$  and  $1745 \text{ cm}^{-1}$ , both of which were assigned to carbonyl stretching; and  
264 a band at  $720 \text{ cm}^{-1}$  caused by the  $-\text{CH}_2$  rocking of the alkyl chain (Ciesik, Koll, &  
265 Grdadolnik, 2006), which jointly confirmed the structure of HC. On the spectrum of  
266 the sample of amylose-HC inclusion complex, the broad O–H vibration ranging from  
267  $3000$  to  $3600 \text{ cm}^{-1}$  was ascribed to the superposition of O–H bonds on the amylose  
268 molecules (Kizil, Irudayaraj, & Eetharaman, 2002). Minor bands of benzene ring  
269 vibration at around  $1600 \text{ cm}^{-1}$ , C=C and C=O on CA at  $1670 \text{ cm}^{-1}$ , and C–H vibration  
270 on the alkyl chain at  $2850$  and  $2920 \text{ cm}^{-1}$  could be observed, indicating the presence  
271 of HC in the inclusion complex. In contrast, the tentative amylose-CA sample showed  
272 no band characteristics of CA, confirming that CA could not complex with amylose  
273 and had been completely washed away.

### 274 *3.2. Photo-stability analysis*

275 UV irradiation is an effective and widely used method for evaluating  
276 photo-stability. The retention of CA, HC, the amylose-HC physical mixture and the  
277 amylose-HC inclusion complex are shown in Fig. 5. After 72 h of UV irradiation,  
278 about 57% of the CA was degraded in the raw CA group, which corresponded to a  
279 43% retention of CA. Lipophilization of CA improved the photo-stability of CA

280 slightly and resulted in a 57% retention after 72 h of UV irradiation. As expected,  
281 simply mixing amylose and HC into a physical mixture did not improve the  
282 photo-stability of HC. In contrast, the retention of CA (in the form of HC) was  
283 significantly higher when HC was encapsulated in the amylose inclusion complex ( $P$   
284  $< 0.05$ ). As much as 90% of the loaded HC was still retained in the amylose-HC  
285 inclusion complex after 72 h of direct UV irradiation. Therefore, the results showed  
286 that amylose inclusion complexation can increase the photo-stability of CA. UV is  
287 electromagnetic radiation with a wavelength between that of visible light and that of  
288 X-rays. Although it is not an ionizing radiation, it can cause the photodegradation of  
289 many compounds, in combination with the actions of oxygen and moisture. By  
290 molecularly encapsulating HC in the crystalline V-type amylose inclusion complex,  
291 its exposure to oxygen and moisture could be largely blocked, thus limiting the extent  
292 of its photodegradation by oxidative and hydrolysis reactions.

### 293 *3.3. In vitro release of CA*

294 The release profiles of CA in SGF and SIF are shown in Fig. 6. A large amount of  
295 CA was detected in the raw CA group within the first 30 min in SGF. In contrast, HC,  
296 the amylose-HC physical mixture and the amylose-HC inclusion complex released a  
297 limited amount of CA in SGF, but burst immediately after being exposed to SIF. This  
298 phenomenon may be ascribed to the absence of lipase and amylase in SGF so that CA  
299 could not be released from HC or the inclusion complex. Similar results have reported  
300 that vitamin D<sub>3</sub> could not be released from starch nanoparticles until pancreatic  
301 amylase had been added to the SIF (Hasanvand, Fathi, Bassiri, Javanmard, &

302 Abbaszadeh, 2015). For raw CA group, it was notable that the CA concentration  
303 decreased after 30 min release in SGF, which may have been caused by the  
304 degradation of CA under acidic conditions (Yang, et al., 2018). Moreover, it appeared  
305 that significantly more CA was released from the inclusion complex sample than from  
306 the references in SIF ( $P < 0.05$ ). These results implied that the amylose inclusion  
307 complex could protect CA from the acidic and enzymatic conditions of the stomach  
308 and thus increase the release of CA in the intestines, where the absorption of phenolic  
309 compounds occurs (Williamson & Clifford, 2017).

#### 310 **4. Conclusions**

311 In this study, an approach for encapsulating CA based on amylose molecules  
312 through lipophilization was investigated. The DSC, XRD, and FTIR results indicated  
313 that the amylose-HC inclusion complex had been formed, but not the amylose  
314 complexation with CA. After 72 h of direct UV irradiation, the CA in the inclusion  
315 complex sample was significantly better retained than that in the other samples. The  
316 release profiles showed that the inclusion complex exhibited higher release rates of  
317 CA during *in vitro* digestion. These results have demonstrated that the molecular  
318 encapsulation of lipophilized bioactive compounds by amylose is a promising  
319 technique for enhancing the stability and bioaccessibility of bioactive compounds,  
320 which could be applied during the production of fortified foods.

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325 **Conflict of interests**

326 The authors have declared no conflicts of interest.

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448 **Figure Captions**

449 Fig. 1. Chemical structure of hexadecyl *p*-coumarate.

450

451 Fig. 2. DSC thermograms of (a) *p*-coumaric acid (CA, 5%, w/v), (b) hexadecyl  
452 *p*-coumarate (HC, 5%, w/v), (c) amylose-CA physical mixture, (d) amylose-HC  
453 physical mixture, (e) tentative amylose-CA inclusion complex, and (f) amylose-HC  
454 inclusion complex.

455

456 Fig. 3. XRD patterns of (a) *p*-coumaric acid (CA), (b) hexadecyl *p*-coumarate (HC),  
457 (c) tentative amylose-CA inclusion complex, and (d) amylose-HC inclusion complex.

458

459 Fig. 4. FTIR spectra of (a) *p*-coumaric acid (CA), (b) hexadecyl *p*-coumarate (HC), (c)  
460 tentative amylose-CA inclusion complex, and (d) amylose-HC inclusion complex.

461

462 Fig. 5. Photo-stability of *p*-coumaric acid (CA), hexadecyl *p*-coumarate (HC),  
463 amylose-HC physical mixture, and amylose-HC inclusion complex. The letters a, b,  
464 and c indicate significant differences at each time point,  $P < 0.05$  ( $a > b > c$ ).

465

466 Fig. 6. The *in vitro* release of *p*-coumaric acid (CA) in simulated gastric and intestinal  
467 fluids from raw CA, hexadecyl *p*-coumarate (HC), amylose-HC physical mixture, and  
468 amylose-HC inclusion complex. The letters a, b, and c indicate significant differences  
469 at each time point,  $P < 0.05$  ( $a > b > c$ ).