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1 **The cadmium binding characteristics of a lactic acid**  
2 **bacterium in aqueous solutions and its application for**  
3 **removal of cadmium from fruit and vegetable juices**

4

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## 23 Abstract

24 Heavy metal cadmium (Cd) is an environmental pollutant that  
25 causes adverse health effects in humans. This toxic metal has been  
26 detected in a wide range of fruit and vegetables. A strain of lactic acid  
27 bacteria, *Lactobacillus plantarum* CCFM8610, was screened out for its  
28 good ability to bind Cd, and this study was designed to investigate the Cd  
29 binding properties of this bacterium, and to evaluate its use for removal  
30 Cd from fruit and vegetable juices. Electron microscopy observations and  
31 energy dispersive X-ray analysis confirmed that the majority of the Cd  
32 was bound to the surface of the bacterial cell. The Cd biosorption of *L.*  
33 *plantarum* CCFM8610 was strongly pH dependent, and carboxyl and  
34 amino groups of the bacterial surface molecules are important in the  
35 binding process. The biosorption was fast and efficient, and could be well  
36 explained by the Langmuir-Freundlich dual isotherm model ( $R^2 = 0.99$ )  
37 and the pseudo second-order kinetic model ( $R^2 = 0.99$ ). After a 2 h  
38 incubation and a simple centrifugation, *L. plantarum* CCFM8610  
39 treatment removed 67% to 82% of the Cd from nine types of fruit and  
40 vegetable juices. Long-period fermentation by *L. plantarum* CCFM8610  
41 (36 h) also significantly decreased Cd concentrations in the juices (56%  
42 to 81%). Our results show that this food-grade bacterial strain could be  
43 used as a potential probiotic for Cd removal from fruit and vegetable  
44 juices.

45 **Keywords:** Lactic acid bacteria; *Lactobacillus plantarum*; Cadmium  
46 removal; Fruit and vegetable juices  
47

## 48 **1. Introduction**

49 Heavy metal cadmium (Cd) is a ubiquitous environmental pollutant  
50 that can contribute to a broad range of adverse health effects in humans.  
51 In recent years, the risk of human exposure to Cd has increased with the  
52 increasing contamination of the food chain and inadequate protection  
53 from Cd-contaminated food <sup>1</sup>. Due to its high rate of soil-to-plant transfer,  
54 Cd accumulation has been observed in a variety of fruit and vegetables  
55 such as apples, strawberries, pears, tomatoes, and cucumbers <sup>2-4</sup>. Even if  
56 the Cd levels in fruit and vegetables are normally low, a high  
57 consumption of fruit and vegetables and their juices can result in an  
58 accumulative contribution to Cd exposure <sup>5</sup>. The Joint FAO/WHO Expert  
59 Committee on Food Additives has set a tolerable weekly intake (TWI) of  
60 Cd at 7  $\mu\text{g kg}^{-1}$  body weight <sup>6</sup>. The Integrated Risk Information System  
61 (IRIS) of United States Environmental Protection Agency (EPA) set the  
62 threshold of oral Cd dose at 1  $\mu\text{g}$  per kg per day <sup>7</sup>. To date, no specific  
63 methods for Cd removal from fruit and vegetable juices have been  
64 developed and chemical processes, such as chemical deposition and ion  
65 exchange, have efficacy and safety issues <sup>8,9</sup>.

66 Several studies have revealed that some lactic acid bacteria (LAB),  
67 including *Lactobacillus rhamnosus*, *L. plantarum*, and *L. reuteri*, are  
68 capable of binding and removing heavy metals such as Cd and lead *in*  
69 *vitro* <sup>10, 11</sup>. Oral administration of *L. rhamnosus* can also prevent the

70 increases in mercury and arsenic blood levels in pregnant women  
71 suspected to have high toxic metal exposures<sup>12</sup>. In our previous study, a  
72 specific lactic acid bacterium, *L. plantarum* CCFM8610, was identified  
73 for its excellent Cd removing ability<sup>13,14</sup>. *In vivo* studies also confirmed  
74 that this strain can sequester Cd in the intestines, thus decreasing the Cd  
75 absorption of mice subjected to chronic dietary Cd exposure<sup>15</sup>. However,  
76 the mechanism of the efficient Cd binding process of this strain have not  
77 yet to be elucidated.

78 *L. plantarum* strains are widely used in the food industry and are  
79 generally regarded as safe<sup>16</sup>. A considerable number of studies have  
80 shown that *L. plantarum* strains can be used for lactic acid fermentation  
81 and to enhance the nutritional, sensory and shelf life properties of fruit  
82 and vegetables<sup>17-19</sup>. However, to our knowledge, no studies have been  
83 carried out so far to evaluate the effects of LAB strains on heavy metal  
84 removal from foodstuffs, such as fruit and vegetable juices.

85 The objective of this study was to investigate the Cd binding  
86 properties of *L. plantarum* CCFM8610 and to assess the potential  
87 application of this LAB strain for removal of Cd from fruit and vegetable  
88 juices.

## 89 **2. Materials and methods**

### 90 *2.1. Bacterial strains and culture*

91 The *L. plantarum* CCFM8610 strain was obtained from the in-house  
92 Culture Collections of Food Microbiology (CCFM), Jiangnan University  
93 (Wuxi, China). It was cultured in de Man, Rogosa and Sharpe (MRS)  
94 broth (Hopebio Company, Qingdao, China) at 37°C for 18 h. All of the  
95 bacteria were subcultured twice before the experiment.

## 96 2.2 *Cd binding assay*

97 The Cd binding experiment was carried out as previously described  
98 <sup>13</sup>. Briefly, *L. plantarum* CCFM8610 cell pellets were suspended in  
99 ultrapure water containing 50 mg/L of Cd as cadmium chloride, to give a  
100 final bacterial density of 1 g/L (wet weight). The cell pellets were  
101 resuspended in sterile saline at an identical bacterial concentration, as an  
102 untreated control. The bacterial suspension samples were then incubated  
103 for 1 h at 37°C, with the pH adjusted to 6.0. The suspension was then  
104 centrifuged at 8000 × *g* for 20 min and a sample was taken from the  
105 supernatant for analysis of the residual Cd concentration by flame atomic  
106 absorption spectrophotometry (Spectr AA 220; Varian). The Cd binding  
107 ability of the strains was expressed as the percentage of Cd removed  
108 (bound by the bacteria) as follows,

$$109 \text{ Removal (\%)} = 100\% \times [(C_0 - C_1)/C_0],$$

110 where  $C_0$  and  $C_1$  are the initial Cd concentration and the residual Cd  
111 concentration after removal, respectively.

## 112 2.2. *Electron microscopy and energy dispersive X-ray analysis*

113 The samples for transmission electron microscopy (TEM)  
114 observation were prepared as previously described<sup>20</sup>. After the Cd  
115 binding assay, bacterial cells were harvested by centrifugation at  $8000 \times g$   
116 for 20 min, washed with phosphate buffer solution (PBS, pH 7.2), and  
117 resuspended in 1 ml of PBS. One hundred microliters of 25%  
118 glutaraldehyde was added to the bacterial suspension and the cells were  
119 left to fix for 1.5 h. The suspensions were centrifuged and washed three  
120 times in sodium cacodylate buffer (0.05 M). The cell pellets were then  
121 mixed at a 1:1 ration with molten 2% low melting point agarose, which  
122 was solidified by chilling and then chopped into small pieces  
123 (approximately  $1 \text{ mm}^3$ ). These samples were left overnight in a 2.5%  
124 glutaraldehyde/0.05 M sodium cacodylate buffer (pH 7.2) and then  
125 transferred to a Leica EM TP tissue processor (Leica Microsystems UK  
126 Ltd., Milton Keynes) where they were washed, post-fixed and dehydrated  
127 with 1 h between each change. The samples were then infiltrated with a  
128 resin (London Resin Company Ltd.) and ethanol mixture, and the tissue  
129 blocks of the samples were placed into gelatine capsules containing fresh  
130 resin and polymerized overnight at  $60^\circ\text{C}$ . The sections (approximately 90  
131 nm thick) were cut using an ultramicrotome (Ultracut E, Reichert-Jung),  
132 and collected on film/carbon-coated copper grids. The sections were  
133 examined and imaged using a FEI Tecnai G2 20 Twin TEM at 200 kV.

134 The samples for scanning electron microscopy (SEM) observation

135 were prepared as previously described, with minor modifications<sup>21</sup>. After  
136 the Cd binding assay, the bacterial cells were harvested by centrifugation  
137 at  $8000 \times g$  for 20 min, washed with PBS (pH 7.2), and fixed with  
138 glutaraldehyde (2.5% v/v) for 4 h. The cells were then washed with PBS  
139 (pH 7.2) three times and dehydrated with graded alcohols. An identical  
140 volume of isoamyl acetate was used to displace the graded alcohols. The  
141 bacterial samples were lyophilized and a Hitachi S-3400N SEM was used  
142 to observe the cellular morphology.

143 The TEM and SEM were equipped with energy dispersive X-ray  
144 (EDX; Edax) microanalysis systems and the elemental composition of the  
145 selected areas observed using the electron microscope was analyzed.

### 146 *2.3. Characterization of the cellular components involved in Cd binding*

147 The Cd binding abilities of the different cellular components of *L.*  
148 *plantarum* CCFM8610 were determined as previously described, with  
149 minor modifications<sup>22,23</sup>. After the Cd binding assay with an initial pH of  
150 6.0, the Cd bound to exopolysaccharides, the external surface of the cell  
151 wall, the interior surface of the cell wall and the external surface of the  
152 cell membrane, and the interior of the protoplast was eluted separately.  
153 The samples were transferred to metal-free digestion vessels (Omni;  
154 CEM) and digested in concentrated HNO<sub>3</sub> using the Microwave  
155 Digestion System (MARS; CEM). The Cd concentration of each cellular

156 component was detected by flame atomic absorption spectrophotometry  
157 and the Cd binding ability was expressed as follows,

158 
$$\text{Cd bound by each cellular component (\%)} = C_1 / C_0 \times 100\%$$
,

159 where  $C_0$  and  $C_1$  are the Cd concentrations of the intact cell and the cell  
160 component, respectively.

#### 161 *2.4. Characterization of the functional groups involved in Cd binding*

162 The bacterial cells were harvested and subjected to chemical  
163 treatments to modify the functional groups<sup>24-26</sup>. Briefly, the carboxyl  
164 groups were neutralized in methanol with HCl (0.1 M), the amino groups  
165 were methylated in formaldehyde and formic acid solution (1:2 v/v),  
166 while the phosphoryl groups were esterified in triethyl phosphite and  
167 nitromethane solution (4:3 v/v). The treated samples were subjected to  
168 the Cd binding assay as described above, with an initial bacterial density  
169 of 10 g/L (dry weight) and an initial pH of 6.0.

170 The effect of the initial pH (2.0-7.0) on the Cd binding of *L.*  
171 *plantarum* CCFM8610 was tested using the same method of Cd binding  
172 assay described above, but with an initial Cd concentration of 5 mg/L.

#### 173 *2.5. Cd binding isotherm study*

174 According to previously reported methods<sup>27</sup>, the harvested cells  
175 were suspended in ultrapure water containing 0.06 to 90 mg/L Cd as

176 cadmium chloride, to give a final bacterial concentration of 1 g/L (dry  
177 weight). The Cd binding assay was then conducted with an initial pH of  
178 6.0 and the equilibrium content of Cd bound by the bacterium was  
179 expressed as follows,

$$180 \quad q_e (\text{Cd content bound by the biomass}) = [(C_i - C_e) \times V] / m,$$

181 where  $C_i$  and  $C_e$  are the initial Cd concentration and the equilibrium Cd  
182 concentration, respectively, and  $V/m = 1 \text{ L/g}$ .

183 The acquired experimental data were fitted to different isotherm  
184 models including the Langmuir<sup>28</sup>, the Freundlich<sup>28</sup>, and the  
185 Langmuir-Freundlich dual models<sup>29</sup>.

### 186 *2.6. Cd binding kinetic study*

187 In accordance with previously reported methods<sup>27</sup>, the harvested  
188 cells were suspended in ultrapure water containing 30 mg/L of Cd as  
189 cadmium chloride, to give a final bacterial concentration of 1 g/L (dry  
190 weight). The Cd binding assay was then conducted with an initial pH of  
191 6.0 and the concentration of Cd in the supernatant was detected at  
192 different time intervals up to 360 min. The acquired experimental data  
193 were fitted to different kinetic models (Table 1).

### 194 *2.7. Cd removal from fruit and vegetable juices*

195 Fresh apples, tomatoes, and cucumbers were collected from three

196 local markets in Jiangsu, Jiangxi, and Hunan Province in China,  
197 respectively. These fruit and vegetables were washed, chopped, and  
198 added to ultrapure water (1:2 w/v). Juice was obtained by pressing the  
199 mixture in a juice extractor. The juice was then centrifuged at  $10000 \times g$   
200 for 20 min, sterilized at  $105 \text{ }^\circ\text{C}$  for 10 min, and glass-bottled. The juices  
201 were labeled as AJ1, AJ2, AJ3 (apple juices), TJ1, TJ2, TJ3 (tomato  
202 juices), CJ1, CJ2, CJ3 (cucumber juices).

203 The viable *L. plantarum* CCFM8610 cells were harvested, washed  
204 with sterilized saline, and used to inoculate the fruit and vegetable juices  
205 at an inoculum level of 4% (v/v), corresponding to ca.  $10^7$  cfu/mL<sup>17,30</sup>.  
206 The juice without the bacterial inoculum was used as a control. All of the  
207 juices were incubated at  $37 \text{ }^\circ\text{C}$  for 2 h or 36 h. After the incubation, the  
208 bacterial growth was measured by colony counting. The juices were then  
209 centrifuged at  $8000 \times g$  for 20 min to remove the Cd-bound bacterial  
210 biomass and the supernatant was collected for biochemical analysis.

211 The antioxidative activities of the juices were measured using a  
212 DPPH (1,1-diphenyl-2-picrylhydrazyl) scavenging assay<sup>31</sup>. The juices  
213 were digested in concentrated  $\text{HNO}_3$  with the Microwave Digestion  
214 System, and the Cd concentration was measured by flame atomic  
215 absorption spectrophotometry.

216 *2.8. Statistical analysis*

217 Data are expressed as the mean  $\pm$  the standard error of the mean  
218 (SEM). A minimum of three independent experiments were carried out  
219 for each assay. The differences between the means of the test were  
220 analyzed using one-way analysis of variance, followed by Tukey's post  
221 hoc test. A  $p$  value of  $<0.05$  was considered to be statistically significant.  
222 The statistical analysis of the data obtained was performed using SPSS  
223 16.0 for Windows software. For the isotherm and kinetic studies, all of  
224 the equation parameters were evaluated using MATLAB R2010b  
225 software and a correlation coefficient ( $R^2$ ) test was used to measure the  
226 fitness of the data to the models.

### 227 **3. Results**

#### 228 *3.1. Electron microscopy observation and EDX analysis*

229 TEM micrographs of *L. plantarum* CCFM8610 before and after the  
230 Cd binding are presented in Fig. 1. Obvious deposits of Cd were observed  
231 on the surface of the cells after binding (Fig. 1B), whereas no Cd was  
232 visible in the micrographs of untreated cells (Fig. 1A). For the EDX  
233 analysis, no Cd signal could be detected in the control sample (Fig. 1C),  
234 but a clear peak for Cd was observed in Cd treated cells (Fig. 1D),  
235 indicating the presence of Cd due to biosorption.

236 The SEM micrographs revealed that Cd exposure caused anomalous  
237 aggregation of the *L. plantarum* CCFM8610 cells (Fig. 2B), whereas no

238 morphological changes in the control biomass were observed (Fig. 2A).  
239 The EDX spectra also confirmed an additional Cd peak of Cd-treated  
240 cells (Fig. 2D), which was absent in the control sample (Fig. 2C).

### 241 3.2. Cd binding abilities of different cell components

242 The exopolysaccharides fraction of *L. plantarum* CCFM8610 did not  
243 bind Cd (Fig. 3). The greatest amount of Cd accumulated on the external  
244 surface of the cell wall ( $40.54\% \pm 0.80\%$ ) and in the space between the  
245 cell wall and the plasma membrane ( $54.30\% \pm 2.93\%$ ). Only  
246 approximately 7% of the Cd entered the protoplast.

### 247 3.3. Effects of functional groups on Cd binding

248 When the amino groups of *L. plantarum* CCFM8610 cell surface  
249 molecules were methylated, the bacterial cells' ability to bind Cd was  
250 hampered and the removal of Cd decreased from 48.16% to 5.95% (Fig.  
251 4). There was also a decrease in Cd removal when carboxyl groups were  
252 neutralized ( $p < 0.05$ ). In contrast, the chemical modification of  
253 phosphoryl groups did not cause a significant difference in Cd removal  
254 ( $p > 0.05$ ).

255 The initial pH was critical to the Cd binding ability of *L. plantarum*  
256 CCFM8610 (Fig. 5). The Cd removal remained negligible when the pH  
257 was below 2.0, but increased significantly with the incremental rise in pH.

258 The results for pH values over 7.0 were excluded from the analysis  
259 because  $\text{Cd}^{2+}$  was found to precipitate under high pH conditions.

### 260 3.4. *Cd binding isotherm*

261 As shown in Fig. 6, the removal of Cd by *L. plantarum* CCFM8610  
262 was evaluated by plotting the amount of Cd bound by the bacteria ( $q_e$ )  
263 against the equilibrium Cd concentration of the metal ( $C_e$ ). The Cd  
264 binding ability increased with an increase of Cd concentration in the  
265 solution, ultimately reaching the equilibrium value. Data were further  
266 analyzed using different isotherm models (Table 2). Considering the  
267 highest  $R^2$  value, the Cd biosorption by the strain showed best fit to the  
268 Langmuir-Freundlich dual isotherm model ( $R^2 = 0.9928$ ).

### 269 3.5. *Cd binding kinetics*

270 As indicated in Fig. 7, the Cd binding process of *L. plantarum*  
271 CCFM8610 was efficient and fast. Approximately 90% of the binding  
272 process was completed in less than 100 min, and the saturation value was  
273 reached by about 300 min. Data were further analyzed by the pseudo first  
274 and second-order rate models (Table 1), with the latter model showing  
275 better fitness ( $R^2 = 0.9954$ ) than the former ( $R^2 = 0.9749$ ).

### 276 3.6. *Removal of Cd from fruit and vegetable juices by L. plantarum*

277 As shown in Table 3, a significant decrease in the pH value was  
278 observed in all of the juices after *L. plantarum* CCFM8610 treatment.  
279 The 36 h-fermentation caused more significant changes in the pH than the  
280 2 h treatment but the extent of reduction was also dependent on the nature  
281 of the juice. There was no statistically significant difference in the viable  
282 bacterial cell numbers between the 0 h and 2 h timepoints ( $p > 0.05$ ). For  
283 the 36 h treatment, the viable count of the strain reached over  $1 \times 10^8$   
284 cfu/mL in each juice (except for CJ3, with a count of  $0.95 \times 10^8$  cfu/mL),  
285 which is significantly higher than that at 0 h and 2 h timepoints.

286 The effects of *L. plantarum* CCFM8610 treatment on Cd removal  
287 from juices are shown in Tables 4 and 5 at 2 h and 36 h time points  
288 respectively. The presence of Cd was observed in all juices collected from  
289 different regions of China. Compared with the control groups, the  
290 treatment of *L. plantarum* CCFM8610 significantly decreased the Cd  
291 levels in each juice ( $p < 0.05$ ). After 2 h, the minimum Cd removal  
292 (67.12%) was found in the tomato juice and the maximum removal  
293 (82.87%) was observed in the cucumber juice (Table 4). For the 36 h  
294 treatment, Cd removal rate varied between 56.18% and 81.79%  
295 depending on the type of juice (Table 5).

296 It was also noticed that the 36 h treatment significantly increased the  
297 free radical scavenging ability of most of the juices (Table 6), whereas the  
298 2 h treatment only increased such ability in only two of the tomato juices.

## 299 4. Discussion

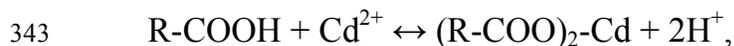
300 As a consequence of environmental Cd contamination, this  
301 hazardous heavy metal has become a threat to food safety. In China, Cd  
302 has been detected in a wide range of fruit and vegetables<sup>32-34</sup>. Our data on  
303 the levels of Cd in nine fruit and vegetables collected from different  
304 regions in China (Tables 4 and 5) further support this finding. As Cd is a  
305 non-essential but extremely toxic metal for humans, the concentrations of  
306 Cd in fruit and vegetables should be controlled to a minimum. The  
307 present study confirmed that a lactic acid bacteria strain, *L. plantarum*  
308 CCFM8610, can be used effectively to decrease the Cd levels in fruit and  
309 vegetable juices. This represents the first study identifying the use of a  
310 food-grade microorganism to reduce Cd levels in foodstuffs. After  
311 exposure of juices to *L. plantarum* CCFM8610, it was possible to remove  
312 over 56% of the Cd in the juices, which can help to keep the oral intake  
313 of Cd of consumers under the limits set by FAO/WHO or EPA<sup>6,7</sup>.

314 The Cd binding characteristics of *L. plantarum* CCFM8610 were  
315 investigated first. Compared with the control samples, additional Cd  
316 signals were observed in the EDX spectra after Cd exposure, confirming  
317 the presence of Cd in the cell biomass due to biosorption (Figures 1 and  
318 2). The morphological alterations of the strain after Cd binding, observed  
319 in the SEM micrographs, may be a result of the change in surface charge  
320 and the degeneration of the surface proteins caused by Cd exposure,

321 leading to the anomalous aggregation and enhanced Cd binding ability of  
322 the cells. A previous study reported similar phenomenon in an  
323 *Acidiphilium symbioticum* strain after Cd binding and indicated such a  
324 mechanism as a form of self-protection by the bacterium <sup>27</sup>. The TEM  
325 micrographs showed that the majority of the Cd passes through the  
326 surface polysaccharides of the strain and deposits on the surface of the  
327 cell, which is consistent with the differential Cd binding capacities of  
328 specific cell components (Fig. 3). It was noted that the deposition of Cd is  
329 discontinuous, indicating the involvement of specific biosorption sites on  
330 the cell surface. Similar deposition of Cd on the perimeter of the cell was  
331 also observed in a sulfate-reducing bacterium <sup>35</sup>.

332 To evaluate the possible functional groups involved in Cd binding,  
333 chemical modification was used to block the carboxyl, amino and  
334 phosphoryl groups on the surface structures of *L. plantarum* CCFM8610.  
335 The chemical modifications of the former two groups reduced the Cd  
336 binding ability of the strain by 18% and 42%, respectively, indicating that  
337 these two functional groups are important in the Cd binding process (Fig.  
338 4). These results are consistent with previous studies on the effects of  
339 functional groups of *A. symbioticum* and *Bacillus subtilis* on heavy metal  
340 binding and the possible ion exchange mechanisms involved could be as  
341 follows <sup>27, 36</sup>,





344 where R represents other molecular components on the cell surface and X  
345 represents compounds that can complex with  $\text{Cd}^{2+}$ .

346 As the electronegative carboxyl groups are abundantly available on  
347 the cell surface, they actively participate in the binding process of  $\text{Cd}^{2+}$   
348 with simultaneous release of protons<sup>27</sup>. The nitrogen atom of the amino  
349 groups can bind to Cd following electron pair sharing. Nitrogen donates a  
350 lone pair of electrons in the process of Cd binding, which facilitates the  
351 formation of stable metal complex<sup>27,37</sup>. Previous research has shown that  
352 Cd removal by *L. fermentum* decreased significantly after blocking of the  
353 phosphoryl groups<sup>25</sup>. However, the chemical modification of the  
354 phosphoryl groups in this study did not cause significant changes in Cd  
355 removal by *L. plantarum* CCFM8610. The Cd biosorption of *L.*  
356 *plantarum* CCFM8610 was strongly pH-dependent and the optimal pH  
357 for Cd binding was between 6.0 and 7.0 (Fig. 5). Similar results have  
358 been reported for other lactic acid bacteria strains, indicating that ion  
359 exchange is at least partly responsible for the binding process<sup>38,39</sup>.

360 The isotherm and kinetic models were established to further  
361 understand the binding characteristics of *L. plantarum* CCFM8610  
362 (Tables 1 and 2). The isotherm experimental data of Cd binding fit the  
363 Langmuir model well ( $R^2 = 0.97$ ), and the theoretical monolayer Cd  
364 biosorption capacity ( $Q_{max}$ ) of the strain was calculated as 24.69 mg/g

365 (dry biomass), which is higher than previously reported commercial LAB  
366 strains such as *L. casei* Shirota (12.1 mg/g) and *L. rhamnosus* GG (13.2  
367 mg/g)<sup>38</sup>. Similar differences in Cd binding ability between LAB strains,  
368 including *L. plantarum* CCFM8610 and *L. rhamnosus* GG, were also  
369 identified in our previous study<sup>13</sup>. Considering the highest  $R^2$  value, the  
370 biosorption showed the best fit to the Langmuir-Freundlich dual model  
371 ( $R^2 = 0.99$ ). This is in accordance with the heavy metal binding  
372 characterization of other bacteria, indicating the contribution of both  
373 physical and chemical binding mechanisms<sup>27, 40</sup>. The kinetic analysis  
374 suggests a rapid binding process that best fits the pseudo second-order  
375 rate model, which is consistent with previous reports<sup>27, 38</sup>. The proper  
376 correlation with pseudo second-order rate model also indicates the  
377 involvement of chemical absorption during the binding process<sup>41, 42</sup>.

378 The potency of this bacterium as a food-grade Cd absorbent was  
379 examined using 9 different Cd containing fruit and vegetable juices  
380 obtained from different regions of China. After exposure of these juices to  
381 *L. plantarum* CCFM8610 for only 2 h followed by a simple  
382 centrifugation to remove the Cd-bound bacterial biomass, it was possible  
383 to remove between 67% to 82% of the Cd present in the juices. The strain  
384 was also tested as a starter culture and the juices were fermented for 36 h.  
385 This treatment also significantly decreased Cd concentration of the juices  
386 (56% to 81%). The dramatic decrease in the pH of the juices after the 36

387 h incubation (Table 3) could have caused negative effects on the binding  
388 ability of the bacterium. However, as the viable cell number significantly  
389 increased after the 36 h incubation, more biomass was involved in Cd  
390 binding than in the 2 h treatment, enhancing the Cd removal and  
391 offsetting the reduced efficiency due to lower pH. The 36 h incubation  
392 also improved the antioxidant ability of the juices compared with the 2 h  
393 incubation (Table 6), which could be due to the increase of total  
394 flavanones in the juices after a long fermentation by this bacterium<sup>31,43</sup>.  
395 As oxidative stress has been reported to be an important toxic mechanism  
396 of Cd exposure<sup>44</sup>, fermentation could enhance the antioxidant ability of  
397 juices against Cd-induced toxicity. Therefore, although such fermentation  
398 processes will add the cost of juice production, the improved safety by Cd  
399 reduction and enhanced antioxidant properties of the vegetable and fruit  
400 juices may make such applications still worthwhile. A further study to  
401 investigate the organoleptic properties of these probiotic-treated juices is  
402 now in progress, to fully evaluate the consumer's acceptance of such  
403 products.

## 404 **5. Conclusions**

405 The Cd binding characteristics of a food-grade microorganism, *L.*  
406 *plantarum* CCFM8610, and the application of this strain to Cd removal  
407 from fruit and vegetable juices were evaluated in this study. The Cd

408 biosorption by *L. plantarum* CCFM8610 was a fast, efficient, and  
409 pH-dependent process, which followed the Langmuir-Freundlich dual  
410 isotherm model and showed the best fit to the pseudo second-order rate  
411 kinetic model. The *L. plantarum* CCFM8610 treatment (both 2 h and 36 h)  
412 significantly decreased the Cd concentration in nine types of juices. Our  
413 results show that this strain could be used as a potential probiotic for Cd  
414 removal from fruit and vegetable juices.

### 415 **Acknowledgements**

416 This work was supported by the National Natural Science  
417 Foundation of China Key Program (No. 31530056), the National Natural  
418 Science Foundation of China (No. 31470161, 31371721), the BBSRC  
419 Newton Fund Joint Centre Award, the 111 Project B07019, and the  
420 Program for Changjiang Scholars and Innovative Research Team in  
421 University (IRT1249).

### 422 **References**

- 423 1. G. F. Nordberg, K. Nogawa, M. Nordberg and L. Friberg, in *Handbook on the*  
424 *Toxicology of Metals*, eds. G. F. Nordberg, B. A. Fowler, M. Nordberg and L.  
425 Friberg, Academic Press, Burlington, MA, 3rd edn., 2011, ch. 23, pp. 446-451.
- 426 2. F. L. Coco, P. Monotti, F. Cozzi and G. Adami, *Food Control*, 2006, **17**, 966-970.
- 427 3. O. Muñoz, J. M. Bastias, M. Araya, A. Morales, C. Orellana, R. Rebolledo and D.

- 428 Velez, *Food Chem. Toxicol.*, 2005, **43**, 1647-1655.
- 429 4. Z. Krejpcio, S. Sionkowski and J. Bartela, *Pol. J. Environ. Stud.*, 2005, **14**,  
430 877-881.
- 431 5. R. Tahvonen, *Food Addit. Contam.*, 1998, **15**, 446-450.
- 432 6. FAO/WHO, Evaluation of certain food contaminants: 64<sup>th</sup> report of the Joint  
433 FAO/WHO Expert Committee on Food Additives, WHO Technical Report Series  
434 930, Geneva, 2006,  
435 [http://apps.who.int/iris/bitstream/10665/43258/1/WHO\\_TRS\\_930\\_eng.pdf](http://apps.who.int/iris/bitstream/10665/43258/1/WHO_TRS_930_eng.pdf),  
436 accessed December, 2015.
- 437 7. Integrated Risk Information System of United States Environmental Protection  
438 Agency,  
439 [http://cfpub.epa.gov/ncea/iris/iris\\_documents/documents/subst/0141\\_summary.p](http://cfpub.epa.gov/ncea/iris/iris_documents/documents/subst/0141_summary.pdf)  
440 [df](http://cfpub.epa.gov/ncea/iris/iris_documents/documents/subst/0141_summary.pdf), accessed November, 2014.
- 441 8. E. Maliou, M. Malamis and P. Sakellarides, *Water Sci. Technol.*, 1992, **25**,  
442 133-138.
- 443 9. W. W. Ngah and M. Hanafiah, *Bioresour. Technol.*, 2008, **99**, 3935-3948.
- 444 10. T. Halttunen, M. Collado, H. El-Nezami, J. Meriluoto and S. Salminen, *Lett.*  
445 *Appl. Microbiol.*, 2008, **46**, 160-165.
- 446 11. J. Bhakta, Y. Munekage, K. Ohnishi and B. Jana, *Int. J. Environ. Sci. Technol.*,  
447 2012, **9**, 433-440.
- 448 12. J. E. Bisanz, M. K. Enos, J. R. Mwanga, J. Changalucha, J. P. Burton, G. B.  
449 Gloor and G. Reid, *MBio*, 2014, **5**, e01580-01514.

- 450 13. Q. Zhai, R. Yin, L. Yu, G. Wang, F. Tian, R. Yu, J. Zhao, X. Liu, Y. Q. Chen and  
451 H. Zhang, *Food Control*, 2015, **54**, 23-30.
- 452 14. Q. Zhai, G. Wang, J. Zhao, X. Liu, F. Tian, H. Zhang and W. Chen, *Appl.*  
453 *Environ. Microbiol.*, 2013, **79**, 1508-1515.
- 454 15. Q. Zhai, G. Wang, J. Zhao, X. Liu, A. Narbad, Y. Q. Chen, H. Zhang, F. Tian and  
455 W. Chen, *Appl. Environ. Microbiol.*, 2014, **80**, 4063-4071.
- 456 16. B. Foligné, C. Daniel and B. Pot, *Curr. Opin. Microbiol.*, 2013, **16**, 284-292.
- 457 17. P. Filannino, L. Azzi, I. Cavoski, O. Vincentini, C. G. Rizzello, M. Gobbetti and  
458 R. Di Cagno, *Int. J. Food Microbiol.*, 2013, **163**, 184-192.
- 459 18. R. Di Cagno, R. Coda, M. De Angelis and M. Gobbetti, *Food Microbiol.*, 2013,  
460 **33**, 1-10.
- 461 19. P. Filannino, Y. Bai, R. Di Cagno, M. Gobbetti and M. G. Gänzle, *Food*  
462 *Microbiol.*, 2015, **46**, 272-279.
- 463 20. E. Dertli, I. J. Colquhoun, A. P. Gunning, R. J. Bongaerts, G. Le Gall, B. B.  
464 Bonev, M. J. Mayer and A. Narbad, *J. Biol. Chem.*, 2013, **288**, 31938-31951.
- 465 21. R. Chakravarty and P. C. Banerjee, *Extremophiles*, 2008, **12**, 279-284.
- 466 22. A. M. Massadeh, F. A. Al-Momani and H. I. Haddad, *Biol. Trace Elem. Res.*,  
467 2005, **108**, 259-269.
- 468 23. S. Deb, S. Ahmed and M. Basu, *B. Environ. Contam. Tox.*, 2013, **90**, 323-328.
- 469 24. G. Panda, S. Das, S. Chatterjee, P. Maity, T. Bandopadhyay and A. Guha,  
470 *Colloids Surf. B.*, 2006, **50**, 49-54.
- 471 25. H. Teemu, S. Seppo, M. Jussi, T. Raija and L. Kalle, *Int. J. Food Microbiol.*,

- 472 2008, **125**, 170-175.
- 473 26. S. S. Majumdar, S. K. Das, T. Saha, G. C. Panda, T. Bandyopadhyoy and A. K.  
474 Guha, *Colloids Surf. B.*, 2008, **63**, 138-145.
- 475 27. R. Chakravarty and P. C. Banerjee, *Bioresour. Technol.*, 2012, **108**, 176-183.
- 476 28. K. Vijayaraghavan, T. Padmesh, K. Palanivelu and M. Velan, *J. Hazard. Mater.*,  
477 2006, **133**, 304-308.
- 478 29. S. Chatterjee, S. K. Das, R. Chakravarty, A. Chakrabarti, S. Ghosh and A. K.  
479 Guha, *J. Hazard. Mater.*, 2010, **174**, 47-53.
- 480 30. A. L. F. Pereira, T. C. Maciel and S. Rodrigues, *Food Res. Int.*, 2011, **44**,  
481 1276-1283.
- 482 31. B. Escudero-López, I. Cerrillo, G. Herrero-Martín, D. Hornero-Méndez, A.  
483 Gil-Izquierdo, S. Medina, F. Ferreres, G. Berná, F. Martín and M.-S.  
484 Fernández-Pachón, *J. Agric. Food. Chem.*, 2013, **61**, 8773-8782.
- 485 32. G. Wang, M.-Y. Su, Y.-H. Chen, F.-F. Lin, D. Luo and S.-F. Gao, *Environ. Pollut.*,  
486 2006, **144**, 127-135.
- 487 33. H. Zhang, J. Chen, L. Zhu, G. Yang and D. Li, *PLOS One*, 2014, **e108572**.
- 488 34. X. Bi, L. Ren, M. Gong, Y. He, L. Wang and Z. Ma, *Geoderma*, 2010, **155**,  
489 115-120.
- 490 35. N. Naz, H. K. Young, N. Ahmed and G. M. Gadd, *Appl. Environ. Microbiol.*,  
491 2005, **71**, 4610-4618.
- 492 36. T. Beveridge and R. Murray, *J. Bacteriol.*, 1980, **141**, 876-887.
- 493 37. S. K. Das, A. R. Das and A. K. Guha, *Environ. Sci. Technol.*, 2007, **41**,

- 494 8281-8287.
- 495 38. T. Halttunen, S. Salminen and R. Tahvonen, *Int. J. Food Microbiol.*, 2007, **114**,
- 496 30-35.
- 497 39. F. Ibrahim, T. Halttunen, R. Tahvonen and S. Salminen, *Can. J. Microbiol.*, 2006,
- 498 **52**, 877-885.
- 499 40. W. Huang and Z.-m. Liu, *Colloids Surf. B.*, 2013, **105**, 113-119.
- 500 41. V. C. Srivastava, M. M. Swamy, I. D. Mall, B. Prasad and I. M. Mishra, *Colloids*
- 501 *Surf. A.*, 2006, **272**, 89-104.
- 502 42. B. Bueno, M. Torem, F. Molina and L. De Mesquita, *Miner. Eng.*, 2008, **21**,
- 503 65-75.
- 504 43. E.-J. Yang, S.-I. Kim, S.-Y. Park, H.-Y. Bang, J. H. Jeong, J.-H. So, I.-K. Rhee
- 505 and K.-S. Song, *Food Chem. Toxicol.*, 2012, **50**, 2042-2048.
- 506 44. J. Liu, W. Qu and M. B. Kadiiska, *Toxicol. Appl. Pharmacol.*, 2009, **238**,
- 507 209-214.
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## 509 **Figure legends**

510 **Fig. 1.** Transmission electron microscopic images of *Lactobacillus plantarum*  
511 CCFM8610 before and after Cd binding. (A) Untreated biomass. (B) Biomass after  
512 Cd binding. (C) Energy dispersive X-ray (EDX) spectra of untreated biomass. (D)  
513 EDX spectra of biomass after Cd binding. Scale bar = 100 nm. The experiment was  
514 carried out in aqueous solution containing Cd as cadmium chloride.

515 **Fig. 2.** Scanning electron microscopic images of *L. plantarum* CCFM8610 before and  
516 after Cd binding. (A) Untreated biomass. (B) Biomass after Cd binding. (C) Energy  
517 dispersive X-ray (EDX) spectra of untreated biomass. (D) EDX spectra of biomass  
518 after Cd binding. Scale bar = 10.0  $\mu\text{m}$ . The experiment was carried out in aqueous  
519 solution containing Cd as cadmium chloride.

520

521 **Fig. 3.** Cd binding ability of the cellular component of *L. plantarum* CCFM8610.  
522 Values are mean  $\pm$  SEM of three determinations. Significant differences ( $p < 0.05$ )  
523 between the cellular components are indicated with different letters above the bars.  
524 The experiment was carried out in aqueous solution containing Cd as cadmium  
525 chloride.

526 **Fig. 4.** Effects of chemical treatments of amino, carboxyl and phosphoryl groups on  
527 the Cd binding ability of *L. plantarum* CCFM8610. Values are mean  $\pm$  SEM of three  
528 determinations. Significant differences ( $p < 0.05$ ) between the different chemically

529 treated groups are indicated with different letters above the bars. The experiment was  
530 carried out in aqueous solution containing Cd as cadmium chloride.

531 **Fig. 5.** Effects of initial pH on Cd binding ability of *L. plantarum* CCFM8610. Values  
532 are mean  $\pm$  SEM of three determinations. The experiment was carried out in aqueous  
533 solution containing Cd as cadmium chloride.

534 **Fig. 6.** Adsorption isotherm of Cd binding by *L. plantarum* CCFM8610. Values are  
535 mean  $\pm$  SEM of three determinations. EXP, experimental data acquired in the present  
536 study; LM, Langmuir Model; FM, Freundlich Model; LFM, Langmuir-Freundlich  
537 Model. The experiment was carried out in aqueous solution containing Cd as  
538 cadmium chloride.

539 **Fig. 7.** Cd binding of *L. plantarum* CCFM8610 at different time points. Values are  
540 mean  $\pm$  SEM of three determinations. The experiment was carried out in aqueous  
541 solution containing Cd as cadmium chloride.

542

543 **Tables**544 **Table 1** kinetic models used in the present study

Kinetic models	
1. Pseudo first-order kinetic model	$q_t = q_e - [q_e \times \exp(-K_1 t)]$
2. Pseudo second-order kinetic model	$t / q_t = 1 / K_2 q_e^2 + t / q_e$

545  $q_t$  (mg/g) represents the Cd content bound by *Lactobacillus plantarum* CCFM8610 biomass at time  
 546 point “t”.  $q_e$  (mg/g) represents the equilibrium content of the Cd bound by the biomass.  $K_1$  and  $K_2$  are  
 547 the biosorption constants in the pseudo first-order kinetic model and the pseudo second-order kinetic  
 548 model, respectively.

550 **Table 2** Biosorption constants from simulations with different isotherm models

Isotherm models	Constants	
1. Langmuir $q_e = \frac{Q_{\max} b_L C_e}{1 + b_L C_e}$	$Q_{\max}$	24.69 (mg/g dry biomass)
	$b_L$	0.2494
	$R^2$ (nonlinear)	0.9679
2. Freundlich $q_e = K_F C_e^{1/n_F}$	$K_F$	6.575
	$n_F$	3.006
	$R^2$ (nonlinear)	0.9834
3. Langmuir-Freundlich dual $q_e = \frac{K_{LF} C_e^{1/n}}{1 + a_{LF} C_e^{1/n}}$	$K_{LF}$	7.941
	$a_{LF}$	0.2038
	n	1.926
	$R^2$ (nonlinear)	0.9928

551  $q_e$  (mg/g) represents the equilibrium content of the Cd bound by the biomass.  $C_e$  (mg/L) represents the  
 552 equilibrium Cd concentration. The experiment was carried out in aqueous solution containing Cd as  
 553 cadmium chloride.

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559 **Table 3** Effects of *L. plantarum* CCFM8610 treatment (2 h and 36 h) on pH and viable cell number of  
560 the juices

Juices	pH			Viable counts ( $10^7$ cfu/mL)		
	0 h	2 h	36 h	0 h	2 h	36 h
AJ1	$3.96 \pm 0.003^a$	$3.87 \pm 0.003^b$	$3.46 \pm 0.007^c$	$1.30 \pm 0.04^a$	$1.42 \pm 0.02^a$	$14.87 \pm 0.19^b$
AJ2	$3.94 \pm 0.003^a$	$3.90 \pm 0.003^b$	$3.39 \pm 0.003^c$	$1.38 \pm 0.06^a$	$1.57 \pm 0.03^a$	$15.63 \pm 0.19^b$
AJ3	$3.98 \pm 0.009^a$	$3.90 \pm 0.012^b$	$3.40 \pm 0.006^c$	$1.30 \pm 0.03^a$	$1.63 \pm 0.02^a$	$13.97 \pm 0.22^b$
TJ1	$4.36 \pm 0.021^a$	$4.20 \pm 0.008^b$	$3.17 \pm 0.009^c$	$1.39 \pm 0.06^a$	$1.80 \pm 0.11^a$	$27.37 \pm 0.70^b$
TJ2	$4.33 \pm 0.009^a$	$4.23 \pm 0.000^b$	$3.20 \pm 0.007^c$	$1.30 \pm 0.02^a$	$1.64 \pm 0.02^a$	$20.70 \pm 0.57^b$
TJ3	$4.39 \pm 0.006^a$	$4.24 \pm 0.009^b$	$3.18 \pm 0.017^c$	$1.43 \pm 0.02^a$	$1.62 \pm 0.03^a$	$21.70 \pm 0.72^b$
CJ1	$5.13 \pm 0.000^a$	$4.90 \pm 0.006^b$	$3.68 \pm 0.015^c$	$1.39 \pm 0.02^a$	$1.44 \pm 0.02^a$	$10.87 \pm 0.19^b$
CJ2	$5.09 \pm 0.003^a$	$4.93 \pm 0.029^b$	$3.61 \pm 0.012^c$	$1.25 \pm 0.03^a$	$1.35 \pm 0.03^a$	$14.27 \pm 0.56^b$
CJ3	$5.13 \pm 0.009^a$	$4.95 \pm 0.003^b$	$3.64 \pm 0.019^c$	$1.37 \pm 0.03^a$	$1.46 \pm 0.02^a$	$9.50 \pm 0.35^b$

561 Values are mean  $\pm$  SEM of three determinations. The superscript letters indicate statistically significant  
562 differences at a *p* value of  $< 0.05$  in comparisons between different time points for each juice group. AJ,  
563 TJ and CJ indicate apple juices, tomato juices and cucumber juices, respectively.

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566 **Table 4** Cd removal from juices by *L. plantarum* CCFM8610 treatment (2 h)

Juices	Cd concentration ( $\mu\text{g/mL}$ )		Cd removal (%)
	Control	<i>L. plantarum</i> CCFM8610 treated	
AJ1	$0.51 \pm 0.006^a$	$0.095 \pm 0.014^b$	$81.42 \pm 2.74$
AJ2	$1.21 \pm 0.033^a$	$0.27 \pm 0.005^b$	$78.08 \pm 0.16$
AJ3	$0.72 \pm 0.013^a$	$0.13 \pm 0.002^b$	$82.28 \pm 0.33$
TJ1	$0.50 \pm 0.005^a$	$0.13 \pm 0.002^b$	$74.59 \pm 0.21$
TJ2	$0.58 \pm 0.006^a$	$0.16 \pm 0.009^b$	$71.71 \pm 1.30$
TJ3	$0.59 \pm 0.012^a$	$0.19 \pm 0.003^b$	$67.12 \pm 0.64$
CJ1	$0.70 \pm 0.008^a$	$0.15 \pm 0.005^b$	$78.10 \pm 0.62$
CJ2	$1.71 \pm 0.013^a$	$0.29 \pm 0.006^b$	$82.87 \pm 0.25$
CJ3	$2.90 \pm 0.008^a$	$0.52 \pm 0.005^b$	$82.07 \pm 0.22$

567 Values are mean  $\pm$  SEM of three determinations. The superscript letters indicate statistically significant  
568 differences at a *p* value of  $< 0.05$  in comparisons between control and *L. plantarum* CCFM8610 treated  
569 samples for each juice group. AJ, TJ and CJ indicate apple juices, tomato juices and cucumber juices,  
570 respectively.

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579 **Table 5** Cd removal from juices by *L. plantarum* CCFM8610 treatment (36 h)

Juices	Cd concentration ( $\mu\text{g/mL}$ )		Cd removal (%)
	Control	<i>L. plantarum</i> CCFM8610 treated	
AJ1	$0.51 \pm 0.005^a$	$0.097 \pm 0.007^b$	$81.16 \pm 1.26$
AJ2	$1.25 \pm 0.036^a$	$0.26 \pm 0.013^b$	$78.89 \pm 1.30$
AJ3	$0.70 \pm 0.017^a$	$0.17 \pm 0.004^b$	$75.54 \pm 0.82$
TJ1	$0.51 \pm 0.002^a$	$0.19 \pm 0.016^b$	$62.98 \pm 0.21$
TJ2	$0.58 \pm 0.004^a$	$0.26 \pm 0.014^b$	$56.18 \pm 2.62$
TJ3	$0.61 \pm 0.005^a$	$0.22 \pm 0.012^b$	$64.35 \pm 1.97$
CJ1	$0.72 \pm 0.011^a$	$0.18 \pm 0.004^b$	$75.44 \pm 0.25$
CJ2	$1.62 \pm 0.055^a$	$0.32 \pm 0.010^b$	$80.21 \pm 0.97$
CJ3	$2.95 \pm 0.030^a$	$0.54 \pm 0.002^b$	$81.79 \pm 0.25$

580 Values are mean  $\pm$  SEM of three determinations. The superscript letters indicate statistically significant  
581 differences at a  $p$  value of  $< 0.05$  in comparisons between control and *L. plantarum* CCFM8610 treated  
582 samples for each juice group. AJ, TJ and CJ indicate apple juices, tomato juices and cucumber juices,  
583 respectively.

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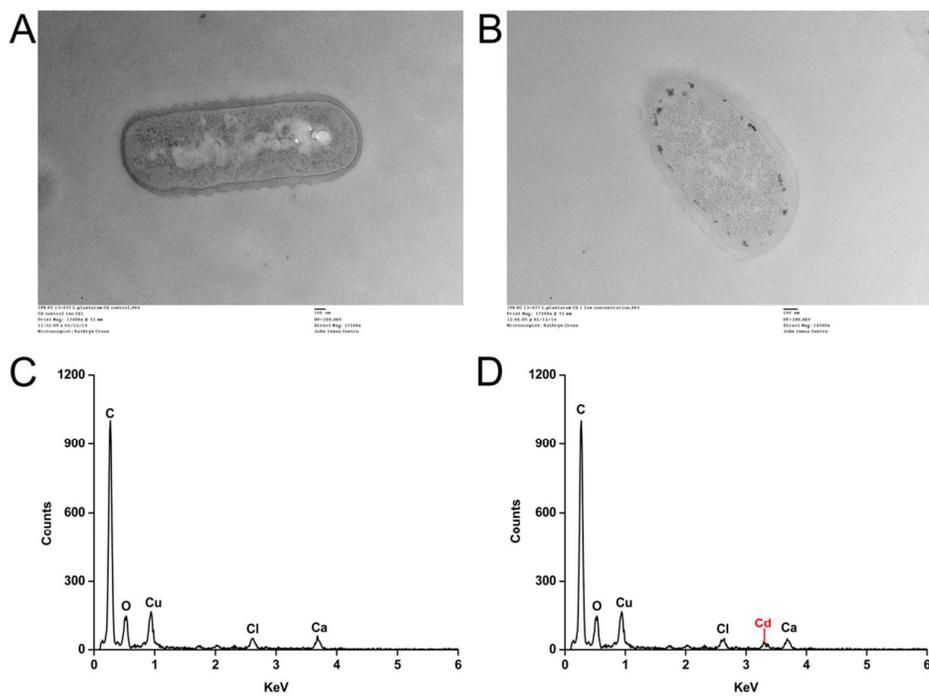
587 **Table 6** Effects of *L. plantarum* CCFM8610 treatment (2 h and 36 h) on DPPH scavenging ability of  
588 the juices

Juices	Scavenging rate of DPPH (%)		
	0 h	2 h	36 h
AJ1	$63.09 \pm 1.51^{ab}$	$59.72 \pm 0.86^a$	$67.90 \pm 1.29^b$
AJ2	$58.48 \pm 0.83^a$	$58.87 \pm 1.34^a$	$61.48 \pm 1.44^a$
AJ3	$64.79 \pm 1.75^{ab}$	$59.38 \pm 2.41^a$	$70.28 \pm 0.52^b$
TJ1	$72.98 \pm 0.81^a$	$71.10 \pm 0.61^a$	$78.51 \pm 0.76^b$
TJ2	$62.39 \pm 0.85^a$	$69.52 \pm 0.30^b$	$72.69 \pm 0.33^c$
TJ3	$58.83 \pm 1.60^a$	$67.20 \pm 0.70^b$	$69.12 \pm 0.59^b$
CJ1	$52.39 \pm 2.32^a$	$51.01 \pm 0.11^a$	$54.33 \pm 1.55^a$
CJ2	$46.80 \pm 0.81^a$	$48.04 \pm 0.68^a$	$52.69 \pm 0.73^b$
CJ3	$54.78 \pm 1.94^a$	$51.22 \pm 0.50^a$	$62.75 \pm 0.75^b$

589 Values are mean  $\pm$  SEM of three determinations. The superscript letters indicate statistically significant  
590 differences at a  $p$  value of  $< 0.05$  in comparisons between different time points for each juice group. AJ,  
591 TJ and CJ indicate apple juices, tomato juices and cucumber juices, respectively.

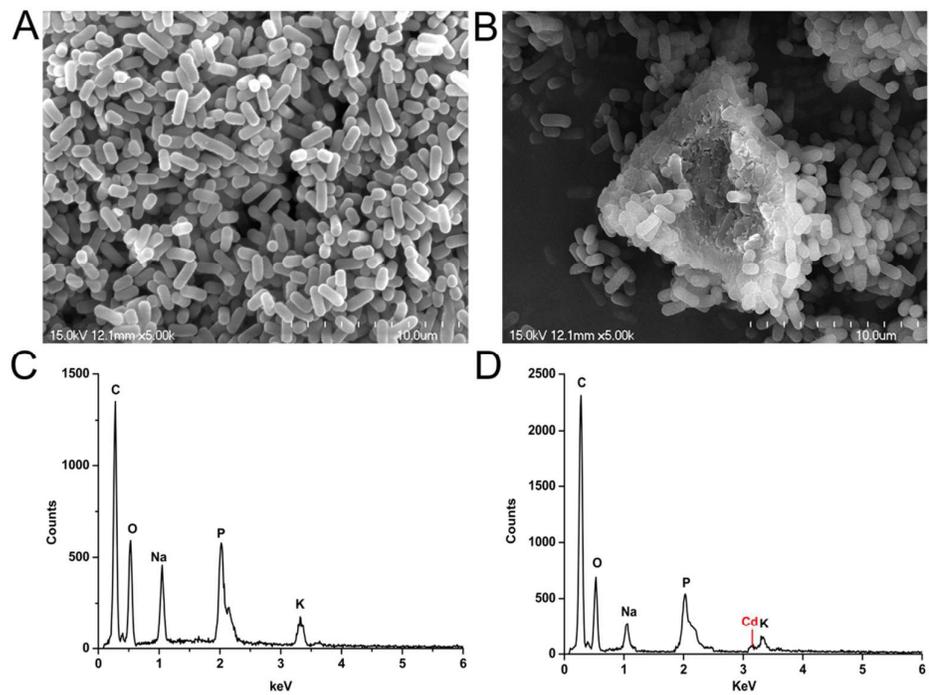
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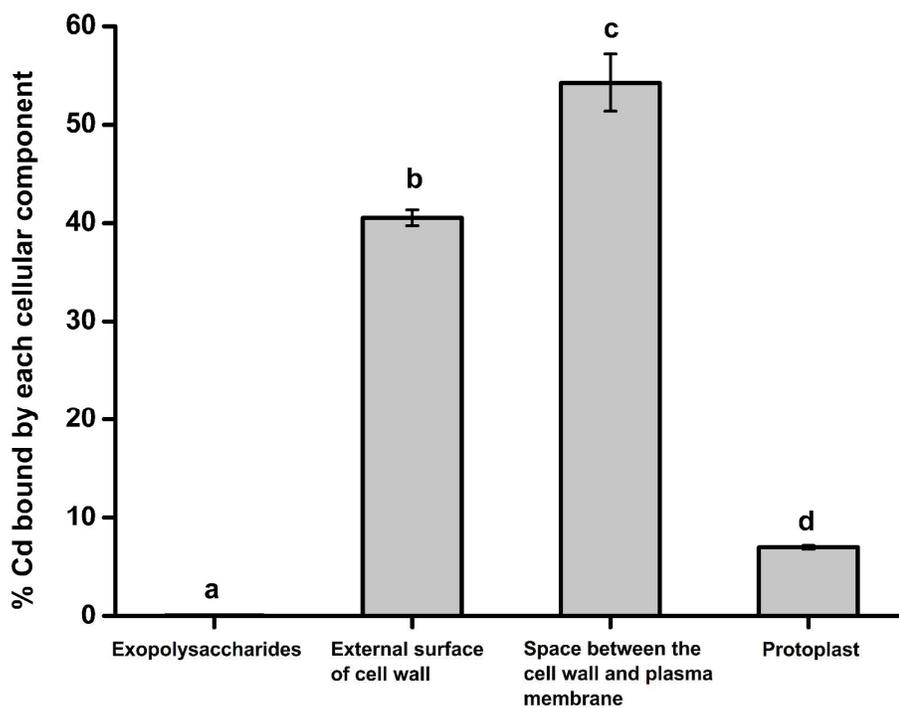
594 **Figure 1**

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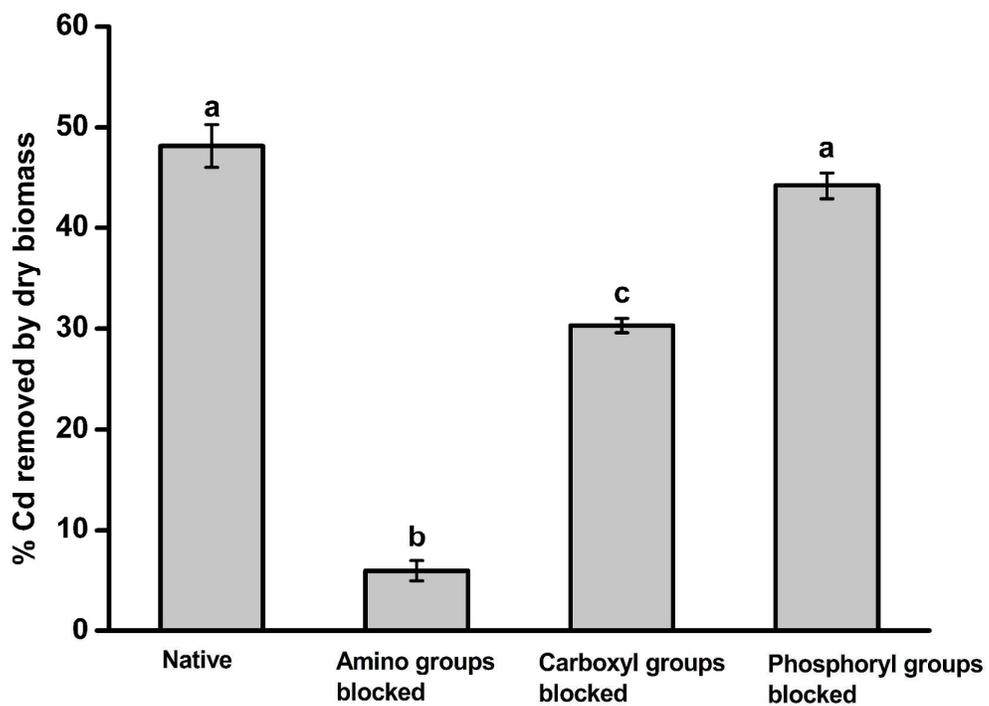
597 **Figure 2**

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599 **Figure 3**

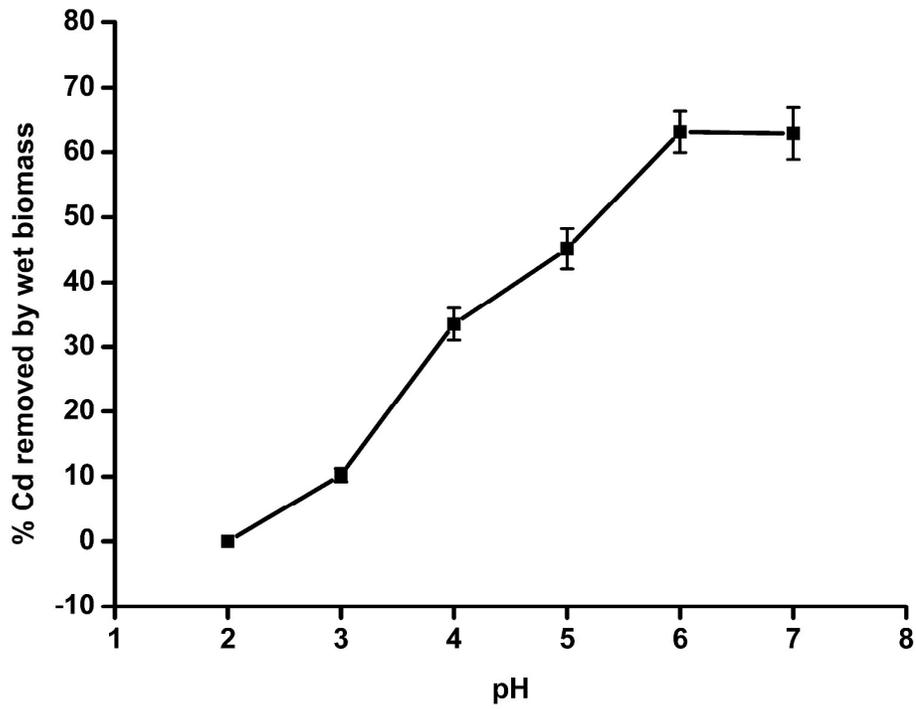
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602 **Figure 4**

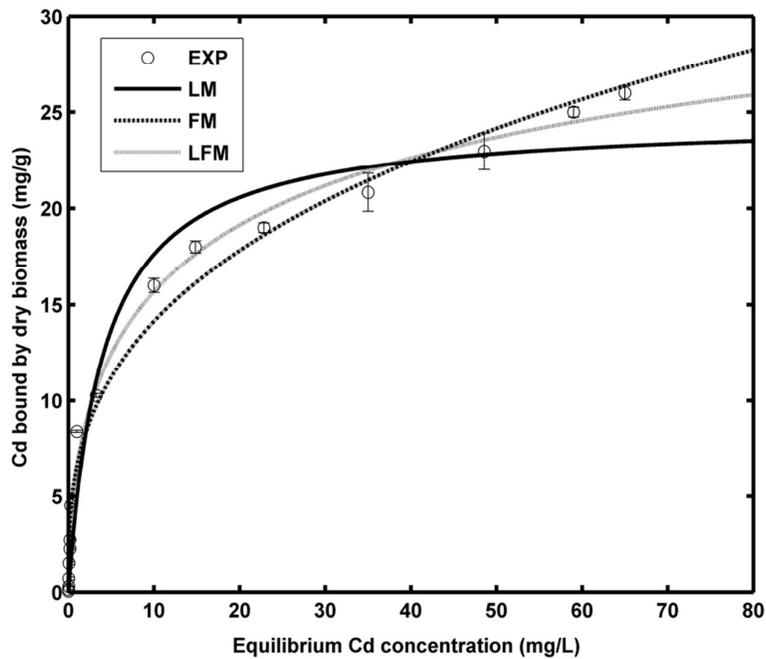
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605 **Figure 5**

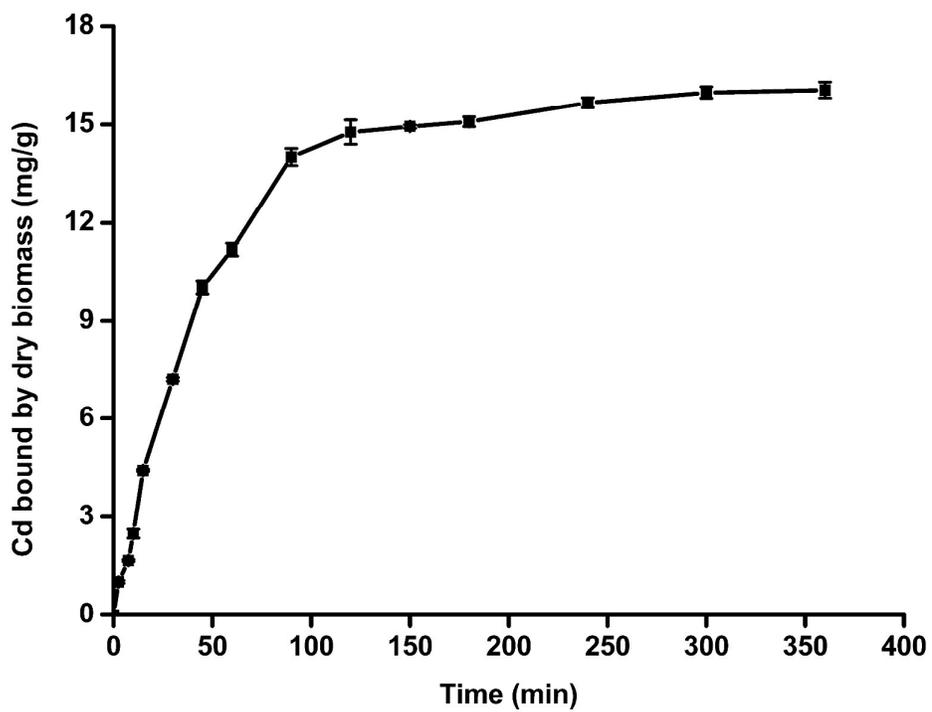
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608 **Figure 6**

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611 **Figure 7**

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621 **Table of Contents**

- Cd binding process:
- pH dependent
  - Carboxyl and amino groups related
  - Fast and efficient
  - Fit to the Langmuir-Freundlich dual isotherm model and the pseudo second-order kinetic model

**Lactic acid bacterium removes of cadmium from fruit and vegetable juices**



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