Intestine-like micro/mesoporous carbon built of chemically modified banana peel for size-selective separation of proteins

Rui-Lin Liu,ab Fu-Yu Yin,a Ji-Fang Zhang,a Jing Zhanga and Zhi-Qi Zhang*ab

An intestine-like micro/mesoporous carbon (ILMC) was fabricated by chemically modifying banana peel and used for the size-selective separation of proteins. The as-prepared ILMC has been exemplarily characterized by FTIR, XRD, SEM, TEM and N2 adsorption measurements. The adsorption property of ILMC was further evaluated with three different molecular size proteins, cytochrome c, bovine serum albumin and lysozyme, at binary and ternary hybrid solutions. The results showed that the ILMC is effective and a highly selective adsorbent for cytochrome c. This simple and procurable ILMC may be utilized as a potential and promising support for immobilizing bio-macromolecules, drug delivery and separation.

Introduction

Carbon materials (CMs) with high porosity and large pore volume are widely employed in adsorption and energy storage and as a carrier for drug delivery system.1 Extensive methods have been used to fabricate various CMs, including hard and soft templating or activation methods.2 Traditional inorganic materials, zeolite and silica, are known templates for casting porous carbon. However, this method is costly, complicated and always involves highly toxic substances, which hinders its application in large scale production. Interestingly, CMs fabricated from waste biomass have shown extensive attention and promising applications as sorption materials.3 Biomass, because of its inexpensive, easy to obtain, rapid regeneration and nontoxic nature, has qualified as a promising starting material for the synthesis of carbonaceous materials.4–7 Till now, some processing techniques, such as hydrothermal carbonization and direct pyrolysis, have been established for fabricating high quality CMs.4,8 However, whether the biomass conversion system is thermochemical or biological, there still does not appear to be a satisfactory and optimal process.4 In particular, the application in the separation and adsorption of giant molecules, such as proteins, is scarce for CMs prepared with low-valued carbon sources.

The adsorption or isolation of proteins from solution onto solid surfaces has attracted much attention because of its scientific importance and application in many areas such as biology, medicine and biotechnology.9 So far, nanochannel titania membrane,10 mesoporous silicas,11 carbon nanotubes and mesoporous carbons,12,13 nanoparticles or nanocomposite,14–17 block copolymer membranes,18 etc. have been used for adsorbing or separating proteins. Although nanoporous carbons are providing a new platform for the separation of proteins,13,19–21 it is difficult to directly apply them in large scale systems because some flaws still exist in the fabrication of mesoporous carbon such as complexity, high cost and toxicity.

Banana peel (BP), as a common biomass waste, which has abundant chemical groups including carboxyl, hydroxyl and amide groups,22 would be easily modified and assembled with various chemicals.23–25 In this work, an inexpensive carbonaceous adsorbent with intestine-like mesostructure was fabricated through chemically modifying BP and was applied to the size-selective isolation of proteins.

Experimental

Materials and chemicals

Crude BP was obtained from a local fruit market in Xi’an, China and thoroughly washed with distilled water for further use. Commercial triblock copolymer, Pluronic F127 (poly(ethylene oxide)-block-poly(propylene oxide)-block-poly(ethylene oxide), PEO106PPO70PEO106, Mw = 12 600) was purchased from Sigma-Aldrich Corp. Al(NO3)3·9H2O, NaOH, hydrofluoric acid (HF), Na2CO3, NaHCO3 and ethanol were purchased from Sinopharm Chemicals Co., Ltd (Shanghai, China). Cytochrome c (Cyt c, purity >98%), bovine serum albumin (BSA, purity >98%) and lysozyme (Lyz, purity >98%) were purchased from Sigma-Aldrich Corp. All chemicals were of analytical grade and were used without further purification. Ultra-pure water (18.2 MΩ cm) was

*Key Laboratory of Analytical Chemistry for Life Science of Shaanxi Province, School of Chemistry and Chemical Engineering, Shaanxi Normal University, Xi’an 710062, China
*a Key Laboratory of Medicinal Resource and Natural Pharmaceutical Chemistry (Shaanxi Normal University), Ministry of Education, Xi’an 710062, China. E-mail: zqzhang@snnu.edu.cn; Fax: +86-29-81530792
produced by a Millipore purification system (Millipore, MA, USA) and was used to prepare all aqueous solutions.

**Fabrication of intestine-like micro/mesoporous carbon**

Intestine-like micro/mesoporous carbon (ILMC) was fabricated using a previously method with some modifications.\(^2\) In brief, about 2 kg BP was washed and cut into small pieces and then completely submerged in 2 L of aluminum nitrate aqueous solution (1.0 mol L\(^{-1}\)) for 5 days at 100 °C. As a result, the resulting macroscopic porous framework complexes (MPFC) were obtained. Subsequently, about 10 g of MPFC was impregnated into 50 mL of F127 (2.0 g) ethanol solution for 15 h at room temperature and then further subjected to thermo-polymerization at 130 °C for 7 h in a vacuum drying oven, and finally the multiple components co-assembled into yellow gel-like composites. Finally, the gel-like composites were heated under N\(_2\) to 500 °C with a heating rate of 1 °C min\(^{-1}\) and maintained at 500 °C for 2 h to decompose the surfactant of F127, and subsequently heated to 650 °C with a heating rate of 3 °C min\(^{-1}\) and maintained at 650 °C for 4 h. The obtained black monoliths were ground and immersed in 15 wt% HF for 24 h to remove the Al component. The black precipitates were washed with deionized water and dried for 24 h at 100 °C.

**Characterizations**

A Bruker Tensor 27 spectrometer with the KBr pellet technique was used to measure the infrared spectroscopy (FTIR) ranging from 400 cm\(^{-1}\) to 4000 cm\(^{-1}\). A Rigaku D/Max-3C X-ray diffractometer with a Cu K\(_{\alpha}\) radiation (\(\lambda = 1.54 \AA\)) was applied for measuring the X-ray diffraction (XRD) pattern at 40 kV and 30 mA with an Inel CPS 120 hemispherical detector. The FEI Quanta 200 scanning electron microscope was used to obtain the scanning electron microscopy (SEM) images at an accelerating voltage of 20 kV. A JEOL JEM-2100 microscope was used to obtain the transmission electron microscopy (TEM) images at 200 kV. Samples were first dispersed in ethanol and then collected using carbon-film-covered copper grids for analysis.

The Micromeritics ASAP 2020 volumetric adsorption analyzer was employed to measure nitrogen sorption isotherms at 77 K. Prior to analysis, the sample was automatically and manually degassed for 9 h and 4 h, respectively, under vacuum at 523 K. The sample was then transferred to the analysis system and cooled in liquid nitrogen. The specific surface area was calculated by the Brunauer–Emmett–Teller (BET) equation based on the adsorption data at relative pressure ranging 0.05–0.3; the total pore volume was evaluated by converting the adsorption volume of nitrogen at relative pressure of 0.976 to equivalent liquid volume of the adsorbate, while the micropore volume was obtained by the t-plot method. Pore size distributions were determined from the adsorption branches of the isotherms using the nonlocal density functional theory (NLDFT) model.

**Zeta potential measurement**

The point of zero charge (pH\(_{PZC}\)) was carried out according to a previously reported method.\(^2\) In brief, the ILMC (0.15 g) was added to 50 mL 0.1 M KNO\(_3\) solution by adjusting the pH values between 2 and 12 with 1 M HCl or 1 M NaOH, and the final pH was measured in triplicate after 48 h under agitation. The difference between the initial (pH\(_{0}\)) and final pH (pH\(_{i}\)) values (\(\Delta \text{pH} = \text{pH}_{i} - \text{pH}_{0}\)) was plotted against the pH. The point of intersection of the resulting curve with abscissa, at which \(\Delta \text{pH} = 0\), provided the pH\(_{PZC}\).

**Isolation of protein**

The size-selective isolation of proteins on ILMC was investigated in the binary mixtures of Cyt c and BSA, Lyz and Cyt c, and the ternary mixture of BSA, Lyz and Cyt c systems. In each adsorption trial, 50.0 mg of ILMC was suspended in 20 mL bicarbonate buffer solution (pH 9.6) with a certain concentration of proteins. The resulting mixture was continuously shaken at 303 K for 16 h at 130 rpm until equilibrium was reached. The mixture was centrifuged for 4 min at 2500 g. The relative percentage of protein adsorbed was measured by UV absorption at 409 nm for Cyt c, and at 280 nm for BSA and Lyz.

**Results and discussion**

**Characterizations of ILMC**

FTIR spectrum of ILMC displayed some typical adsorption peaks (Fig. 1a), which includes the broad band at 3420 cm\(^{-1}\) related to O–H groups, 1618 cm\(^{-1}\) assigned to C=O stretching of carboxylic acid or ester and 1384 cm\(^{-1}\) identical to COO\(^{-}\) anion stretching.\(^2\) XRD spectrum of ILMC displayed two adsorption peaks at 2\(\theta\) = 25 and 44° (Fig. 1b), which were attributed to the (002) and (100) diffractions for graphitized carbon.\(^2\) In addition, the pH\(_{PZC}\) of ILMC was measured using the acid/base titration method, and the pH\(_{PZC}\) was ca. 5.2 (Fig. 1c).

The SEM images showed that the surface of ILMC exhibits a well pronounced intestine-like structure with a series of irregular cavities distributed around the cross section (Fig. 2a–c). The TEM image showed that the ILMC material has abundant 2D nanoporous textures (Fig. 2d–h), which can provide a rapid transfer channel of molecules through the mesostructure.

As shown in Fig. 3a, the nitrogen sorption isotherm features an intermediate between type-I and type-IV curves with a sharp capillary condensation step in the relative pressure range of 0.4–0.6 and H1-type hysteresis loop, which is indicative of uniform cylindrical pores in the size range of mesoporous. The steep increase in the adsorbed volume at low relative pressure was related to the presence of micropores, and the desorption hysteresis at medium relative pressure revealed the existence of developed mesopores.\(^2\) The pore size distribution of ILMC was ascertained by the NLDFT model, which is mainly distributed between 0.5 and 20 nm (Fig. 3b and Table 1), with an average pore size of 2.60 nm. This finding shows that the vast majority of the pores fall in the range of mesoporous for ILMC.

**Isolation of protein**

To explore the as-prepared ILMC as a new protein adsorbent in size-selective isolation, three typical proteins with different...
molecular sizes, including Cyt c (MW 12.3 kDa, pI 9.8, molecular dimensions 2.6 × 3.2 × 3.0 nm), BSA (MW 66.4 kDa, pI 4.8, molecular dimensions 4.0 × 4.0 × 14 nm) and Lyz (MW 14.4 kDa, pI 11.2, molecular dimensions 3.0 × 3.0 × 4.5 nm), were chosen as model molecules (Fig. 4). The relative percentage of adsorbed protein was used for evaluating the adsorption efficiency.

Because the adsorption capacity could be maximized near the pI,11,14 adsorption close to the isoelectric point was first investigated for Cyt c, BSA and Lyz at pH 9.6, pH 4.8 and pH 11.2, respectively. The corresponding results (Fig. 5) showed that the adsorptions of ILMC were poor for BSA (7.80 ± 0.12%, \( n = 3 \)) and Lyz (6.61 ± 0.17%, \( n = 3 \)), while the adsorption for Cyt c (63.87 ± 0.19%, \( n = 3 \)) was very strong. The results mainly attributed to the dimensions of BSA and Lyz are larger than the average mesoporous pore size of ILMC (approximately 3.55 nm), most of the BSA and Lyz were excluded from the pores of the adsorbent. The dimension of Cyt c is lower than the average mesoporous pore size of ILMC; hence, maximal adsorption was acquired. This indicates that the size selectivity is excellent using ILMC with a narrow mesopore size distribution as an adsorbent to separate Cyt c at pH 9.6.

The adsorption of Cyt c is typically determined by electrostatic and hydrophobic interactions.11 Because the net charge of Cyt c is very low at pH 9.6, which resulted in the minimal electrostatic repulsion between the amino acid residues on the surface of Cyt c molecules, the Cyt c molecules can be adsorbed into the pores of the ILMC by hydrophobic interactions.

In order to further understand the influence of electrostatic interaction between adsorbent and protein in the size-selective isolation of Cyt c, the adsorption trials were operated for BSA and Lyz at pH 9.6, respectively. The results showed that the adsorption of ILMC for BSA presented relatively poor efficiency (5.76 ± 0.18%, \( n = 3 \)) than that at pI (Fig. 5a), while the adsorption for Lyz (11.43 ± 0.23%, \( n = 3 \)) was markedly higher than that at pI (Fig. 5b). It could be explained as the electrostatic repulsion between the carbon surface (pI ≈ 5.2, negative charge) and BSA molecules (negative charge) at pH 9.6 leads to negative adsorption of BSA. Lyz holds a little positive charge at
pH 9.6 and makes more contribution to the adsorption on ILMC. In addition, Lyz has a prolate spheroid shape with two characteristic cross-sections: a side of dimensions roughly 3.0 × 4.5 nm and an end of dimensions 3.0 × 3.0 nm, so the size of part of Lyz could match the pores of the adsorbent and resulted in more electrostatic adsorption. Higher percentage of adsorbed Cyt c at pH 7.4 (75.02 ± 0.26%, n = 3) than that at pH 9.6 (Fig. 5c) further demonstrated that the electrostatic interaction between adsorbent and protein affected the adsorption capacity.

Considering that the anchoring ability of COOH groups located inside and at the entrance of the mesoporous cavity might obstruct the desorption of protein molecules from the pore channels of carboxy carbon at pH 7.4, the pI of Cyt c at pH 9.6 was determined to isolate Cyt c from a protein mixture solution. It could be seen that the amount of adsorbed Cyt c in

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**Table 1** Porosity structures of the ILMC prepared from BP

<table>
<thead>
<tr>
<th>Properties</th>
<th>ILMC</th>
</tr>
</thead>
<tbody>
<tr>
<td>BET surface area (m² g⁻¹)</td>
<td>995</td>
</tr>
<tr>
<td>Langmuir surface area (m² g⁻¹)</td>
<td>1364</td>
</tr>
<tr>
<td>Total pore volume (cc g⁻¹)</td>
<td>0.63</td>
</tr>
<tr>
<td>Micropore volume (cc g⁻¹)</td>
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<tr>
<td>Mesopore volume (cc g⁻¹)</td>
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</tr>
<tr>
<td>Average pore size (nm)</td>
<td>2.60</td>
</tr>
<tr>
<td>Average mesoporous size (nm)</td>
<td>3.55</td>
</tr>
</tbody>
</table>

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**Fig. 3** Nitrogen sorption isotherms, cumulative pore volumes and pore size distributions of ILMC before (a and b) and after (c and d) adsorbing Cyt c from a ternary solution of Cyt c, BSA and Lyz.

**Fig. 4** Schematic of the ILMC toward selective adsorption of Cyt c among BSA (pink ellipsoid), Lyz (blue dumbbell-shaped) and Cyt c (green sphere) at pH 9.6.

**Fig. 5** UV-Vis spectra of protein solutions before and after being adsorbed on ILMC. (a) BSA (0.6 mg mL⁻¹) in carbonate buffer solution (pH 9.6) and acetic acid buffer saline (pH 4.8); (b) Lyz (0.4 mg mL⁻¹) in carbonate buffer solution (pH 9.6) and phosphate buffer solution (pH 11.2); (c) Cyt c (0.4 mg mL⁻¹) in carbonate buffer solution (pH 9.6) and phosphate buffer solution (pH 7.4).
the binary solution of Cyt c and Lyz (Fig. 6a) is lower than that in the binary solution of Cyt c and BSA (Fig. 6b). The difference in the selective adsorption of Cyt c from different competitive proteins might indicate that the interaction between Cyt c and Lyz is relatively stronger than that of Cyt c and the large protein BSA.\(^1\) Fig. 6c shows the UV-Vis spectra of Cyt c, BSA and Lyz ternary mixed protein solution before and after the adsorption. The peak at 280 nm for BSA and Lyz decreased by 21.63% and the peak intensity of Cyt c at 409 nm was dramatically reduced (50.05 ± 0.23%, \(n = 3\)) after adsorption, indicating that Cyt c is more easily adsorbed on ILMC than BSA and Lyz. Such selectivity is mainly attributed to numerous large accessible mesopores and the high surface area of ILMC. For better comparison, the commercial activated carbon as a representative micropore matrix was investigated for the adsorption of ternary mixed protein solution, and the result indicated that the selectivity to Cyt c was very poor (data not shown). This further demonstrated that the adsorption selectivity of ILMC to Cyt c is mainly attributed to Cyt c having a smaller spherical size than the ILMC mesopore window.

**Verification for size selective isolation of proteins**

To better validate whether Cyt c molecule enters the mesopore of ILMC, the adsorbent was characterized by nitrogen adsorption before and after Cyt c adsorption (Fig. 3a–d). As expected, the amounts of nitrogen adsorbed (from 428 to 298 cc g\(^{-1}\)), specific surface areas (from 995 to 518 m\(^2\) g\(^{-1}\)) and total pore volumes (from 0.63 to 0.42 cc g\(^{-1}\)) of ILMC decreased after Cyt c adsorption. These numerical values clearly indicate that the Cyt c molecules are adsorbed inside the intestine-like mesopore of ILMC without affecting the structural integrity of the parent material.

**Conclusions**

In this work, for the first time, an ILMC with high surface area (995 m\(^2\) g\(^{-1}\)), large pore volumes (0.63 cc g\(^{-1}\)) and narrow mesopore size distribution (ca. 3.55 nm) was fabricated by chemically modifying BP and was successfully used for the selective isolation of Cyt c from a mixed protein solution of BSA, Lyz and Cyt c. The findings revealed that the pore size of ILMC plays a key role in the selective adsorption of Cyt c from a mixed protein solution, associated with the electrostatic and hydrophobic interactions as well. This low-cost carbonaceous adsorbent offers a very useful medium to separate biomolecules with different molecular sizes. It is believed that further development of this carbon material would benefit controlled release applications, biosensing, nanocarriers and so on.

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**Notes and references**