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## COMMUNICATION

***Escherichia coli* trap in human serum albumin microtubes†**

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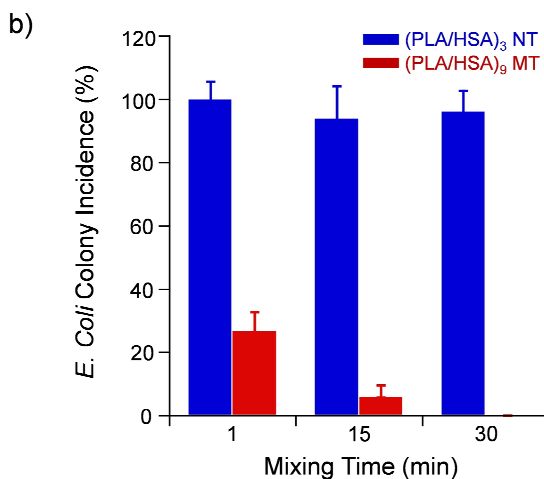
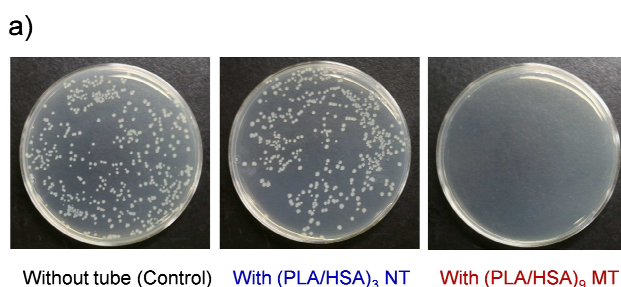
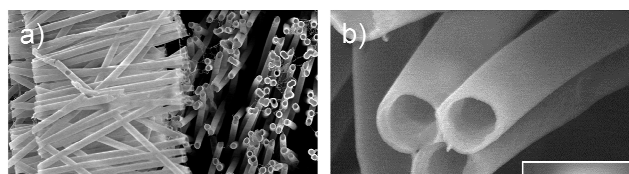
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We describe the template synthesis of human serum albumin microtubes (MTs) and highlight their *Escherichia coli* (*E. coli*) trapping capability with extremely high efficiency. The *E. coli* was loaded into the one-dimensional pore space interior of the tubule. Similar MTs including an Fe<sub>3</sub>O<sub>4</sub> layer also captured *E. coli* and were manipulated by exposure to a magnetic field.

Hollow, cylindrical, nanometer-scale structures comprising biomaterials, i.e. bionanotubes, have attracted considerable attention because of their potential applications as molecular trapping devices, drug delivery containers, and enzymatic reactors.<sup>1–13</sup> Alternate layer-by-layer (LbL) build-up assembly of proteins,<sup>4,6,7,9,10</sup> DNAs,<sup>5,8,11</sup> and antibodies<sup>12,13</sup> in nanoporous membrane enables creation of structurally defined smart nanotubes (NTs) with versatile biochemical reactivities. The one-dimensional (1D) pore space interior of the tubule can be tailored by deposition of a desired material. Therefore, many investigators have explored the loading of nanometer-size entities into the channel, such as pharmaceutical drugs,<sup>9,12</sup> bioactive spheres,<sup>9</sup> inorganic colloids,<sup>11</sup> and infectious viruses.<sup>13</sup> Another challenging subject in this field would be the trapping of a living organism. *Escherichia coli* (*E. coli*), a rod-shaped gram-negative bacterium, is the smallest organism of the micrometer-scale world (0.4–0.7 μm width, 2–4 μm length). Many strains are harmless, but some serotypes can cause severe poisoning in humans, such as enterohemorrhagic *E. coli* O157.<sup>14</sup> If one were able to generate a unique *E. coli* trap in protein microtubes



**Fig. 2** (a) Appearance of LB agar plates spread with *E. coli* dispersion which was incubated with (PLA/HSA)<sub>3</sub> NTs or (PLA/HSA)<sub>9</sub> MTs for 30 min (after 16-h cell culture at 37 °C). (b) Relation between mixing time and colony incidence [ $N_c(\text{NT})$  or  $N_c(\text{MT}) / N_c(\text{Control}) \times 100$ ] ( $n = 3$ ).

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† Electronic Supplementary Information (ESI) available: Experimental section, SEM image of *E. coli*, CLSM image of CTC-*E. coli*, and relation between mixing time and cell viability of *E. coli*. See DOI: 10.1039/x0xx00000x

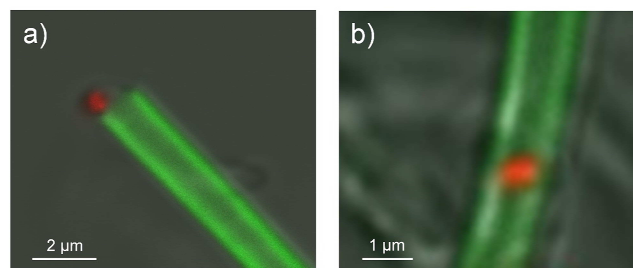
(MTs), then it would have an important contribution not only to bioseparation chemistry, but also on diverse aspects of human health. This paper is the first to describe the synthesis and structure of human serum albumin (HSA)-based MTs, and to highlight their excellent *E. coli* trapping capability.

The MTs were fabricated by template synthesis using electrostatic LbL assembly.<sup>9</sup> Briefly, positively charged poly-L-arginine (PLA) and negatively charged HSA were alternately deposited (nine-cycles) onto the pore wall of a track-etched polycarbonate (PC) membrane (1.2  $\mu\text{m}$  pore size). Subsequent dissolution of the PC framework in *N,N*-dimethylformamide and freeze-drying of the liberated core yielded (PLA/HSA)<sub>9</sub> MTs as white powder. SEM measurements revealed the formation of uniform hollow cylinders with  $1.0 \pm 0.02 \mu\text{m}$  outer diameter and  $148 \pm 5 \text{ nm}$  wall thickness (Fig. 1a, 1b). The tube lengths (ca. 25  $\mu\text{m}$ ) coincided well with the PC membrane pore depth. In contrast, 12-layered (PLA/HSA)<sub>6</sub> MTs were wrinkled and fragile, which implies that six-cycle injection is insufficient to generate stiff MTs. We concluded that at least nine-cycle deposition is necessary to construct a robust 1- $\mu\text{m}$ -width cylinder by our templating synthesis. Based on the general principle of LbL membrane growth, we hypothesized a nine-layered cylinder model. The average thickness of a PLA/HSA bilayer in the MT is estimated as 16.4 nm. If one postulates the dimensions of HSA to be 8 nm from the data of the single-crystal structure<sup>15</sup> and small-angle X-ray scattering analyses,<sup>16</sup> then the PLA layer thickness is calculated as 8.4 nm, which is between the reported values for typical polyelectrolyte layers prepared in the porous template by wet process.<sup>17–19</sup>

The obtained (PLA/HSA)<sub>9</sub> MTs were dispersed in deionized water, yielding a slightly turbid solution. To evaluate the morphology and stability of the MTs in water, the aqueous dispersion was freeze-dried *in vacuo*. SEM images demonstrated that the tubular walls swelled considerably and that their thickness became  $250 \pm 7 \text{ nm}$  in water (Fig. 1c). It is interesting that the outer diameter was unaltered ( $1.0 \pm 0.04 \mu\text{m}$ ). Consequently, the inner pore size diminished to ca. 500 nm. The average thickness of an individual PLA/HSA bilayer was measured as 28 nm in water. Under the assumption that the HSA size did not change ( $\leq 8 \text{ nm}$ ),<sup>15,16</sup> the PLA layer thickness might be 20 nm. This value is apparently larger than that of the dry form, but it is similar to the values of the polyelectrolyte layers in the NTs prepared under pressure conditions.<sup>9,20</sup> We determined the swelling ratio of the PLA layer ( $\alpha_{\text{PLA}}$ : a ratio between the section area in swollen state and that in a dried state) to be 2.1 (see ESI), which is almost identical to the data reported in the literature for general polyelectrolytes (1.2–4.0).<sup>21,22</sup> Morphologies of the (PLA/HSA)<sub>9</sub> MTs were retained for more than 24 h in water at 25 °C.

Then we measured the *E. coli* capture capability of the (PLA/HSA)<sub>9</sub> MT. The dimensions of *E. coli* K12 (XL10-gold) we used were 425 nm width and approximately 2–3  $\mu\text{m}$  length, as revealed by SEM measurements (Fig. S1). The colony incidence of the sample solution after mixing with the MTs was assayed using standard plate count method. First, *E. coli* ( $1 \times 10^8 \text{ CFU/mL}$ , 100  $\mu\text{L}$ ) was added to the aqueous solution of (PLA/HSA)<sub>9</sub> MTs (ca. 150  $\mu\text{g/mL}$ , 900  $\mu\text{L}$ ).<sup>23</sup> The resultant mixture was incubated with gentle rotation at 25 °C for 1–30 min. Then a part of the solution was

spread on LB agar plate and cultured at 37 °C for 16 h. The colonies appearing on the plate [ $N_c(\text{MT})$ ] were markedly fewer than those of identically treated *E. coli* without the tubes [ $N_c(\text{Control})$ ]. To our surprise,  $N_c(\text{MT})$  became completely zero by 30-min mixing with the MTs (Fig. 2a, 2b); the disappearance yield ( $100 - \text{colony incidence}$ ) reached 100%. Incubation with the 6-layered thin (PLA/HSA)<sub>3</sub> nanotubes (NTs) (ca. 200 nm inner diameter) exhibited *no* change in the colony number. We reasoned that the *E. coli* (425 nm width) entered the pore of the MT (ca. 500 nm), although it is too large to enter the narrow NT's channel (ca. 200 nm).



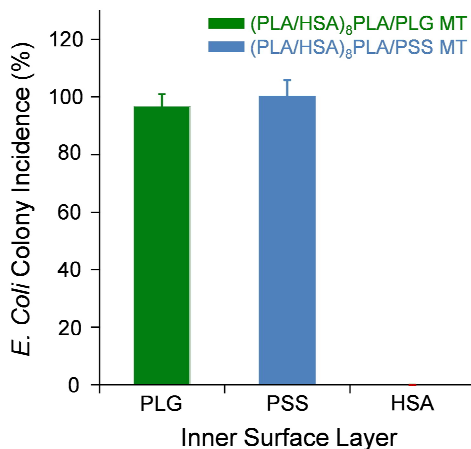
**Fig. 3** Overlapping images of DIC and CLSM of fluorescent MTs incorporating CTC-*E. coli*. (a) CTC-*E. coli* remained at the terminal of the tube and (b) small CTC-*E. coli* reached the depth of the tube. Ex. 488 nm.

The incorporation of *E. coli* into the MT was proved using confocal laser scanning microscopy (CLSM). To visualize the tube, a fluorescein-labeled HSA (f-HSA) was exploited as an intermediate layer component, yielding (PLA/HSA)<sub>7</sub>PLA/f-HSA/PLA/HSA MTs (fluorescent MTs). *E. coli* was also stained with 5-cyano-2,3-ditolyl tetrazolium chloride (CTC). This labeling agent specifically stained the edge of *E. coli* rod (Fig. S2), which is quite helpful to distinguish the *E. coli* position in the tube. In CLSM images of the mixture solution, the tube wall fluoresced green (Em. 522 nm); the hollow structure was clearly visible. CTC-*E. coli* also showed sharp fluorescence in red (Em. 630 nm). Overlapping these pictures with a DIC image demonstrates that CTC-*E. coli* was loaded into the 1D pore space of the MT (Fig. 3). Remarkably, the edge of *E. coli* appeared from the mouse of the tube in most cases, thereby forming “bacterium-corked MT” (Fig. 1e, 3a). The *E. coli* width is only slightly smaller than the pore size (ca. 500 nm). Therefore it cannot enter the deep inside of the tubule. Sometimes, a few *E. coli* become able to diffuse into the channel and reach the tube depth (Fig. 3b).

Does *E. coli* really like the 1D hollow microspace with high density of HSA? To make clear this point, we prepared comparable MTs with different interior surface using poly-L-glutamic acid sodium salt (PLG) and poly(sodium 4-styrenesulfonate) (PSS) (both negatively charged polymers) instead of the last layer of HSA. SEM measurements revealed that the (PLA/HSA)<sub>8</sub>PLA/PLG MTs and (PLA/HSA)<sub>8</sub>PLA/PSS MTs possess the same structures (outer/inner diameters and length) as (PLA/HSA)<sub>9</sub> MTs. Predictably, these MTs without the HSA pore wall could not capture *E. coli* (Fig. 4). We inferred that the inner surface wall of the tube must be comprised of HSA to entrap *E. coli*. HSA can reversibly bind many bacterial species by interacting with expressed surface proteins.<sup>24</sup> Bacteria move in response to a chemical stimulus, i.e. chemotaxis. For

instance, *E. coli* directs its movements according to certain chemicals in the environment.<sup>25</sup> We reasoned that multiple factors contribute to the *E. coli* capture into the (PLA/HSA)<sub>9</sub> MT.

To clarify the fate of the entrapped *E. coli*, the cell viability was measured using WST assay. It is noteworthy that *E. coli* accommodated in the (PLA/HSA)<sub>9</sub> MT was metabolically inactive.



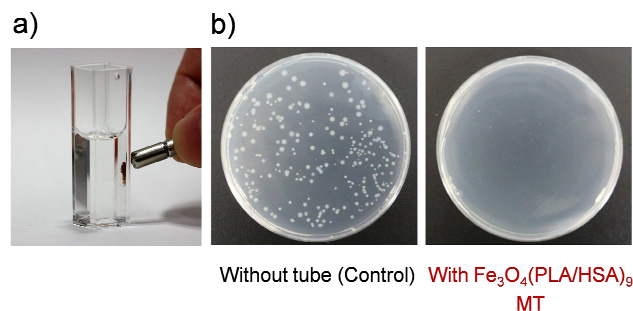
**Fig. 4** Relation between mixing time and colony incidence of *E. coli* ( $n = 3$ ).

After 1 h mixing, almost all *E. coli* lost cell proliferation ability (Fig. S3). One possible explanation is that *E. coli* growth might be blocked by stopping cell deviation. In the narrow space of the MT, bacteria cannot reproduce through asexual reproduction by binary fission. Another proposed mechanism is degradation of the HSA wall by OmpT, which is an aspartyl protease found on the outer membrane of *E. coli*.<sup>26</sup> This protease on the *E. coli*'s surface comes into contact with the HSA inner-wall and cleaves the polypeptide. Subsequently, the exposed cationic PLA layer might cause cytotoxicity.

Moreover, we introduced a magnetite ( $\text{Fe}_3\text{O}_4$ ) nanoparticle layer into the MT. Magnetic-field assisted bioseparation using spherical particles including  $\text{Fe}_3\text{O}_4$  has been an area of particular recent interest because of its diverse medical applications.<sup>27,28</sup> Nevertheless, few reports describe the use of hollow magnetic cylinders for bioseparation.<sup>12,17,29</sup> An HSA-based MT bearing a ferrimagnetic layer would become a magnetically responsive trap for *E. coli*. The magnetic tubes were prepared using a similar LbL assembly procedure with  $\text{Fe}_3\text{O}_4$  nanoparticle. SEM observations of the  $\text{Fe}_3\text{O}_4$ (PLA/HSA)<sub>9</sub> MTs revealed the formation of highly ordered arrays of the MTs with outer diameter of  $1.0 \pm 0.03 \mu\text{m}$  and maximum length of *ca.*  $25 \mu\text{m}$  (Fig. 1d). The wall thickness was  $155 \pm 7 \text{ nm}$ , which is slightly thicker than that observed in the (PLA/HSA)<sub>9</sub> MTs.

Finally, the *E. coli* capture ability of these magnetic MTs were evaluated. As expected, the  $\text{Fe}_3\text{O}_4$ (PLA/HSA)<sub>9</sub> MTs can be collected by exposure to a magnetic field. By bringing a neodymium magnet close to the quartz cuvette including the *E. coli* solution with the  $\text{Fe}_3\text{O}_4$ (PLA/HSA)<sub>9</sub> MTs, the brown tubes were attracted rapidly to the magnet; the solution became colourless (Fig. 5a). Detaching the

magnet from the cuvette liberated the MTs in the aqueous phase. This magnetic-field-induced collection–dispersion was observed to be reversible. Then a part of the upper clear solution in a cuvette, in which the *E. coli* loaded- $\text{Fe}_3\text{O}_4$ (PLA/HSA)<sub>9</sub> MTs were gathered at the bottom by the magnetic field, was spread on an LB agar plate and cultured at  $37^\circ\text{C}$  for 16 h. The number of colonies appearing on the plate became zero after 30-min mixing with  $\text{Fe}_3\text{O}_4$ (PLA/HSA)<sub>9</sub> MTs (Fig. 5b). The disappearance yield reached 100%. We



**Fig. 5** (a) Photograph of collection of *E. coli*-loaded  $\text{Fe}_3\text{O}_4$ (PLA/HSA)<sub>9</sub> MTs by magnetic field. (b) Appearance of LB agar plates spread with *E. coli* dispersion which was incubated with  $\text{Fe}_3\text{O}_4$ (PLA/HSA)<sub>9</sub> MTs for 30 min (after 16-h cell culture at  $37^\circ\text{C}$ ).

concluded that the *E. coli* entered the channel of  $\text{Fe}_3\text{O}_4$ (PLA/HSA)<sub>9</sub> MTs in a similar fashion to that of (PLA/HSA)<sub>9</sub> MTs.

In conclusion, the blood serum protein HSA microtubes ensnared the *E. coli* perfectly. The efficiency of removal by a single treatment with (PLA/HSA)<sub>9</sub> MTs was over  $-7\log$  order. This remarkable result will serve as a trigger to produce a new and productive field of removing device systems for bacteria. For example, elimination of enterohemorrhagic *E. coli* O157, using (PLA/HSA)<sub>9</sub> MTs is expected to be of incredible medical importance. Furthermore, *E. coli*-loaded MTs containing an  $\text{Fe}_3\text{O}_4$  layer were magnetically manipulated in solution. Recombinant HSA is currently manufactured on an industrial scale,<sup>30</sup> which enables production of these protein MTs for practical use.

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