



# Magneto-controlled potentiometric assay for *E. coli* based on cleavage of peptide by outer-membrane protease T



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## ABSTRACT

Rapid, sensitive and reliable *Escherichia coli* (*E. coli*) detection and identification are critically important to protect public health. Here, we describe a magneto-controlled potentiometric assay for specific detection of *E. coli* cells by making use of the *E. coli* outer-membrane protease T (OmpT). OmpT is an endopeptidase that specifically cleaves peptide at dibasic sites. A rationally designed peptide serving as both OmpT substrate and potentiometric signal reporter was immobilized on magnetic beads. The rapid accumulation and extraction of peptide-functionalized magnetic beads on a polymeric membrane doped with an ion exchanger can be achieved using a magnetic force. The magnetic-field-assisted extraction of the peptide into the polymeric membrane ion-sensitive sensor, as confirmed by Fourier transform infrared spectroscopy and X-ray photoelectron spectroscopy, can lead to a rapid, stable and reproducible potential change. OmpT is capable of cleaving the positively charged peptide on the magnetic beads, thus resulting in charge density change. The change in charge density and subsequently the potential change can be readily detected and used for quantification of *E. coli* at levels down to  $5.0 \times 10^3$  CFU mL<sup>-1</sup>. This work provides a versatile, rapid and reliable potentiometric method for *E. coli* detection.

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## 1. Introduction

*Escherichia coli* (*E. coli*) is a gram-negative bacillus and known to pose a serious threat to public health. Some of the *E. coli* strains can cause various intestinal or extraintestinal infections, severe foodborne diseases and even lead to death. [1] Currently, *E. coli* has been used as an indicator for routine analysis of the possible existence of pathogenic microbial species in contaminated water and food sources. [2–4] The standards set by the World Health Organization for drinking water and wastewater are 0 and 1000 CFU 100 mL<sup>-1</sup>, respectively. [5] For the detection of *E. coli*, traditional plate counting method is reliable but labor-intensive and time-consuming. [6] In order to meet the growing demand for sensitive and rapid bacteria detection, various technologies have been developed, such as enzyme-linked immune sorbent assay, [7] polymerase chain reaction, [8] cell counting via flow cytometry, [9] quartz crystal microbalance, [10] nanosensor [11] and so

on. While these detection methods are accurate and sensitive, each technique still has its own limitations. Therefore, the development of simple, sensitive, selective and reliable assays for *E. coli* detection is still drawing a fair amount of attention.

Potentiometric polymeric membrane sensors show advantages of small size, rapid response, low cost and resistance to color and turbidity interferences and are deemed as a promising tool for ion sensing. [12–16] In recent years, with the introduction of receptors such as aptamers, antibodies and molecularly imprinted polymers, potentiometric sensors provide a new means for the detection of bacterial cells. [17] Among these bioreceptors, peptide with unique features such as good affinity, high stability and versatile sequences is an ideal biorecognition element for potentiometric sensing of bacteria. [18,19] Recently, we designed a magneto-controlled extraction of peptide-modified magnetic beads onto/into polymeric membranes for the detection of bacteria. [20] The cooperative ion-pairing interactions between lipophilic ion-exchanger species in the membrane and the magnetic-field-assisted extraction of peptide lead to a non-equilibrium electromotive force response in a similar way to the polyion sensor. [21] Moreover, magnetic beads could eliminate the background interferences and al-

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**Table 1**  
Peptide designed for *E. coli* J96 in this work.

Peptide	Sequence <sup>a</sup>	Charge number <sup>b</sup>
Peptide-1R	biotin-LLLR	+1
Peptide-3R	biotin-LLLR	+3
Peptide-5R	biotin-LLLR	+5
Peptide-9R	biotin-LLLR	+9
Peptide-PR	biotin-LLLRPRPRPR	+6

<sup>a</sup> L, R, and P are glycine, arginine and proline, respectively.<sup>b</sup> Charge numbers were calculated by the NovoPro Bioscience Inc. (Shanghai, China).

low for sensitive and rapid detection of low colony-forming unit bacteria. Although the disposable potentiometric sensor array can achieve multiple bacteria classification and identification in a rapid way, there is significant room for the improvement of selectivity.

Herein, we report on a label-free potentiometric assay for specific detection of *E. coli* based on cleavage of peptide by outer-membrane protease T (OmpT). OmpT is a proteolytic enzyme on the outer membrane of *E. coli* and acts as the key virulence factor of *E. coli* degrading host immune responsibility-related proteins, cell matrix proteins, and interfering with host hemostatic function. [22] OmpT is known to cleave short peptides preferentially at dibasic sites [such as -R (arginine) -R-, -K (lysine) -K-, -K-R-, and -R-K-]. [23–25] Its pyrolysis efficiency was comparable to that of water-soluble proteases such as chymotrypsin. [26] In this work, taking advantage of the charge originated from the zwitterionic and amphipathic properties of peptide, a peptide serving as both OmpT substrate and potentiometric signal reporter was designed and immobilized on magnetic beads. Magnetic field-driven extraction of peptide-modified magnetic beads onto/into polymeric membranes ion-sensitive sensor leads to a rapid, stable and reproducible potential response. OmpT on *E. coli* is capable of cleaving the peptide on the magnetic beads, which can result in a potential change and be used for detection of *E. coli*. It will be shown that *E. coli* can be selectively detected by the proposed magneto-controlled potentiometric assay.

## 2. Experimental section

### 2.1. Materials

Dinonylnaphthalenesulfonate (DNNS), high molecular weight poly(vinyl chloride) (PVC), 2-nitrophenyl octyl ether (*o*-NPOE) and tetradodecylammonium tetrakis (4-chlorophenyl)borate (ETH 500) were purchased from Sigma-Aldrich. Streptavidin-modified magnetic beads (MBs, 1  $\mu$ m, 10 mg mL<sup>-1</sup>) were bought from BioMag Scientific Inc. (Wuxi, China). All the peptides (Table 1) were synthesized by GL Biochem Ltd. (Shanghai, China). *E. coli* BL21 and *E. coli* J96 ATCC® 700336™ were purchased from Bluebio. (Shanghai, China). *Staphylococcus aureus* was purchased from BIOBW Biotechnology Co., Ltd. (Beijing, China). The number of colony-forming units per mL (CFU mL<sup>-1</sup>) was determined by the surface plate counting method. Materials for bacteria culture were purchased from Qingdao Hopebio-Technology Co., Ltd (Qingdao, China). Tetrahydrofuran and other materials were obtained from Sinopharm Chemical Reagent Co., Ltd. Coastal seawater samples were collected from coastal area (Yantai, China, 121.60°E, 37.38°N). Seawater certified reference material for trace metals (NASS-6) was purchased from National Research Council Canada. Aqueous solutions were prepared with freshly deionized water (18.2 M $\Omega$ . cm specific resistance) obtained with a Synergy® UV water purification system.

### 2.2. Electrodes and membranes

The screen-printed ceramic electrodes with carbon ink as indicator electrodes substrate and Ag/AgCl ink as reference electrodes substrate were manufactured by Shenzhen Xinci Technology Co. LTD (Shenzhen, China). It should be noted that the thickness of carbon layers may have an influence on the potentiometric responses of the MBs under same magnetic field. The membrane cocktail solution containing 49 wt% PVC, 49 wt% *o*-NPOE, 1 wt% DNNS, 1 wt% ETH 500 was prepared as described in previous research. [20,27] For the fabrication of the electrodes, 10  $\mu$ L membrane cocktail was drop-casted on the indicator electrodes. The prepared electrodes were finally left to dry. The thickness of the sensing membrane was approximately 50  $\mu$ m. Membrane thicknesses were visually measured with a CX31–32C02 Olympus microscope (Tokyo, Japan).

### 2.3. Potentiometric measurement

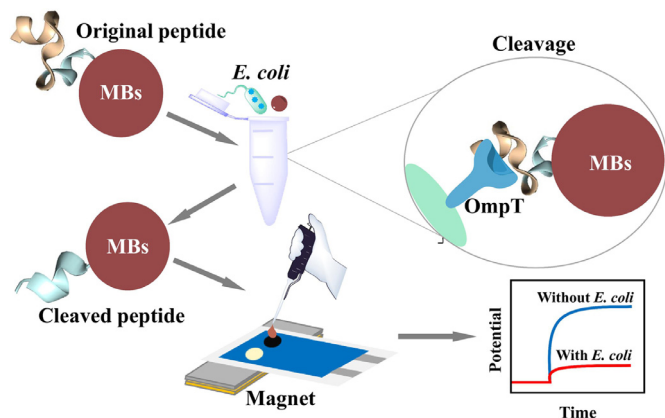
Potentiometric measurements were carried out using a Model PXSJ-226 digital ion analyzer (Shanghai INESA & Scientific Instrument CO. LTD, China) at room temperature in a custom-built cell with 1.0 mL phosphate buffered solutions (PBS, pH 7.4) containing 1.0 mM NaCl. The custom-built cell (40 mm  $\times$  40 mm  $\times$  23 mm) was made of polytetrafluoroethylene with the inner dimensions of 20 mm  $\times$  18 mm  $\times$  15 mm. The screen-printed ion-sensitive electrode (40 mm  $\times$  18 mm  $\times$  0.635 mm) was placed on the bottom of the cell (Fig. S1 in the supporting information). The magnetic field was controlled with the number of the applied magnets (surface magnetic field strength, 750 gauss; magnetic force, 3 kg; size, 40 mm  $\times$  20 mm  $\times$  2 mm).

### 2.4. Characterization of the polymeric membrane

The appearances of membranes without and with addition of MBs-peptide (magnetic beads modified with the peptide) under magnetic force were measured by field-emission scanning electron microscope (FESEM, S-4800, Hitachi, Japan). The elemental compositions of these membranes were analyzed by X-ray photoelectron spectroscopy (XPS, ESCALAB Xi<sup>+</sup>, Thermo Scientific, Japan). Fourier Transform Infrared Spectrometer (Nicolet iS50, Thermo Fisher, USA) was used to confirm the extraction of MBs-peptide into the polymeric membrane. Zeta potential measurements were carried out by Malvern Zetasizer Nano-ZS90 (ZEN3590, Malvern, UK). The molecular dynamics simulation was carried out using Discovery Studio 3.0 (Accelrys Software Inc.) as described in previous research. [28]

### 2.5. *E. coli* detection

The streptavidin-modified magnetic beads (MBs, 10 mg mL<sup>-1</sup>) were incubated with the biotin-labelled peptide (1.0  $\times$  10<sup>-5</sup> M) at the ratio of 1:9 for 30 min. After washing with PBS and magnetic separation, MBs-peptide was obtained. For detection of *E. coli*, MBs-peptide was incubated with *E. coli* at different concentrations for 1 h at 37 °C. The volumes of MBs-peptide and *E. coli* were at the ratio of 1:5. After incubation, washing and separation with magnetic field, 5  $\mu$ L of the incubated MBs-peptide was dropped on the indicator electrodes for potentiometric measurements. Before measurements, the electrodes were conditioned in 1.0 mM PBS containing 1.0 mM NaCl for 300 s to obtain a stable potentiometric response. Potential differences of MBs-peptide before and after incubation with *E. coli* were used for quantification of *E. coli*.

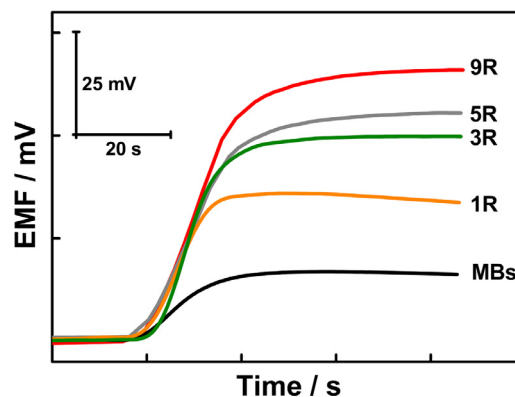


**Scheme 1.** Schematic illustration of the magneto-controlled potentiometric assay for *E. coli* sensing based on the cleavage of peptide on magnetic beads. MBs, streptavidin-modified magnetic beads.

### 3. Results and discussion

#### 3.1. Mechanism of the magneto-controlled potentiometric assay for *E. coli* detection

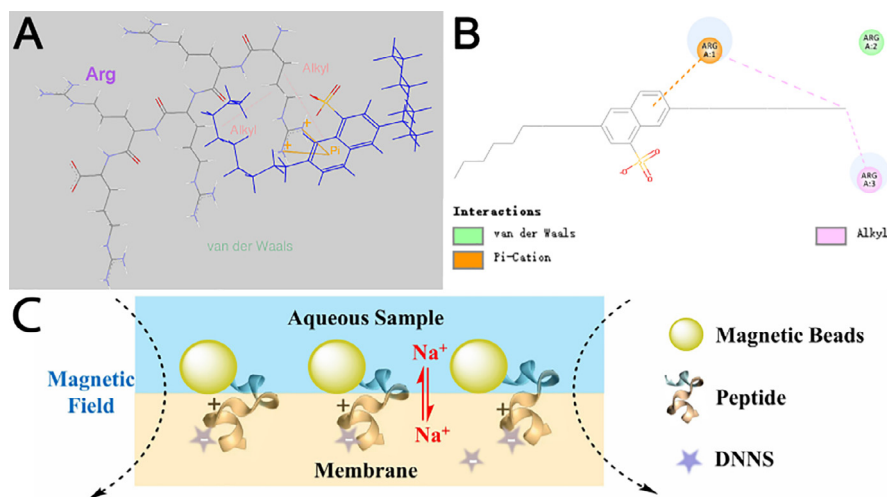
In order to achieve sensitive potentiometric measurements, the peptides that can be digested by OmpT were rationally designed. OmpT is known as a hydrolytic enzyme on the outer membrane of *E. coli* and can preferentially cleave peptide at dibasic sites (-R-R-, -K-K-, -K-R-, and -R-K-). [23] The pyrolysis efficiency of OmpT is comparable to water-soluble proteases. If proline (P) is located on the carboxyl side of the cleavage site, hydrolysis of the -R-P- will be weaker for protease such as trypsin. [26,29] Moreover, previous studies have shown that arginine (R) -rich protamine can induce a large and reproducible potential response on a polymeric membrane electrode doped with ion-exchanger. [27] Therefore, besides synthetic peptides with cleavage site (-R-R-) and different amounts of arginine (Table I), a synthetic peptide with proline located on the carboxyl side of the cleavage site was also designed. To immobilize the peptide on the magnetic beads via biotin-streptavidin interactions, biotin was added to the N-terminal of the sequences. In addition, three glycine acted as a spacer to impart chain flexibility were also added.



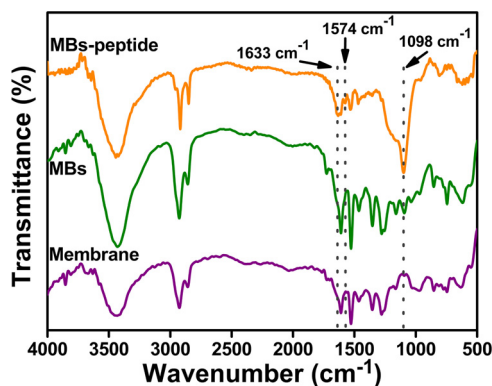
**Fig. 1.** Potentiometric response of ISE to MBs and MBs-peptide in 1.0 mL PBS (1.0 mM pH 7.4) containing 1.0 mM NaCl. 5  $\mu$ L of MBs and MBs-peptide (10 mg mL<sup>-1</sup>) was used. MBs, streptavidin-modified magnetic beads; 9R, 5R, 3R, and 1R represent MBs modified with peptide-9R, peptide-5R, peptide-3R, and peptide-1R, respectively. The arrow at 20 s represents the sample addition.

The sensing principle is illustrated in Scheme 1. External magnetic fields enhance the accumulation and extraction of hydrophilic peptide into the polymeric membrane via the formation of ion pairs with the ion exchanger (i.e., DNNS) in the membrane phase. As a result, phase boundary potential at the membrane sample/interface achieves a relatively significant and rapid change. In the presence of *E. coli*, OmpT on the surface of *E. coli* enables the cleavage of peptide at dibasic sites (-R-R-), which leads to a charge density change on the surface of MBs-peptide. Such changes of surface charge can be sensitively detected by polymeric membrane ion-sensitive sensors under magnetic field. The sensing protocol is label-free and can eliminate the procedure for bacteria lysis. Moreover, the peptide functionalized magnetic beads are capable of controlling the enzymatic hydrolysis process and subsequent potentiometric sensing.

Recently, we have shown that peptides immobilized on magnetic beads could be attached to or even extracted into polymeric membrane and lead to a rapid potential response in the presence of magnetic field. [20] To further investigate the magneto-controlled potentiometric sensing mechanism, it would be ideal to design peptides with the precise amino acids and charge density (Table I). As shown in Fig. 1, peptide-modified MBs can lead to



**Fig. 2.** (A) Molecular docking of DNNS (blue) to the receptor peptide-5R. (B) The interaction between DNNS and amino acids in peptide-5R. (C) Illustration of the response mechanism of MBs-peptide on the polymeric membrane ion-selective electrode. (For interpretation of the references to color in this figure legend, the reader is referred to the web version of this article.)



**Fig. 3.** FTIR spectra of bare membrane (purple), membrane dropped with MBs (green) and membrane dropped with MBs-peptide-9R (orange) under magnetic force. Membranes dropped with MBs and MBs-peptide-9R were washed with freshly deionized water and dried before characterization. (For interpretation of the references to color in this figure legend, the reader is referred to the web version of this article.)

a rapid, stable and reproducible potential response on the polymeric membrane ion-sensitive sensor. More importantly, with the amount of arginine in peptide increasing, the potentiometric responses of the electrode increase.

Molecular docking analysis reveals that the arginine in peptide can interact with the ion exchanger DNNS via van der Waals force,  $\pi$ -cation, and alkyl interactions (Fig. 2A, B). Moreover, the binding strength to DNNS can enhance with increasing the amount of arginine in the peptide (Related interaction forces and CDocker interaction energy are listed in Table S1). Therefore, peptides with higher charge density are more favorably extracted into the polymeric membrane to induce the ion exchange between the peptide in the sample solution and sodium ions in the membrane. The ion exchange process results in the potentiometric response (Fig. 2C). This is analogous to polymeric membrane-based potentiometric polyion sensors where the potential response is dependent on both the charge density and molecular weight. [30] However, a potential difference of ca. 38 mV was observed for the present system when measuring the same amount of MBs-peptide in 1.0 mM PBS containing 1.0 and 10 mM NaCl, respectively (Fig. S2). The potential difference is smaller than that predicted by the equilibrium potential response model (a Nernstian potential response for  $\text{Na}^+$ ). This is probably due to the high hydrophilicity of the peptide and the relatively weak interaction between the peptide and DNNS in the membrane phase. [20]

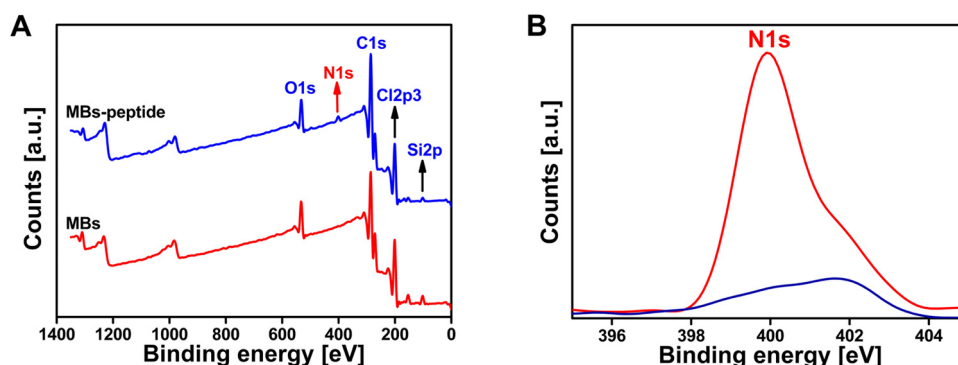
### 3.2. Confirmation of the ion-pair formation in membrane

As discussed in our previous communication, ion-pair formation between peptide on the MBs and ion exchanger within the polymer membranes leads to the potentiometric responses. [20] In order to verify this hypothesis, the infrared spectra of the membrane without and with addition of MBs-peptide under magnetic force were recorded. After dropping with MBs-peptide, strong washing with freshly deionized water and drying, the membrane shows three peaks at  $1098\text{ cm}^{-1}$  (C–O stretching),  $1633\text{ cm}^{-1}$  (C=N stretching) and  $1574\text{ cm}^{-1}$  (C=N bending). These peaks are characteristics of arginine in peptide, but were not obvious in the spectrum of the bare membrane and membrane dropped with MBs (Fig. 3). [31] Therefore, the lipophilic cation exchanger within the membrane phase can likely interact with the arginine in the peptide and leads to the accumulation and even extraction of the MBs-peptide into the polymeric membrane.

Extraction of peptide into the polymeric membrane were also characterized by X-ray photoelectron spectroscopy (XPS). Nitrogen 1s peak centered at  $400.2\text{ eV}$  is often assigned as amide bonds ( $-\text{CO}-\text{NH}-$ ) (Fig. 4). [32] The appearance of  $-\text{CO}-\text{NH}-$  component in arginine of peptide indicates the extraction of the peptide into membrane. Besides, field-emission scanning electron microscopy (FE-SEM) also shows that MBs-peptide can be adsorbed on the membrane (Fig. S3). All these results above indicate that peptide on the MBs at least partially can be extracted into the polymeric membrane in the presence of ion exchanger and magnetic force.

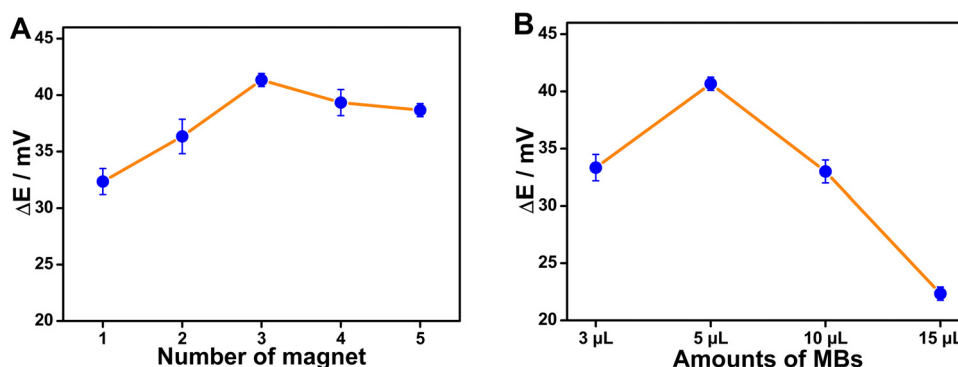
### 3.3. Optimization for potentiometric measurement of MBs-peptide

According to our previous research, the membrane containing PVC and *o*-NPOE in a weight ratio of 1:1, 1 wt% DNNS and 1 wt% ETH 500 was used for further experiments. [20] In order to increase the total electromotive force (EMF) response, experimental conditions including the magnetic field intensity, amount of MBs-peptide, and background solution were optimized. The potential difference between MBs-peptide and MBs was used for optimization. Magnetic field was controlled by the number of magnets. Since the layer thicknesses of aqueous diffusion could be modulated by the magnetic field, with the number of magnets increasing, potential change will increase (Fig. 5A). Then, it will decrease, which is due to the reduced ion-exchange process. The EMF response of the electrode to different amounts of MBs-peptide was investigated. As shown in Fig. 5B, potential difference decreased when the amount of MBs-peptide exceeds  $5\text{ }\mu\text{L}$  because of the high background signal originated from the extraction of the MBs. Therefore,  $5\text{ }\mu\text{L}$  MBs-peptide was used in this study. Moreover, a low background electrolyte (1.0 mM PBS, pH 7.4 containing 1.0 mM NaCl) was used to obtain a sensitive EMF response (Fig. S2).

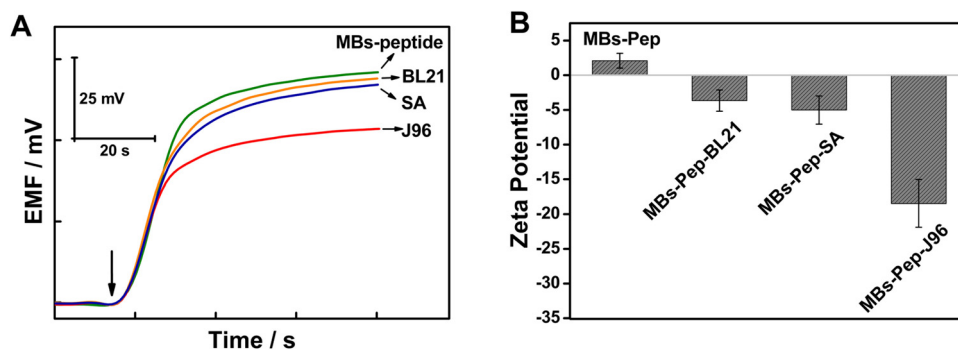


**Fig. 4.** XPS spectra of membranes dropped with MBs and MBs-peptide-9R under magnetic force: (A) survey scan; (B) XPS spectrum of nitrogen (N1s). Membranes dropped with MBs and MBs-peptide-9R were washed with freshly deionized water and dried before characterization.

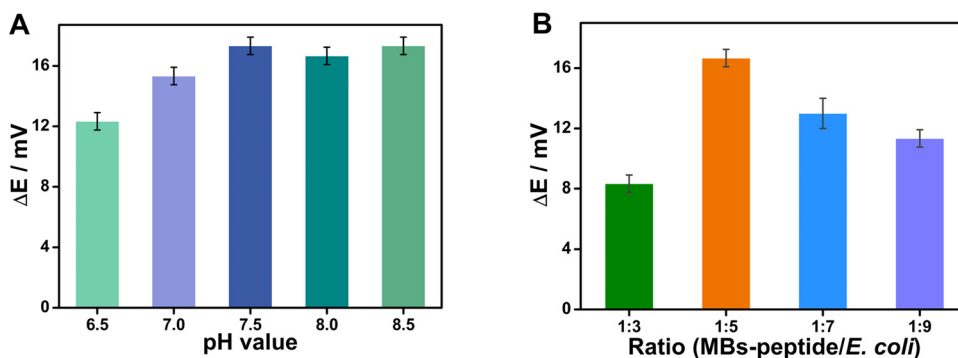




**Fig. 5.** Effects of (A) number of magnets and (B) amounts of MBs on the potential difference between MBs-peptide-9R and MBs. 5  $\mu$ L of MBs and MBs-peptide-9R (10 mg mL<sup>-1</sup>) were used. The potential difference between MBs-peptide and MBs was used for optimization. Error bars represent one standard deviation for three measurements.



**Fig. 6.** (A) Potential responses of MBs-peptide-9R and MBs-peptide-9R after incubation with  $1.0 \times 10^6$  CFU mL<sup>-1</sup> *E. coli* BL21 (BL21), *Staphylococcus aureus* (SA) and *E. coli* J96 (J96) at 37 °C for 1 h. After incubation, separation with magnetic field, and washing with PBS, MBs-peptide-9R was resuspended in 5  $\mu$ L PBS (1.0 mM pH 7.4) containing 1.0 mM NaCl. The arrow at 20 s represents the sample addition. (B) Zeta potentials of MBs-peptide-9R and MBs-peptide-9R after incubation with  $1.0 \times 10^6$  CFU mL<sup>-1</sup> *E. coli* BL21, SA and *E. coli* J96. Error bars represent one standard deviation for three measurements.



**Fig. 7.** Effects of (A) pH and (B) the ratio between volumes of MBs-peptide-9R and *E. coli* on the potential difference between MBs-peptide-9R and MBs-peptide-9R after incubation with  $1.0 \times 10^6$  CFU mL<sup>-1</sup> *E. coli* J96 at 37 °C for 1 h. After incubation, separation with magnetic field, and washing with PBS, MBs-peptide-9R was resuspended in 5  $\mu$ L PBS (1.0 mM pH 7.4) containing 1.0 mM NaCl. The potential differences of MBs-peptide-9R on the electrode before and after incubation with *E. coli* J96 were used to evaluate the cleavage efficiencies of OmpT. Error bars represent one standard deviation for three measurements.

### 3.4. Evaluation of enzyme digestion

OmpT is on the surface of all wild-type *E. coli* strains. [33] The pyrolysis efficiency of OmpT is known to be comparable to trypsin. [26] Our preliminary experiments have shown that the peptide on the magnetic beads can be digested in the presence of trypsin (Fig. S4). In this study, a strain of wild-type *E. coli* (*E. coli* J96) expressing OmpT at a high level was selected as a model. *E. coli* BL21 lacks the genes necessary to synthesize OmpT on its surface was used as a negative control. [33] To improve the sensitivity of the potentiometric assay, peptide-PR was initially designed as a substrate. However, compared with the potential change of MBs-peptide-9R on the electrode before and after digestion by OmpT, much lower

potential change of MBs-peptide-PR was observed (Fig. S5). Therefore, peptide-9R was selected for further experiments. As shown in Fig. 6A, the potential change of MBs-peptide-9R on the electrode was obvious (16 mV) after incubation with  $1.0 \times 10^6$  CFU mL<sup>-1</sup> *E. coli* J96. However, for the control strains such as *E. coli* BL21 and *Staphylococcus aureus* (*S. aureus*), a potential change of 3 mV was observed, which indicates that the OmpT plays a key role in the cleavage of peptide on the magnetic beads. The background potential change is probably due to the nonspecific interaction of the negatively charged bacteria.

Since the zeta potential analyzer can be used to conveniently measure the surface charge change, zeta potentials of MBs-peptide and MBs-peptide after incubation with *E. coli* J96 and control

**Table 2**  
Comparison of several enzyme-based techniques for *E. coli* detection.

Enzyme	Methods	LOD (CFU mL <sup>-1</sup> )	Assay Time (h)	Ref.
$\beta$ -Galactosidase	Colorimetry	10 <sup>4</sup>	2.5	34
$\beta$ -Galactosidase	Luminescence	40	8	35
$\beta$ -D-Galactosidase	Amperometry	10 <sup>5</sup>	1	36
$\beta$ -D-Glucuronidase	Fluorometry	10 <sup>5</sup>	2	37
$\beta$ -D-Glucuronidase	Amperometry	5 × 10 <sup>4</sup>	3	38
OmpT <sup>a</sup>	Fluorometry	10 <sup>5</sup>	1	33
OmpT	Potentiometry	5 × 10 <sup>3</sup>	1	This work

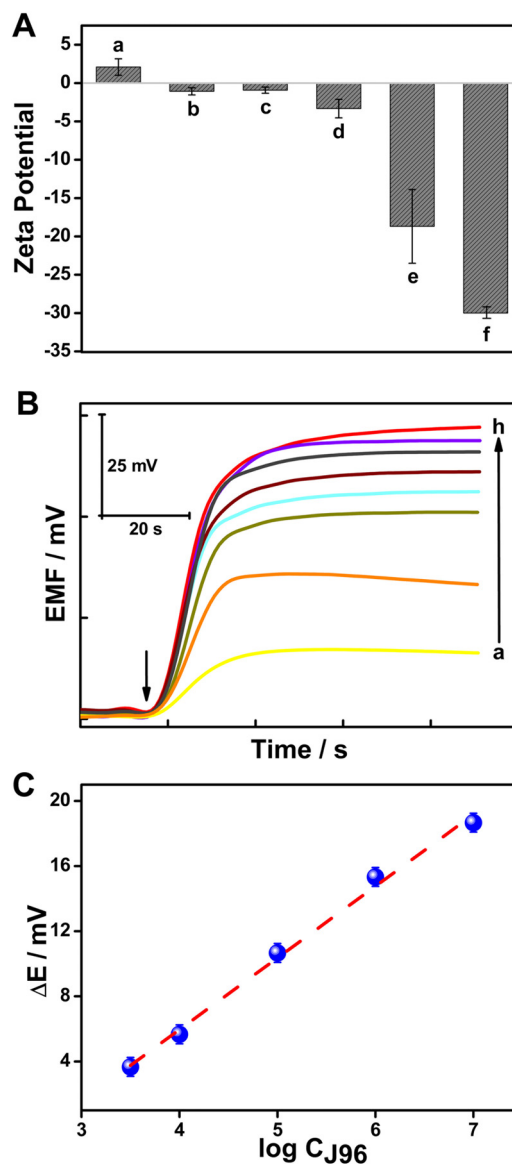
<sup>a</sup> OmpT, outer-membrane protease T.

strains (*E. coli* BL21 and *S. aureus*) were also investigated. As shown in Fig. 6B, zeta potential value of the MBs-peptide-9R changes from the initial positive value ( $2.09 \pm 1.07$  mV) to a negative value (e.g.,  $-18.47 \pm 3.42$  mV in the presence of  $1.0 \times 10^6$  CFU mL<sup>-1</sup> *E. coli* J96), which indicates the cleavage of peptide on magnetic beads. In contrast, nonspecific interaction of the control strains can only leads to a zeta potential decrease of ca. 5 mV. These results revealed the feasibility and selectivity for *E. coli* J96 sensing with peptide-9R modified magnetic beads.

### 3.5. Potentiometric detection of *E. coli*

In order to achieve sensitive potentiometric measurements, experimental parameters including pH and ratio between MBs-peptide-9R and *E. coli* were optimized. As shown in Fig. 7A, the cleavage efficiencies of OmpT improved with increasing pH values (from 6.5 to 8.5), and the maximum efficiency of OmpT was achieved at pH 7.5. This result is roughly consistent with a previous report. [25] Experiments also show that the maximum cleavage efficiency of OmpT was obtained by incubating MBs-peptide-9R with *E. coli* J96 ( $10^6$  CFU mL<sup>-1</sup>) at a ratio of 1:5 (Fig. 7B). Moreover, as a counterbalance of sensitivity and detection time, the incubation time of 1 h was used.

The proposed potentiometric *E. coli* assay was measured based on the surface charge change on the MBs-peptide. Therefore, the surface charge change of MBs-peptide can be quantified by the zeta potentials and be used for the *E. coli* detection. As shown in Fig. 8A, in the presence of *E. coli* J96, the surface positive charge of MBs-peptide-9R decreased, resulting in the decrease of zeta potential of MBs-peptide-9R. More importantly, as *E. coli* J96 concentration increases from  $1.0 \times 10^4$  to  $1.0 \times 10^7$  CFU mL<sup>-1</sup>, an increase in zeta potential change (from  $-0.92 \pm 0.40$  mV to  $-28.50 \pm 2.12$  mV) was observed. Indeed, by increasing the concentration of *E. coli* J96 cells, more peptides on the magnetic beads can be digested and thus reduce the surface charge or charge density, resulting in the steadily decrease of the potential response of the MBs-peptide-9R on the electrode. A linear relationship between the potential change and logarithmic value of *E. coli* J96 concentrations was found ranging from  $5.0 \times 10^3$  to  $1.0 \times 10^7$  CFU mL<sup>-1</sup>. The magneto-controlled potentiometric assay is able to detect the *E. coli* J96 at concentrations down to  $5.0 \times 10^3$  CFU mL<sup>-1</sup> (Fig. 8B, C). The detection limit of several similar techniques for *E. coli* strains are summarized and compared in Table 2. The proposed system is comparable with or even better than other enzyme-based assays. [33–38] More importantly, peptide functionalized magnetic beads offer the benefits of simplicity, speed, stable and flexibility to perform potentiometric detection based on surface charge change. The detection time is shorter than most commercially available detection kits for *E. coli*. [33,34,37,38] Note that, by prolonging the incubation time and using a pre-culture step, [33,39] the limit of detection can be further improved to satisfy the World Health Organization standard for *E. coli* content in drinking water. Since some other bacteria including *Shigella flexneri*, *Salmonella enterica* and *Yersinia pestis* can also express sim-



**Fig. 8.** (A) Zeta potentials of (a) MBs-peptide-9R and MBs-peptide-9R after incubation with (b)  $1.0 \times 10^3$ ; (c)  $1.0 \times 10^4$ ; (d)  $1.0 \times 10^5$ ; (e)  $1.0 \times 10^6$ ; (f)  $1.0 \times 10^7$  CFU mL<sup>-1</sup> *E. coli* J96 at 37 °C for 1 h. After incubation, separation with magnetic field, and washing with PBS, MBs-peptide-9R was resuspended in 500  $\mu$ L PBS (1.0 mM pH 7.4) containing 1.0 mM NaCl. (B) The potentiometric response of (a) MBs; (b) MBs-peptide-1R and MBs-peptide-9R after incubation with (c)  $1.0 \times 10^7$ ; (d)  $1.0 \times 10^6$ ; (e)  $1.0 \times 10^5$ ; (f)  $1.0 \times 10^4$ ; (g)  $5.0 \times 10^3$ ; (h) 0 CFU mL<sup>-1</sup> *E. coli* J96. The arrow at 20 s represents the sample addition. (C) Potential changes of MBs-peptide-9R after incubation with *E. coli* J96 at concentrations range of  $5.0 \times 10^3$ – $1.0 \times 10^7$  CFU mL<sup>-1</sup>. Error bars represent one standard deviation for three measurements.

ilar outer-membrane proteases, [40] the proposed method can be used to detect multiple bacteria by designing peptides with different cleavage sites. Compared to a number of serine proteases and metalloproteases, OmpT exhibits a remarkable tolerance of lysine/arginine methylation and acetylation. [25] Moreover, previous reports have shown that by changing the residues of the nearest-neighbor positions at the cleavage site, the catalytic efficiency of OmpT can be significantly improved. [41] Therefore, the substrate specificity of the OmpT might be further enhanced by changing the residue types or using methylated arginine.

To further investigate the feasibility of the magneto-controlled potentiometric assay for direct detection of *E. coli* cells in real samples, coastal seawater samples and standard seawater spiked with  $1.0 \times 10^6$  CFU mL<sup>-1</sup> *E. coli* J96 suspensions were tested. Compared to the standard samples ( $1.0 \times 10^6$  CFU mL<sup>-1</sup> *E. coli* J96 in PBS), a potential change decrease of 4 mV for standard seawater samples and 6 mV for coastal seawater samples were observed (data are not shown here). Some negatively charged species including anionic surfactants, negative charged biomolecules or organisms in real samples may electrostatically interact with positively-charged peptide and lead to the background signal. While both the sample matrix and interferences presented in sample may affect the performance of the assay, [23] the interferences can be eliminated by using a filtration system that facilitates separation and preconcentration of the target organism as described in our previous research. [39]

#### 4. Conclusions

In conclusion, a magneto-controlled potentiometric assay for *E. coli* J96 based on outer-membrane protease T (OmpT) have been developed. We demonstrated the feasibility of utilizing a magnetic force for the extraction of the peptide into an ion-exchanger-based polymeric membrane. The magnetic-field-assisted accumulation and extraction of the peptide into the membrane can induce an electromotive force response in a similar way to the polyion sensor. By making use of the OmpT on the surface of *E. coli* and the specific cleavage of peptide at the center of two pairs of arginine, *E. coli* can be detected accurately without any amplification or enrichment in 5000 CFU mL<sup>-1</sup> level with high selectivity. Owing to the simplicity, flexibility and versatility of the peptide functionalized magnetic beads, we envision that the magneto-controlled potentiometric assay holds great potential for rapid, sensitive and selective bacteria analysis.

#### Declaration of Competing Interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

#### Credit authorship contribution statement

**Han Zhang:** Methodology, Investigation, Data curation, Writing – original draft. **Junsong Mou:** Methodology, Software. **Jiawang Ding:** Conceptualization, Writing – review & editing, Supervision, Funding acquisition. **Wei Qin:** Supervision, Funding acquisition, Project administration.

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#### Supplementary materials

Supplementary material associated with this article can be found, in the online version, at doi:10.1016/j.electacta.2021.138408.

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