



Cite this: *RSC Adv.*, 2019, 9, 23658

# A spherical poly(acrylic acid) brush–enzyme block with high catalytic capacity for signal amplification in digital biological assays†

Yibei Wang,<sup>a</sup> Zehang Gao,<sup>bcd</sup> Jingwei Yi,<sup>a</sup> Hongbo Zhou,<sup>b</sup> Xiaoxia Fang,<sup>a</sup> Hong Xu,<sup>id</sup>\*<sup>a</sup> Jianlong Zhao<sup>b</sup> and Hongchen Gu<sup>id</sup>\*<sup>a</sup>

Ultrasensitive determination of some ultra-low abundance biological molecules closely related to diseases is currently a wide concern and urgent issue to be addressed. Here, a spherical poly(acrylic acid)–alkaline phosphatase (SP–AKP) signal amplification block using spherical poly(acrylic acid) brush nanoparticles (SP) as the immobilized carriers was designed and synthesized optimally first. The results show that a single SP–AKP with high enzyme binding capacity and high catalytic ability (up to about 4800 effective free AKP per SP–AKP) has much greater fluorescence signal amplification ability than a single free AKP or SiO<sub>2</sub>–COOH–AKP. Then, a droplet generation microfluidic chip was prepared successfully, and the SP–AKP was loaded and confined in a 14 pL droplet by adjusting its concentration to ensure at most one SP–AKP was encapsulated in each droplet according to Poisson's theory. Finally, the fluorescence signals produced by 4-methylumbelliferyl phosphate (4-MUP) catalyzed *via* SP–AKP within 6 min were sufficient to be detected by a fluorescence microscope. Thus, the digital signal distribution of “1/0” (signal/background) was obtained, making this SP–AKP signal amplification block a promising enzyme label for potential high sensitivity digital biological detection applications.

Received 6th May 2019  
 Accepted 20th July 2019

DOI: 10.1039/c9ra03404h

[rsc.li/rsc-advances](http://rsc.li/rsc-advances)

## Introduction

The enzyme-linked immunosorbent assay (ELISA) is a biological immunoassay technique widely used in clinical analysis and laboratory medicine.<sup>1</sup> It utilizes the specific binding of antigen–antibody and the high catalytic ability of enzymes to determine the target concentration rapidly by means of absorbance, fluorescence and chemiluminescence, *etc.* In recent years, it has been used in the determination of proteins, antigens, antibodies, such as Human Chorionic Gonadotropin (HCG),<sup>2</sup> insulin,<sup>3</sup> thyrotropin,<sup>4</sup> hepatitis B surface antigen (anti-HBV),<sup>5</sup> *Escherichia coli* enterotoxin,<sup>6</sup> HIV antibody (anti-HIV),<sup>7</sup> *etc.*

However, there are many kinds of biomarkers with extremely low abundance and great potential for clinical applications cannot be detected by current technology. They are like the part of a huge iceberg that sinks under the sea

(usually making up more than 80 percent<sup>8</sup> of the total volume), so few of them are available for us to know and utilize. For example, some patients with severe neurodegenerative diseases (such as Alzheimer's disease, Parkinson's disease, *etc.*) have specific “proteins” in their brains (like Aβ1-42, protein tau *etc.*<sup>9-11</sup>), due to the Blood–Brain Barrier (BBB), these specific “markers” find it difficult to penetrate the dense BBB into the blood, making their concentrations in the blood as low as femtogram or sub-picograms per milliliter. At present, the most sensitive ELISA detection technology is only at a level of picograms to tens of picograms per milliliter, which makes such “markers” with important clinical diagnostic and prognostic value unable to be applied in clinic.

In addition, digital detection as a single molecule biological detection method with the characteristics of small reaction volume and high throughput is emerging, which uses the microfluidic technology to encapsulate a single immune complex into a large quantity of separate microcavities<sup>9,12,13</sup> or microdroplets<sup>14-18</sup> for independent signals. Thus, absolute quantitative analysis of the biological samples can be achieved, especially suitable for the detection of ultra-low abundance target biomolecules. Compared with the microarray-based detection, microdroplet-based digital detection is more likely to have low detection limit and wide linear range owing to its infinite number of droplets, becoming a more promising digital detection method. However, the size of the droplet is relatively

<sup>a</sup>School of Biomedical Engineering, Med-X Research Institute, Shanghai Jiao Tong University, Shanghai 200030, PR China. E-mail: xuhong@sjtu.edu.cn; hcgu@sjtu.edu.cn

<sup>b</sup>Shanghai Institute of Microsystem and Information Technology, Chinese Academy Science, Shanghai 200030, PR China

<sup>c</sup>School of Information Science and Technology, ShanghaiTech University, Shanghai 201210, China

<sup>d</sup>University of Chinese Academy of Sciences, Beijing 100049, China

† Electronic supplementary information (ESI) available. See DOI: 10.1039/c9ra03404h



large (tens of microns) and the fluorescence signal from a single enzyme-catalysed droplet is difficult to detect. Therefore, it has become an urgent issue to improve the loading capacity and total catalytic activity of labelled enzyme in microdroplet immunoassay.

For that matter, efforts have been made by some researchers in adopting various nanoparticle materials as the immobilized carriers of enzymes to enhance the detection sensitivity by amplifying the detection signals with a large quantity of enzymes, such as gold nanoparticles (AuNPs),<sup>19</sup> polymer particles (PS microspheres),<sup>20</sup> mesoporous silica nanoparticles (MSN),<sup>21</sup> magnetic particles.<sup>22</sup> However, they have common problems of limited binding capacity and greatly decreased activity of immobilized enzymes. A spherical polyelectrolyte brush is a new kind of material formed when one end of linear polyelectrolyte chains densely attaches to spherical carriers such as nanoparticles.<sup>23–26</sup> It has high protein binding capacity, and maintains good biological activity due to the unique three-dimensional space structure formed by soft high-density polyelectrolyte chains grafted on surface.<sup>27–29</sup>

In this article, we established an effective fluorescence signal amplification enzyme label block based on the spherical poly(acrylic acid) brush (SP) for microdroplet digital detection. Scheme 1 shows the procedures of the protein immobilization and digital detection. First, alkaline phosphatase (AKP) was covalently immobilized into SP *via* a “chemical conjugation after electrostatic entrapment” (CCEE) method<sup>30</sup> to obtain the signal amplification block SP-AKP, and it reacted with the substrate 4-MUP to produce fluorescein 4-methylumbelliferone (4-MU) which emitted blue fluorescence excited by laser at 360 nm. Second, water-in-oil droplets with a diameter of 30  $\mu\text{m}$  were generated by flow focusing method (two oil phases, one substrate phase and one SP-AKP phase). Moreover, by controlling the concentration of the obtained SP-AKP, each droplet contained at most one SP-AKP particle. After reaction, the digital signals “1” and “0” (“1” for bright droplets that contain the SP-AKP, “0” for dark empty droplets) were observed by a fluorescence microscope and recorded by a CCD camera, and the concentration of the SP-AKP was obtained through

quantitative statistics and Poisson distribution formula. In addition, we also made comparative analysis of its catalytic activity to study the signal amplification ability of SP-AKP.

## Experimental section

### Materials

Alkaline phosphatase (AKP, from bovine intestinal mucosa) and 4-methylumbelliferyl phosphate (4-MUP) was purchased from Sigma-Aldrich. EDC (*N*-(3-dimethyl-aminopropyl)-*N'*-ethylcarbodiimide hydrochloride) was obtained from Aladdin Reagent (Shanghai) Co., Ltd. NHS (*N*-hydroxysuccinimide) was obtained from Thermo Scientific. All other reagents used in this work were the products of China National Medicines Group Shanghai Chemical Reagents Company.

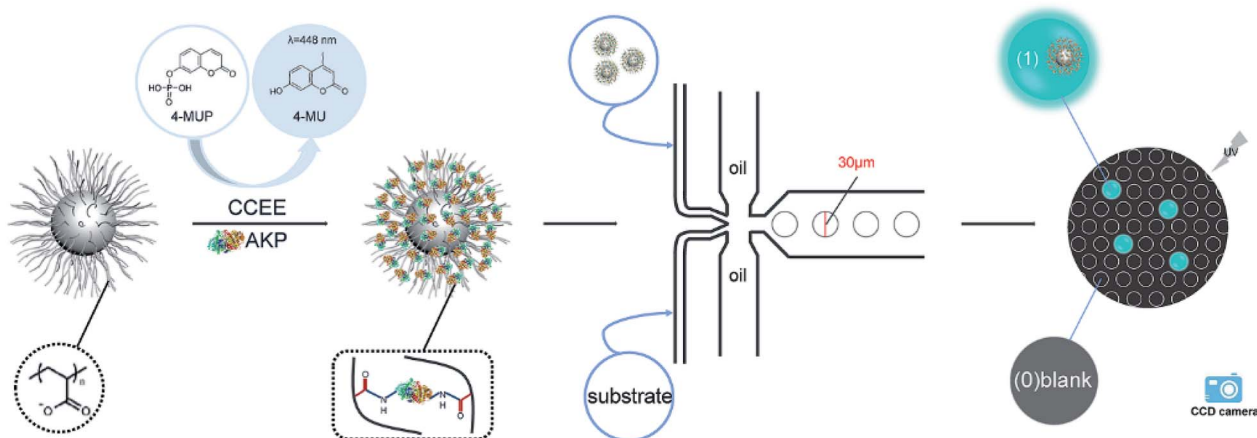
Tris(hydroxymethyl)aminomethane buffer (Tris-HCl, 0.1 M, pH = 9.0) and 2-(*N*-morpholino) ethanesulfonic acid buffer (MEST, 10 mM, pH = 5.0, containing 0.05 wt% Tween-20) were used in the protein binding experiments.

SP-300, the SP with high grafting density ( $\sigma = 0.42 \text{ nm}^{-2}$ ) of acrylic acid with a polymerization degree of 300 onto silica nanoparticles with a diameter of 116 nm, and conventional carboxylated  $\text{SiO}_2$  nanoparticles ( $\text{SiO}_2\text{-COOH}$ ) with a similar size were both used as immobilized enzyme carriers to evaluate their signal amplification abilities. The synthesis and characterization of SP-300 was done as described recently.<sup>31</sup> The preparation of  $\text{SiO}_2\text{-COOH}$  was conducted according to the literature.<sup>32</sup> Key structural parameters are listed and defined in Table 1 as follows.

The TEM images of SP-300 and SP-AKP were obtained by a 120 kV transmission electron microscope. The size distributions, PDI and zeta potential of SP-300 and SP-AKP were measured by Zetasizer Nano ZS *via* dynamic light scattering (DLS) in water.

### Protein immobilization

As conventional methods of coupling AKP on  $\text{SiO}_2\text{-COOH}$ , 400  $\mu\text{L}$  of 50  $\text{mg ml}^{-1}$  of NHS/EDC solution was added into each reaction system including 0.5  $\text{mg SiO}_2\text{-COOH}$  nanoparticles to



Scheme 1 Illustration of the protein immobilization and digital detection.



activate carboxyl to form NHS ester intermediates. Then 400  $\mu\text{l}$  AKP with different concentration gradients (3, 6, 9, 12, 15 and 18  $\text{mg ml}^{-1}$ ) were added to the reaction system respectively. The amino groups on the protein AKP attacked NHS intermediates to form amide bonds.

CCEE method was adopted to immobilize AKP into the SP, 400  $\mu\text{l}$  AKP with different concentration gradients (3, 6, 9, 12, 15 and 18  $\text{mg ml}^{-1}$ ) were first entrapped into 0.5 mg SP in each MEST reaction system *via* electrostatic interaction respectively. Then, the particle–enzyme complexes were washed two times by MEST before the conjugation with 300  $\mu\text{l}$  of 0.5 mM EDC in MEST to react at room temperature for 2 h on a rotary mixer. After conjugation, the particle–enzyme complexes were re-dispersed in 500  $\mu\text{l}$  Tris–HCl.

The protein concentration was quantified by BCA protein quantification method. The protein concentrations in the original protein solution, supernatant, and washing solution in each step were measured to obtain the binding capacity on the particles *via* subtraction method. The activity of free AKP, SP–AKP and  $\text{SiO}_2\text{–COOH–AKP}$  were acquired by microplate reader (SpectraMax\_i3). The relative activity of the enzyme was determined by kinetic fluorometry, a fluorescent value was detected every 30 seconds. The relative activity was expressed by the slope of the catalytic curve ( $\text{RFU s}^{-1}$ ).

### Droplets generation and signal detection

The droplets generation and detection platform were constructed, and divided into generation device and detection system. The generation device included PDMS chip, injection tubes/needles, vacuum pump and nitrogen pump, and the detection system included fluorescence microscope, mercury lamp, filter, high-speed monochrome CCD camera.

The droplets were generated by the flow focusing method (shown in Fig. S1a and b<sup>†</sup>). AKP/SP–AKP and substrate entered into the two dispersion phases respectively, and the continuous phase (mineral oil and surfactant EM90, Triton X-100) was sheared from both sides to obtain water-in-oil droplets. AKP/SP–AKP reacted with the substrate in droplets at room temperature.

The detection of droplets was mainly achieved by optical photography. Blue fluorescence signal of 448 nm was generated under near-ultraviolet light of a 130 W short-arc mercury lamp with 25% light power and a 380–400 nm cut-off filter. Fluorescence images were captured by the high-speed monochrome CCD camera under the fluorescence microscope in a very short time (exposure time of 100 ms and bandpass filter of 424–469 nm).

## Results and discussion

### Covalent protein immobilization on SP

Due to the large numbers of carboxyl functional groups on poly(acrylic acid) (PAA) chains, traditional NHS/EDC progress<sup>33</sup> seems to be a good choice for protein immobilizations in SP. But previous experiments have shown several drawbacks of the NHS/EDC process used in 3D materials like SP listed as follows:

(I) the inner sites of the SP could not be fully utilized due to the steric hindrance. Therefore, the protein binding capacity in covalent immobilization is only less than simple electrostatic adsorption. (II) it is possible that EDC/NHS activation reaction lead to the collapse of polymer chains from stretching state, which is because that the negative charge of carboxyl ionization on PAA chains would turn into a neutral NHS state, losing the advantage of three-dimensional structure of the SP. (III) NHS ester hydrolysis might bring some side reactions, resulting in a complex and uncontrollable coupling process.<sup>30</sup>

To solve above problems, a tailored–designed bio-conjugation method entitled “chemical conjugation after electrostatic entrapment” (CCEE) method was developed previously by our group<sup>30</sup> and was used in this study. First, the SP was synthesized by RAFT polymerization with a 116 nm  $\text{SiO}_2$  core and the chain length of the PAA was 92 nm. As a comparison, conventional carboxylated  $\text{SiO}_2$  nanoparticles ( $\text{SiO}_2\text{–COOH}$ ) with a similar core size were synthesized *via* modified Stöber method<sup>32</sup> and the surface carboxyl groups were obtained by grafting PAA polymers with average molecular weight of 5000 onto the surface of  $\text{SiO}_2$  nanoparticles. Key structural parameters of SP and  $\text{SiO}_2\text{–COOH}$  are listed in Table 1. As is reported, electrostatic adsorption is the main factor affecting the binding capacity during the immobilization process,<sup>30</sup> and AKP could be entrapped into the 3D structure of SP in MEST buffer of pH 5.0 *via* electrostatic interaction (the isoelectric point of SP and AKP are 4.0 and 5.7 respectively). Then, the obtained particle–protein intermediate complexes were separated by centrifugal and dispersed in MEST by ultrasonic to remove the excess unadsorbed proteins. Next, EDC was added to form amide bonds by covalently conjugating carboxyl groups on SP with residual amino groups of AKP enzyme. Finally, the SP–enzyme complexes (SP–AKP) were washed and dispersed in Tris–HCl buffer for further application.

Fig. 1a and b shows the TEM images of SP and the SP–AKP after covalent immobilization of AKP by CCEE method. The  $\text{SiO}_2$  core presented dark black can be observed clearly around with light grey PAA chains shell. When undergoing TEM characterization, the SP-300/SP–AKP particles were on copper grid in a dry state, and PAA chains grafted onto the silica nanoparticles collapsed into polymer shell, forming a certain gap between the SP particles; after coupling with AKP, the thickness of polymer shell around the  $\text{SiO}_2$  core increased distinctly, and the distance between the SP particles widened (shown in Fig. 1b and inset

Table 1 Comparison of structural parameters between SP-300 and  $\text{SiO}_2\text{–COOH}$ <sup>a</sup>

	$d^b$ (nm)	$D^c$ (nm)	$L^d$ (nm)	$N^e$ ( $\mu\text{mol g}^{-1}$ )
SP-300	116	300	92	4.425
$\text{SiO}_2\text{–COOH}$	121	211	—	2.513

<sup>a</sup>  $\text{SiO}_2\text{–COOH}$  with the same size as SP-300 was selected as an example. <sup>b</sup> Diameter of  $\text{SiO}_2$  core was determined by dynamic light scattering (DLS). <sup>c</sup> Diameter of  $\text{SiO}_2$  nanoparticles was determined by dynamic light scattering (DLS). <sup>d</sup> Lengths of group chains,  $L = (D - d)/2$ . <sup>e</sup> The amount of carboxyl was measured by conductometric titration.



images). This is due to a large number of enzyme molecules acting as cross-linking agents combined onto the PAA chains, preventing the chains from collapsing and exhibiting an increase in thickness in the dry state. It proves that the enzyme molecules were successfully immobilized into the inner parts of the SP, making full use of the sites on the PAA chains.

Fig. 1c shows the size distributions of SP-300 and SP-AKP particles by dynamic light scattering (DLS). They are approximately 300 nm in size with very low polydispersity index (PDI) in pure water (PDI < 0.08 and PDI < 0.02 respectively), indicating that the as-synthesized SP-300 and SP-AKP nanoparticles have monodisperse size with less aggregations. In the process of CCEE method, electrostatic adsorption is the dominant reaction,<sup>30</sup> the flexible chain segments can constantly adjust the binding positions of enzyme molecules to make full use of the sites and achieve efficient binding. In addition, Fig. 1d shows the state of SP-AKP in Tris-HCl buffer (pH 9.0) with excellent dispersion stability, which could be easily separated from the buffer after centrifugation for 6 min.

### Catalytic effect of SP-AKP

After adding substrate 4-MUP into SP-AKP ( $1 \text{ mg ml}^{-1}$ ), ultra-bright blue fluorescence could be observed by the naked when excited under 360 nm (shown in Fig. 1d), demonstrating that the AKP molecules on SP maintain high catalytic capacity.

According to the previous researches, the enzymes immobilized on the carriers would be partially inactivated due to deformation. In order to study the catalytic effect, we measured the catalytic activity of AKP and SP-AKP at different 4-MUP concentrations and studied the variation of catalytic rate with varying substrate concentration, which allowed us to determine the kinetic parameters for respective enzymes.

As shown in Fig. 2a and b, the catalytic rates of free AKP and SP-AKP first both increase linearly with the increasing substrate concentration, then tend to balance and reach a relatively stable rate. By fitting the data points with the Michaelis-Menten equation, the Michaelis-Menten constant ( $K_m$ ) of free AKP and SP-AKP are determined to be  $10.96 \mu\text{M}$  and  $16.84 \mu\text{M}$  respectively. Since their  $K_m$  values are relatively close, the substrate affinity of SP-AKP does not decline too much compared with free AKP. In addition, the turnover number ( $K_{\text{cat}} = V_{\text{max}}/[E]$ ,  $[E]$  is the concentration of AKP or SP-AKP) is  $6.91 \text{ s}^{-1}$  for AKP and  $2.11 \text{ s}^{-1}$  for SP-AKP respectively, thus, we can draw a conclusion that the relative activity of SP-AKP decrease to about 30.6% of free AKP.

### Establishment of signal amplification block based on SP-AKP

Since AKP on SP maintains relatively high catalytic activity and substrate affinity, the catalytic capacity of SP-AKP was studied in depth to prove the feasibility of the established fluorescent

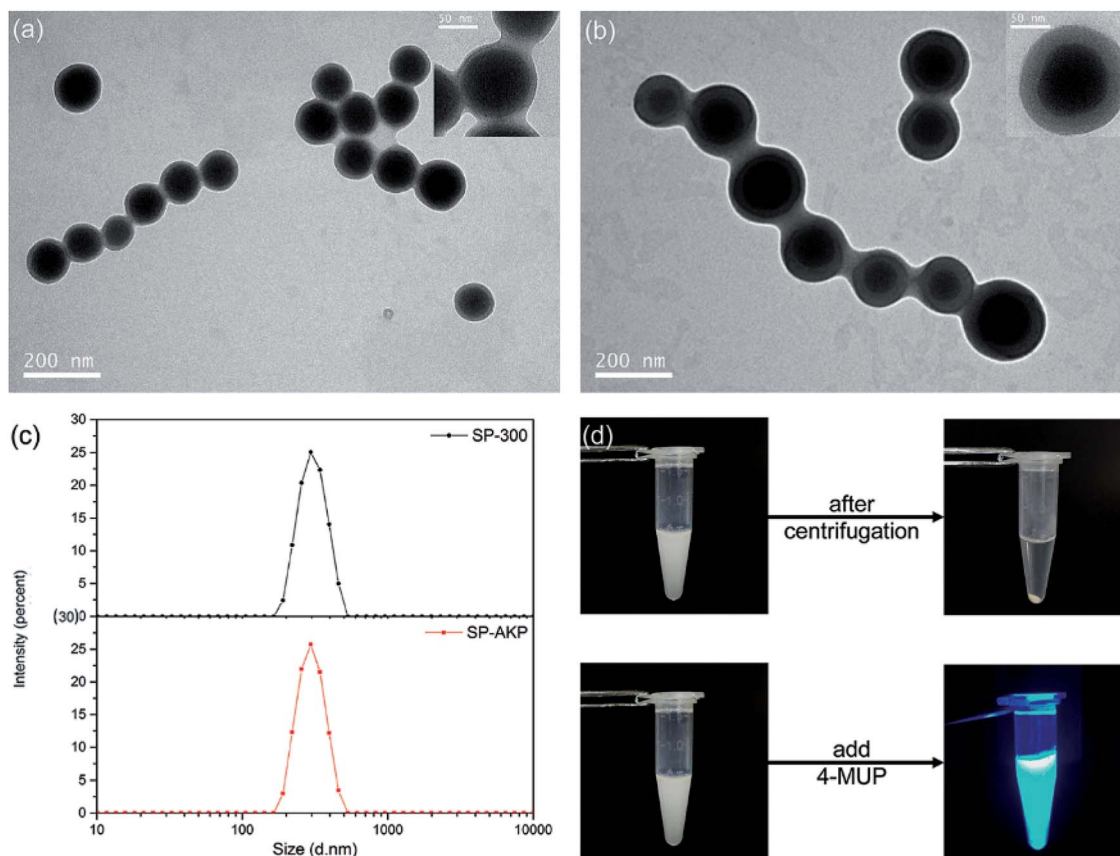


Fig. 1 TEM images of SP (a) before and (b) after covalent immobilization with AKP by CCEE (with a close-up of one particle at top left). (c) The corresponding DLS size distributions of particles SP-300 and SP-AKP. (d) Photos of the SP-AKP in aqueous suspension (left), after centrifugation and the fluorescent products (right).



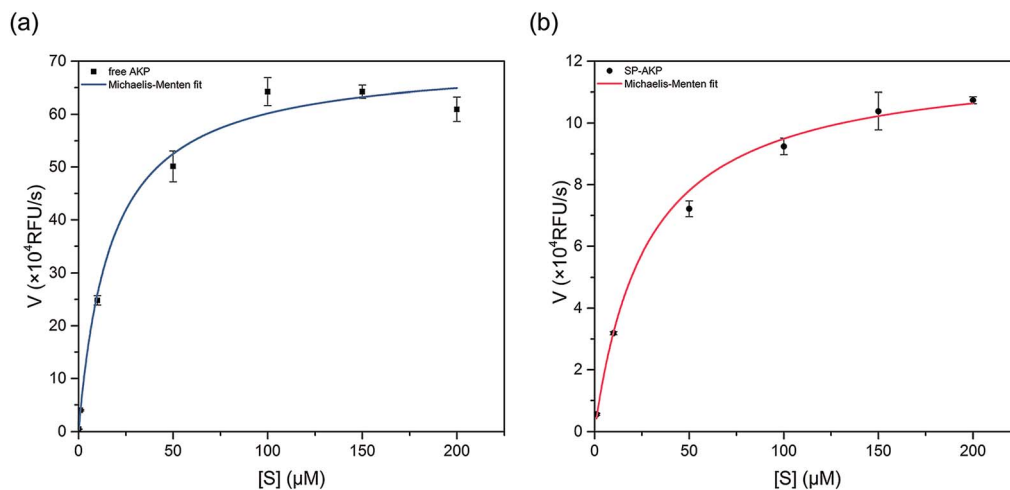


Fig. 2 Dependence of the catalytic rate of (a) free AKP and (b) SP-AKP on 4-MUP concentration. The activity was measured in the presence of 0.1 M Tris-HCl and 0.5 mM  $\text{MgCl}_2$  at pH 9.0. Data points were fitted to the Michaelis-Menten equation to give a  $K_m$  of free AKP and the SP-AKP of 10.96  $\mu\text{M}$  and 16.84  $\mu\text{M}$  respectively.

signal amplification enzyme label block. In the meantime, to further verify the signal amplification effect, we compared the SP-AKP with conventional  $\text{SiO}_2$ -COOH-AKP in a similar size in terms of protein binding capacity and relative activity. As shown in Fig. 3, the binding capacity of SP-AKP increases with the increasing added amount of AKP, and it can be well controlled. Besides, the relative total activity of SP-AKP and  $\text{SiO}_2$ -COOH-AKP both increases with the increasing binding capacity, and the maximum relative total activity are  $4.90 \times 10^5$  RFU  $\text{s}^{-1}$ ,  $4.57 \times 10^3$  RFU  $\text{s}^{-1}$ , respectively. Fig. 4b shows SP and the  $\text{SiO}_2$ -COOH can be combined with up to about 1800  $\mu\text{g}$  AKP per mg SP ( $\sim 16$  000 AKP per SP) and 800  $\mu\text{g}$  AKP per mg  $\text{SiO}_2$ -COOH ( $\sim 6$ 000 AKP per nanoparticle) respectively, on these conditions, the relative total activity of  $\text{SiO}_2$ -COOH-AKP is only 1% of SP-AKP. Furthermore, at the similar binding amount (794.24  $\mu\text{g}$   $\text{mg}^{-1}$  SP and 779.57  $\mu\text{g}$   $\text{mg}^{-1}$   $\text{SiO}_2$ -COOH), the relative total activity of  $\text{SiO}_2$ -COOH-AKP is about 50 times lower than SP-AKP. In conclusion, one single SP can combine a large number of AKP (up to about 4800 effective free AKP) and makes the total

activity of a single SP-AKP much higher than that of a single AKP, which means this SP-AKP has great potential to be a signal amplification enzyme label block.

The above results prove that the SP can immobilize a large number of enzymes and maintain higher enzyme activity than the traditional silica nanoparticles with hard surface, which endows the SP-AKP with a strong fluorescence signal amplification ability. This is because the hard surface structure of the conventional carboxylated  $\text{SiO}_2$  particles ( $\text{SiO}_2$ -COOH) limits the binding capacity of the enzyme to only one layer of enzymes on its surface, and the enzymes are prone to deformation. On the contrary, SP has a special 3D spatial structure, as well as soft long PAA chains with abundant carboxyl groups (shown in Fig. 4a), and thus the enzymes can maintain certain natural structures and catalytic activity. Circular dichroism (CD) of free AKP and SP-AKP further indicates the immobilized enzymes in SP maintains good protein structures (shown in Fig. S2†).

In general, fluorescence products catalysed by SP-AKP are usually dispersed in the solution. However, conventional bioassays that uses enzyme labels, such as ELISA, are typically reacted in large volume wells (*i.e.*,  $\sim 100$   $\mu\text{L}$ ), which means millions of enzymes at femto molar concentration can be detected through existing detection techniques. Therefore, when the abundance of biomolecules in the reaction system is extremely low, it is difficult to be detected in final system. In recent years, microdroplets produced by microfluidic chip technology characterized by small volume, fast production speed, non-diffusion of samples, cross-contamination avoiding between samples, stable reaction conditions and rapid mixing have been widely used in the biomedical fields such as DNA, protein, enzyme detection and drug delivery. Thanks to microfluidic technology, the reaction is confined to picoliter-volume droplets, and the fluorescent signals produced by a single target in a short time are possible to be detected by a fluorescence microscope as the concentration of fluorescent molecules can be much higher.<sup>34</sup>

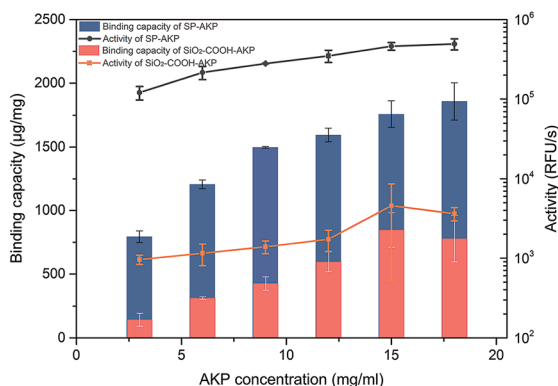


Fig. 3 Binding capacity and activity of SP-AKP and  $\text{SiO}_2$ -COOH-AKP and in dependence of concentration of AKP. Histogram shows binding capacity and line chart shows activity.



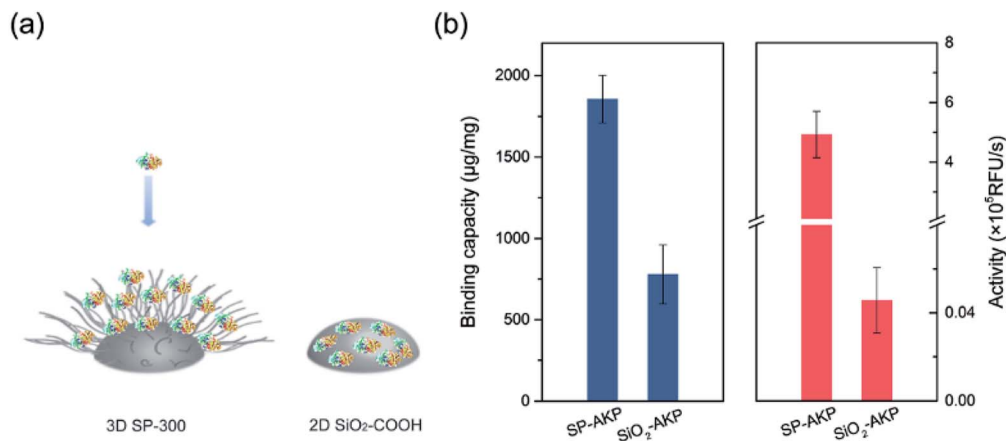


Fig. 4 (a) Comparison on protein immobilization mechanisms between 3D SP-300 and 2D SiO<sub>2</sub>-COOH. (b) Comparison of SP-AKP and SiO<sub>2</sub>-COOH-AKP in terms of protein binding capacity and activity.

Therefore, we constructed a microdroplet generation and detection platform suitable for the SP-AKP signal amplification block, which produced a large quantity of uniform and stable water-in-oil microdroplets. Single signal amplification block based on single SP-AKP nanoparticle could be separated from each other to form an independent microreactor to realize digital fluorescent signal expression. Thus, microdroplets were produced by flow focusing method. The substrate and enzyme entered from two different water dispersed phase channels at the same time, while the continuous oil phase (mineral oil and surfactant) turned around the cross channel, and squeezed two water dispersed phase to form droplets with the diameter of 30 μm. By keeping the concentration of substrate equivalent, the average number of SP-AKP in a single droplet can be changed by adjusting the concentration of SP-AKP. According to the Poisson distribution<sup>9,34</sup> (eqn (1),  $P(X)$ : the probability that there are exactly  $X$  SP-AKP nanoparticles per droplet,  $\lambda$ : the average number of SP-AKP nanoparticles per droplet), at very low concentration, most droplets have no SP-AKP, a few droplets

contain one SP-AKP, and very few ones contain more than one SP-AKP.

$$P(X) = \frac{\lambda^X}{X!} e^{-\lambda}. \quad (1)$$

Fig. 5 depicts the fluorescence images of free AKP ( $\lambda = 0.1$ , the average number of free AKP molecules per droplet) and SP-AKP ( $\lambda = 0.1$  in theory, the average number of SP-AKP particles per droplet). There are obvious and stable digital signal “1/0” in SP-AKP system, while no such signals in free AKP system even if the catalytic time was extended to 1 h (the grey value of all droplets in Fig. 5a are close to the background), and the signal-to-background ratio of a single SP-AKP is much higher than a single enzyme. It means that the fluorescence signals generated by a single enzyme could not be detected, while the SP-AKP, similarly as a single labelled building block, can produce sufficient fluorescence molecules in a 14 pL droplet to over the detection limit of CCD we used here in a very short time due to there are up to about 16 000 AKP molecules inside one SP-AKP

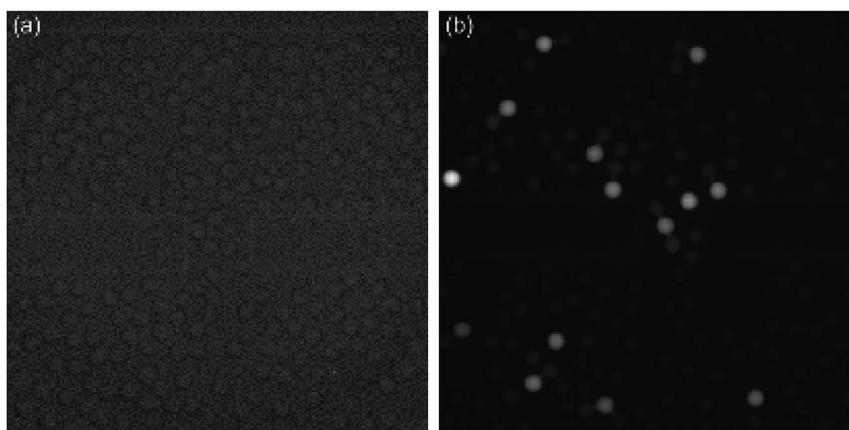


Fig. 5 The fluorescence images of (a) free AKP ( $\lambda = 0.1$ ) and (b) the SP-AKP ( $\lambda = 0.1$ ) in 14 pL droplets after a 6 min incubation with 0.25 mM 4-MUP. The exposure time was 100 ms. The brightness and contrast of two images were automatically adjusted by ImageJ software to clearly distinguish all droplets.



(corresponding to 4800 effective free AKP), demonstrating the necessity of signal amplification in droplet-based digital bioassay.

Moreover, we analysed the fluorescence intensity of signal “1/0” droplets, and statistics are shown in Fig. S3 and Table S1.† It is verified to satisfy the Poisson distribution. The total number of droplets is 6096,  $P(X = 0) = 91.04\%$ , and the actual  $\lambda$  is 0.0939 calculated by Poisson distribution formula which is almost consistent with theoretical feeding. Thus, according to this discrete probability distribution, we can obtain  $\lambda$  through  $P(X = 0)$ , and then the concentration of SP-AKP can be obtained according to the volume of droplet.

## Conclusions

In summary, a sensitive immunoassay signal amplification system (SP-AKP) based on the spherical poly(acrylic acid) brush (SP) has been developed by a CCEE method, with the characteristics of uniform, stable, good dispersion and easy separation. The results show that SP have higher enzyme binding capacity than traditional nanoparticles with hard surface, such as  $\text{SiO}_2\text{-COOH}$ , and the relative total activity of SP-AKP is 100 times higher than  $\text{SiO}_2\text{-COOH-AKP}$ . And then we built a microdroplet platform to separate the particle-enzyme block from each other, and obtained a considerable number of uniform droplets. The fluorescent signal distribution of “1/0” expressed in droplets further verifies the single SP-AKP with high catalytic capacity and remarkable fluorescence signal amplification ability in comparison to the single free AKP and  $\text{SiO}_2\text{-COOH-AKP}$ . These results indicate that the established SP-AKP signal amplification block is a promising label material for ultra-sensitive digital biological detection.

## Conflicts of interest

There are no conflicts to declare.

## Acknowledgements

This research is supported by the National Natural Science Foundation of China (Grant No. 21874091), and SJTU funding (Grant No. 17X10090003).

## Notes and references

- R. M. Lequin, *Clin. Chem.*, 2005, **51**, 2415–2418.
- H. T. Kuo, J. Z. Yeh, C. M. Jiang and M. C. Wu, *J. Immunol. Methods*, 2012, **381**, 32–40.
- M. J. Macdonald and J. P. Gapinski, *Metab., Clin. Exp.*, 1989, **38**, 450–452.
- A. E. Ohwovoriole and T. J. Wilkin, *J. Immunol. Methods*, 1988, **115**, 99–104.
- O. E. Tsitsilonis, A. Thrasyvoulides, A. Balafas, J. F. Voutsas, M. Papamichail and P. Lymberi, *J. Pharm. Biomed. Anal.*, 2004, **34**, 811–822.
- S. Shan, D. Liu, Q. Guo, S. Wu, R. Chen, K. Luo, L. Hu, Y. Xiong and W. Lai, *J. Dairy Sci.*, 2016, **99**, 7025–7032.
- G. Filice, L. Soldini, P. Orsolini, E. Razzini, R. Gulminetti, D. Campisi, L. Chiapparoli, E. Cattaneo and G. Achilli, *Microbiologica*, 1991, **14**, 185–194.
- Erenna SMC, [https://www.instrument.com.cn/netshow/SH101341/down\\_826643.htm](https://www.instrument.com.cn/netshow/SH101341/down_826643.htm), 2017.
- D. M. Rissin, D. H. Wilson and D. C. Duffy, *The Immunoassay Handbook*, 4th edn, 2013, vol. 1, ch. 4, pp. 223–242.
- M. J. Savage, K. Juliya, W. Abigail, T. Katherine, K. Rachel, C. M. Tanesha, J. W. Maxwell, N. G. Hatcher, S. J. Haugabook and W. Guoxin, *J. Neurosci.*, 2014, **34**, 2884–2897.
- R. Tarawneh, D. Head, S. Allison, V. Buckles, A. M. Fagan, J. H. Ladenson, J. C. Morris and D. M. Holtzman, *JAMA Neurol.*, 2015, **72**, 656–665.
- K. Leirs, P. Tewari Kumar, D. Decrop, E. Perez-Ruiz, P. Leblebici, B. Van Kelst, G. Compennolle, H. Meeuws, L. Van Wesenbeeck, O. Lagatie, L. Stuyver, A. Gils, J. Lammertyn and D. Spasic, *Anal. Chem.*, 2016, **88**, 8450–8458.
- L. Chang, D. M. Rissin, D. R. Fournier, T. Piech, P. P. Patel, D. H. Wilson and D. C. Duffy, *J. Immunol. Methods*, 2012, **378**, 102–115.
- M. L. S. Haakan and N. Joensson, *Angew. Chem.*, 2009, **121**, 2556–2559.
- L. Mazutis, J. Gilbert, W. L. Ung, D. A. Weitz, A. D. Griffiths and J. A. Heyman, *Nat. Protoc.*, 2013, **8**, 870–891.
- Z. Guan, Y. Zou, M. Zhang, J. Lv, H. Shen, P. Yang, H. Zhang, Z. Zhu and C. James Yang, *Biomicrofluidics*, 2014, **8**, 014110.
- A. Rakszewska, J. Tel, V. Chokkalingam and W. T. S. Huck, *NPG Asia Mater.*, 2014, **6**, e133.
- S. Tian, Z. Zhang, J. Chen, M. Du, Z. Li, H. Yang, X. Ji and Z. He, *Talanta*, 2018, **186**, 24–28.
- Y. Zhou, X. L. Tian, Y. S. Li, F. G. Pan, Y. Y. Zhang, J. H. Zhang, L. Yang, X. R. Wang, H. L. Ren and S. Y. Lu, *Biosens. Bioelectron.*, 2011, **26**, 3700–3704.
- Y. Li, F. Gao, W. Wei, J. B. Qu, G. H. Ma and W. Q. Zhou, *J. Mol. Catal. B: Enzym.*, 2010, **66**, 182–189.
- C. Lei, C. Xu, A. Nouwens and C. Yu, *J. Mater. Chem. B*, 2016, **4**, 4975–4979.
- M. Tudorache and C. Bala, *Sensors*, 2008, **8**, 7571–7580.
- M. Ballauff, *Prog. Polym. Sci.*, 2007, **32**, 1135–1151.
- A. Weiss, X. Guo and M. Ballauff, *Macromolecules*, 1999, **32**, 6043–6046.
- Q. de Robillard, X. Guo, M. Ballauff and T. Narayanan, *Macromolecules*, 2000, **33**, 9109–9114.
- X. Guo and M. Ballauff, *Phys. Rev. E: Stat., Nonlinear, Soft Matter Phys.*, 2001, **64**, 051406.
- A. Wittemann, B. Haupt and M. Ballauff, *Phys. Chem. Chem. Phys.*, 2003, **6**, 1671–1677.
- S. Rosenfeldt, A. Wittemann, M. Ballauff, E. Breininger, J. Bolze and N. Dingenouts, *Phys. Rev. E: Stat., Nonlinear, Soft Matter Phys.*, 2004, **70**, 061403.
- G. Jackler, A. Wittemann, M. Ballauff and C. Czeslik, *Spectroscopy*, 2004, **18**, 289–299.
- Z. Qu, K. Chen, H. Gu and H. Xu, *Bioconjugate Chem.*, 2014, **25**, 370–378.



- 31 Z. Qu, F. Hu, K. Chen, Z. Duan, H. Gu and H. Xu, *J. Colloid Interface Sci.*, 2013, **398**, 82–87.
- 32 W. Stöber, A. Fink and E. Bohn, *J. Colloid Interface Sci.*, 1968, **26**, 62–69.
- 33 S. Sam, L. Touahir, A. J. Salvador, P. Allongue, J. N. Chazalviel, A. C. Gougetlaemmel, d. V. C. Henry, A. Moraillon, F. Ozanam and N. Gabouze, *Langmuir*, 2010, **26**, 809–814.
- 34 D. M. Rissin and D. R. Walt, *Nano Lett.*, 2006, **6**, 520.

