

Research Paper

Colorimetric *in situ* assay of membrane-bound enzyme based on lipid bilayer inhibition of ion transport

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Abstract

Membrane-bound enzymes (MBEs), which make up a very high proportion of intracellular enzymes, catalyze a variety of activities that are currently analyzed by various techniques after purification. However, due to their amphipathic character, the purification of MBEs is difficult. Therefore, the most productive approach represents *in situ* analysis of MBEs in the cellular membrane.

Methods: In this study, using membrane-bound α -glucosidase (α -Glu) as an example, we have developed a colorimetric *in situ* assay for MBEs based on the inhibitory effect of lipid bilayer on ion transport. The enzyme substrate could mediate the self-assembly of phospholipid PEG derivative around magnetic nanospheres that were modified with boronic acid. The formation of lipid bilayer could inhibit the leaking of iron ions under acidic conditions. However, the product of the catalytic reaction had no capability for self-assembly of the lipid bilayer, leading to the release of iron ions from the magnetic nanospheres under acidic pH.

Results: The colorimetric *in situ* assay for MBEs could not only analyze the activity of membrane-bound α -Glu located on Caco-2 cells but could also evaluate the α -Glu inhibitors in cell medium.

Conclusions: The simple, economic, and efficient method proposed here offers a potential application for high-throughput testing of α -Glu and its inhibitors. Our study also outlines a strategy for exploring the colorimetric method to detect the activities of MBEs *in situ*.

Key words: membrane-bound enzyme, *in situ* analysis, lipid bilayer, ion transport, inhibitor evaluation

Introduction

Membrane-bound enzymes (MBEs) associated with the plasma membrane [1], endoplasmic reticulum [2, 3], membranes of mitochondria and chloroplasts [4, 5], and the membranes of other organelles [6] constitute a large proportion of intracellular enzymes. These enzymes catalyze a variety of activities including translocation [7] and information transfer [8] and act on locally concentrated substrates [9]. Importantly, MBEs are targets of over 50% of recent medical drugs [10-12]. Currently, the activities of MBEs are analyzed through purification [13, 14], followed by

determination with colorimetric [15, 16], fluorescent [17-19], chemiluminescent [20], and electrochemical techniques [21], as well as Western blotting [22].

In contrast to the soluble enzymes, MBEs are difficult to purify because of their amphipathic nature [23, 24]. For purification, MBEs are usually solubilized disrupting the integrity of the membrane and resulting in changes in stability, affinity, specificity for substrates and effectors, optimum pH, and other kinetic properties [23, 25]. Therefore, the most constructive approach is to analyze MBEs *in situ* in cellular membranes. Recently, fluorescent enzyme

substrates have been developed to label and test the activity of MBEs *in situ*. For instance, Wang et al. developed two novel fluorescent *in situ* targeting probes that specifically target γ -glutamyl-transpeptidase (GGT) in tumor cells and monitored the GGT activity in living cells [26]. However, it is rare for a colorimetric method to detect the activity of MBEs at the cellular level *in situ*.

The lipid bilayer is a thin polar membrane composed of two layers of lipid molecules. It is the barrier that prevents favorable ions and other substances from diffusing into other areas [27]. The bilayer is particularly impermeable to ions and allows cells to regulate the ion concentrations [28]. A variety of artificial lipid bilayers, such as liposomes and planer lipid bilayers, have been prepared and used to investigate ion translocation [29-31]. Also, black lipid membrane [32], solid lipid bilayer [33], hybrid lipid bilayer [34, 35], and polymer cushioned lipid bilayer [36] have been constructed to study the surface chemistry of the cells [37].

In this study, using α -glucosidase (α -Glu) as an example, we propose a new strategy for MBE analysis in cell medium. α -Glu is located in the brush border of the small intestine that acts on $\alpha(1\rightarrow4)$ bonds and its inhibitors show a broad spectrum of activity for the treatment of diabetes [38]. The enzyme can catalyze the hydrolysis of the substrate, para-aminophenyl- α -D-glucopyranoside [39], which forms stable complexes of cyclic esters with boronic acid groups [15, 40]. It is predicted that the substrate can assist the assembly of reactive phospholipid PEG derivatives in the vicinity of magnetic nanoparticle clusters modified by boronic acid and inhibit the leakage of iron ions into the solution and the subsequent catalyzed breakdown of hydrogen peroxide [41]. Thus, we designed a novel colorimetric strategy based on the substrate-mediated formation of the lipid bilayer and inhibition of ion transport for the analysis of MBEs. The slight change in the substrate can modulate the density of lipid bilayer around nanoparticles cluster and the subsequent release of iron ions. Also, compared with current colorimetric methods, which detect the concentrations of enzyme substrates or products in the complicated cell medium [42], the catalyzed property of iron ions confers high sensitivity to the *in situ* assay. Furthermore, the magnetic separation capability of the nanoparticles cluster allows for the removal of interfering substances thereby enhancing the analytical accuracy [43]. Importantly, this method for the analysis of MBE activity can be used in a high-throughput fashion with a simple colorimetric reader.

Materials and Methods

α -Glu (EC 3.2.1.20, from *Saccharomyces cerevisiae*), alkaline phosphatase (ALP), ovalbumin (Ova), sodium acetate, trisodium citrate, 3-aminophenylboronic acid (APBA), 1-(3-dimethylaminopropyl)-3-ethylcarbodiimide hydrochloride (EDC), 2,2'-azino-bis(3-ethylbenzothiazoline-6-sulphonic acid) diammonium salt (ABTS), N-hydroxysuccinimidepurum (NHS), N-2-hydroxyethylpiperazine-N'-2-ethanesulfonic acid (HEPES), 2-(N-morpholino)ethanesulfonic acid (MES), and trizma base were purchased from Sigma (Shanghai, China). para-Aminophenyl- α -D-glucopyranoside (pAPG) was acquired from Beijing Chemsynlab Pharmaceutical Co. Ltd (Beijing, China). 1,2-Distearoyl-sn-glycero-3-phosphoethanolamine (DSPE)-conjugated polyethylene Glycol 10000 and NHS (DSPE-PEG-NHS) were obtained from Ponsure Biotechnology Co. Ltd. (Shanghai, China). Vybrant® DiO cell-labeling solution and Cyanine 3 amine were purchased from Thermofisher (Shanghai, China). $\text{FeCl}_3 \cdot 6\text{H}_2\text{O}$, bovine serum albumin (BSA), α -amylase, lactoferrin (LF), lipase, trypsin, glucose oxidase (GO), hemoglobin, gallic acid (GA), quercetin (Que), maltopentaose (Mal), and ethylene glycol were obtained from Sinopharm Chemical Reagent Co. Ltd. (Shanghai, China). All other reagents used were of analytical grade. Caco-2 cells were acquired from Shanghai Genechem Co. Ltd. (Shanghai, China). All buffers and aqueous solutions were prepared with ultrapure water purified with a Millipore Milli-Q water purification system (Barnstead, USA) with a specific resistance of 18 M Ω cm. The solution of α -Glu with different concentrations was prepared in HEPES buffer (20 mM, pH 6.8).

Preparation of assembled magnetic nanospheres (AMNSs) and APBA/AMNSs

AMNSs were prepared according to the method reported previously [44]. APBA was immobilized onto the surface of AMNSs by using EDC as a linker [27]. Detailed experimental conditions have been described in Supplementary Material.

Construction and characterization of the lipid bilayer (LB)

20 μL of pAPG (15 μM) in HEPES buffer (20 mM, pH 7.4) was mixed with 20 μL of DSPE-PEG-NHS (500 μM) dissolved in 20 mM HEPES buffer (100 mM NaCl, 5 mM CaCl_2 , pH 7.4). The mixture was incubated for 50 min at room temperature (25 $^\circ\text{C}$), to yield DSPE-PEG-NHS/pAPG. Subsequently, 2.28 μL of NaOH (200 mM) was added to the mixture for adjusting the pH of the solution to 9.4. Next, 5 μL of 2 mg/mL APBA/AMNSs was added to the solution

and the resulting mixture was incubated for 13 min at room temperature with gentle shaking to form the stable complex through boronate ester bond between APBA and pAPG. Finally, the mixture was washed three times with HEPES buffer (20 mM, pH 9.4) to produce a lipid bilayer around AMNSs (LB/pAPG/APBA/AMNSs).

The lipid bilayer was further characterized by using laser scanning confocal fluorescence microscopy, fluorescence spectroscopy, dynamic light scattering, and Fourier transform infrared spectrometer. Detailed conditions have been described in Supplementary Material.

Detection of released iron ions

100 μ L of 100 mM MES solution was added to LB/pAPG/APBA/AMNSs with 15 μ M of pAPG concentration. Five minutes later, the supernatants resulting from magnetic separation were used for the detection of iron ions. The content of iron ions was measured using an inductively coupled plasma optical emission spectrometer (JY 2000-2, Horibia, France).

Analysis of enzyme activity

10 μ L of purified α -Glu solution (0.01-2.1 U/mL) or cell suspension (0-0.5 \times 10⁸ cells/mL) was added to 20 μ L of pAPG solution, and the resulting mixture was kept at 37 $^{\circ}$ C for 20 min. Subsequently, 1.5 μ L of NaOH (50 mM) was added to adjust the pH value to 7.4. Next, 20 μ L of DSPE-PEG-NHS (500 μ M) was added and after 50 min the mixture was washed three times followed by addition of 2.28 μ L of NaOH (200 mM) to adjust the pH value to 9.4. Then, 5 μ L of 2 mg/mL APBA/AMNSs solution was added. After 13 min, the reaction mixture was separated by magnet and washed three times to obtain LB/pAPG/APBA/AMNSs. Finally, 100 μ L of 100 mM MES buffer solution, 20 μ L of 5 mM ABTS, and 20 μ L of 4 mM H₂O₂ were added to LB/pAPG/APBA/AMNSs. Five minutes later, the supernatants resulting from magnetic separation were used for UV-Vis spectroscopic measurements, and the absorbance values were recorded with a UV-Vis spectrometer (Shimadzu Co., Kyoto, Japan).

Analysis of enzyme inhibitors

In the inhibition assay, 10 μ L gallic acid or quercetin solutions with different concentrations were premixed with purified α -Glu (10 μ L, 2.1 U/mL) or cell culture (10 μ L, 0.5 \times 10⁸ cells/mL) for 20 min at 37 $^{\circ}$ C. Next, 20 μ L of pAPG solution was added and the solution was incubated for 20 min at 37 $^{\circ}$ C. Subsequently, LB/pAPG/APBA/AMNSs were obtained and the absorbance values were recorded.

The inhibitory ratio was calculated as follows:

$$\text{Inhibitory ratio (\%)} = \frac{A_{414} - A_{414}^*}{A_{414}^0 - A_{414}^*} \times 100\%$$

where A_{414} was the absorbance obtained in the presence of enzyme and inhibitor, A_{414}^* was the absorbance obtained with enzyme only, and A_{414}^0 was the absorbance obtained without enzyme and inhibitor.

Results and Discussion

As α -Glu substrate, pAPG was composed of two moieties, i.e. phenyl ring and saccharide. The 2'-OH and 4'-OH in saccharide fraction of pAPG could react with boric acid on the surface of AMNSs, forming a stable six-membered ring of the complex (**Figure 1A**) [45]. As shown in **Figure 1B** (black curve), the spectrum of APBA/AMNSs exhibited two peaks at 1105 cm⁻¹ and 1016 cm⁻¹, which could be attributed to B-O-H stretching and deformation vibrations, respectively [46]. The two peaks disappeared for pAPG/APBA/AMNSs and the other two peaks at 1045 cm⁻¹ and 1180 cm⁻¹, corresponding to B-O-C stretching and deformation vibrations, appeared inversely (**Figure 1B**, red curve). Furthermore, the mean hydrodynamic diameter increased from 262.53 nm to 295.22 nm and the zeta potential decreased from -34.89 mV to -36.25 mV (**Figure 1C**). On the other hand, the primary amino group in phenyl ring fraction of pAPG could readily crosslink with NHS ester moiety of DSPE-PEG-NHS (**Figure 1A**) [47]. As shown in **Figure 1B** (red curve), the broad band around 3443 cm⁻¹ was assigned to the stretching vibration of -NH₂ group in pAPG fraction for pAPG/APBA/AMNSs [48]. For LB/pAPG/APBA/AMNSs, the band became narrow (**Figure 1B**, blue curve). Also, a strong peak at about 2917 cm⁻¹ appeared and belonged to -CH₂CH₂O- in the PEG fraction of DSPE-PEG-NHS [48]. Moreover, the negative charge increased from -36.25 mV to -41.58 mV after the linkage of DSPE-PEG-NHS (**Figure 1C**) due to the presence of phosphate groups in the structure of DSPE-PEG-NHS. As reactive phospholipid PEG derivative, DSPE-PEG-NHS could spontaneously form liposomes in the aqueous environment [49]. Therefore, it appeared that pAPG could assist the self-assembly of DSPE-PEG-NHS, leading to the formation of lipid bilayer surrounding of AMNSs (**Figure 1A**). As shown in **Figure 1C**, an approximate increase of 76 nm in the mean hydrodynamic diameter for LB/pAPG/APBA/AMNSs (371.47 nm) compared to that of pAPG/APBA/AMNSs (295.22 nm), indicated the coating of lipid bilayer [50].

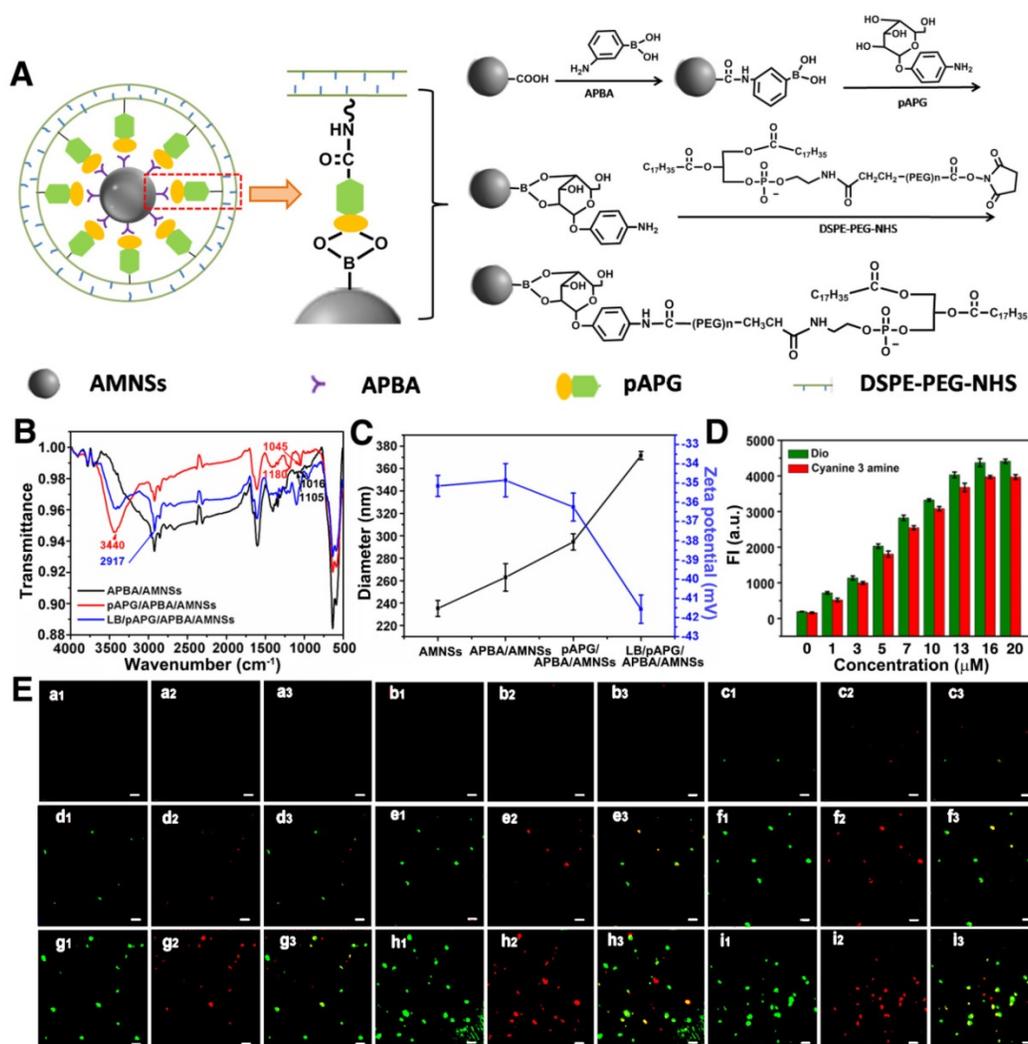


Figure 1. (A) Schematic illustration of the mechanism of lipid bilayer formation around AMNSs. (B) FT-IR spectra as well as (C) particle size distribution and zeta potential of AMNSs, APBA/AMNSs, pAPG/APBA/AMNSs, and LB/pAPG/APBA/AMNSs. (D) Fluorescence intensities of maximum emission peaks at 515 nm and 580 nm for LB/pAPG/APBA/AMNSs with different concentrations of pAPG (0, 1, 3, 5, 7, 10, 13, 16, and 20 μM) using Dio and Cyanine 3 amine as molecular probes, respectively. (E) Fluorescence images of LB/pAPG/APBA/AMNSs with different pAPG concentrations ((a) 0, (b) 1, (c) 3, (d) 5, (e) 7, (f) 10, (g) 13, (h) 16, (i) 20 μM) separately using Dio and Cyanine 3 amine as molecular probes: (1) green fluorescence, (2) red fluorescence, and (3) merged fluorescence.

To further validate the existence of lipid bilayer, two different fluorescent dyes, Dio and Cyanine 3 amine, were used for fluorescence image and fluorescent spectra measurement. As a long-chain carbocyanine dye, Dio is essentially insoluble in water and becomes inserted into the lipid bilayer with its alkyl chains and is widely used for the characterization of membranes [51]. Therefore, it was assumed that the formation of lipid bilayer would result in the exposure of NHS moiety in the molecular structure of DSPE-PEG-NHS. NHS could react with amino group of Cyanine 3 amine generating green fluorescence near AMNSs [52]. In the absence of pAPG, no fluorescence could be observed (Figure 1D-Ea) because of the lack of lipid bilayer close to APBA/AMNSs. With the increase in pAPG concentrations from 1 μM to 16 μM, the fluorescent intensities increased (Figure 1D) and the green and

red fluorescence gradually became brighter (Figure 1Eb1-h1 for green fluorescence and Figure 1Eb2-h2 for red fluorescence). This could be due to the incremental formation of lipid bilayer mediated by pAPG at the surface of AMNSs. However, fluorescence intensities were unchanged with further increase in the pAPG concentration from 16 μM to 20 μM (Figure 1D) indicating that AMNSs were totally covered by lipid. These results confirmed the formation of lipid bilayer around AMNSs.

It has been reported that under acidic conditions, iron ions could be leached from AMNSs [53]. pAPG-assisted lipid bilayer could protect AMNSs against acid leaching due to its inhibitory effect on the transport of hydrogen and iron ions [27] (Figure 2A). As displayed in Figure 2B, the concentration of iron ions was maximized for APBA/AMNSs, pAPG/APBA/AMNSs, and α-glucose/APBA/AMNSs,

indicating no inhibitory effect of APBA, pAPG, and α -glucose on the leakage of iron ions in the acidic solution. For LB/pAPG/APBA/AMNSs, the concentration of iron ions was unchanged in the absence pAPG compared with that for APBA/AMNSs, because of the absence of lipid bilayer around AMNSs. With the increase of pAPG concentration from 1 μ M to 20 μ M, the concentration of iron ions gradually decreased from 37.52 mg/L to 0.353 mg/L. This could be due to the increased self-assembly of DSPE-PEG-NHS and the formation of lipid bilayer which could not only prevent the transport of hydrogen ions to inhibit the leaching of iron ions, but also hinder the release of leached iron ions from the outer layer of AMNSs into the solution.

The mechanism for α -Glu assay is illustrated in **Figure 3A**. The enzyme could catalyze the breakage of α -1,4 glycosidic bond, resulting in the hydrolysis of pAPG into pAP and α -glucose [39]. The α -glucose consisting of the *cis* 2,4-diols group, could interact with boronic acid group at the outer layer of APBA/AMNSs forming a stable six-membered ring of the complex. After magnetic separation, α -glucose/APBA/AMNSs obtained could release iron ions into the solution under acidic pH, because of the no barrier effect around the AMNSs. In the presence of H_2O_2 , the iron ions could catalyze the oxidation of ABTS to the colored product (ABTS^{•+}) with a maximum absorbance peak at 414 nm producing the dark green supernatant [41, 54]. On the contrary, without α -Glu, pAPG as a linker could mediate the self-assembly of DSPE-PEG-NHS and subsequent formation of lipid bilayer around the AMNSs which was confirmed by the above experiment (**Figure 1**). At acidic pH, the lipid bilayer could inhibit the accumulation of hydrogen ions onto the surface of AMNSs and

prevent the leaching of iron ions (**Figure 2**). In this case, ABTS could not be oxidized due to the deficiency of iron ions in the H_2O_2 solution and a weak absorbance and light green supernatant liquid could be observed. Thus, α -Glu activity was closely related to the concentration of iron ions in the solution and a new method for enzyme analysis based on inhibition of ion transport in the lipid bilayer could be developed. Considering specific cleavage of the substrate by the enzyme and magnetic separation property of AMNSs, we could analyze enzyme activity specifically, accurately, and efficiently.

To validate the feasibility of our proposed method for enzyme activity analysis, a series of experiments were conducted and the results are presented in **Figure 3B**. In the presence of H_2O_2 , ABTS could be mildly oxidized resulting in a low absorbance peak at 414 nm (**Figure 3B**, blank curve) and a nearly colorless reaction solution (vial 1). pAPG, α -Glu, and the mixture of both had a negligible effect on the oxidation of ABTS as there was almost no change in the absorbance value (**Figure 3B**, red, blue, and green curves) or the color of solution (vials 2, 3, and 4). However, addition of iron ions led to a high absorbance value (**Figure 3B**, pink curve) and dark green solution (vial 5) that could be attributed to the catalyzing effect of iron ions on the oxidation of ABTS by H_2O_2 . pAPG and the enzyme had nearly no impact on the catalyzed oxidation of iron ions as no change in absorbance values (**Figure 3B**, yellow and navy curves) and the color of the solution (vials 6 and 7) was observed. Like iron ions, addition of APBA/AMNSs and pAPG/APBA/AMNSs yielded high absorbance value (**Figure 3B**, dark red and rose red curves) and dark green solution (vials 8 and 9) indicating the oxidation effect catalyzed by the two

composites due to the leaching of iron ions from APBA/AMNSs and pAPG/APBA/AMNSs under acidic pH. As shown in **Figure 2B**, iron ions could be separately leached from APBA/AMNSs and pAPG/APBA/AMNSs into the solution. Addition of LB/pAPG/APBA/AMNSs resulted in a low absorbance value (**Figure 3B**, pale green curve) and light solution color (vial 10), implying that it decreased the number of iron ions in the solution. The phenomena could be explained by the formation of lipid bilayer originating from the self-assembly of DSPE-PEG-NHS. On the one hand, the lipid bilayer could inhibit the permeation of hydrogen ions into AMNSs to prevent the leaching of iron ions [27, 28].

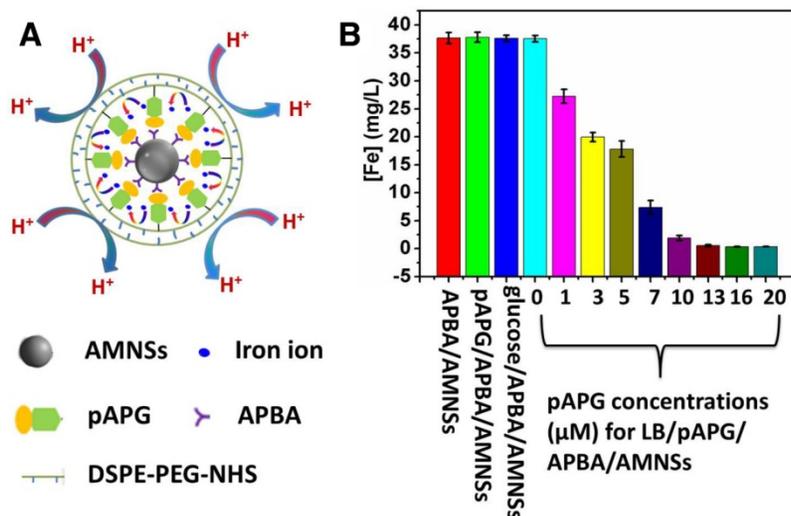


Figure 2. (A) Schematic illustration for the release of iron ions inhibited by the pAPG-mediated lipid bilayer. (B) The release of iron ions for APBA/AMNSs, pAPG/APBA/AMNSs, glucose/APBA/AMNSs, and LB/pAPG/APBA/AMNSs with different concentrations of pAPG (0, 1, 3, 5, 7, 10, 13, 16, and 20 μ M).

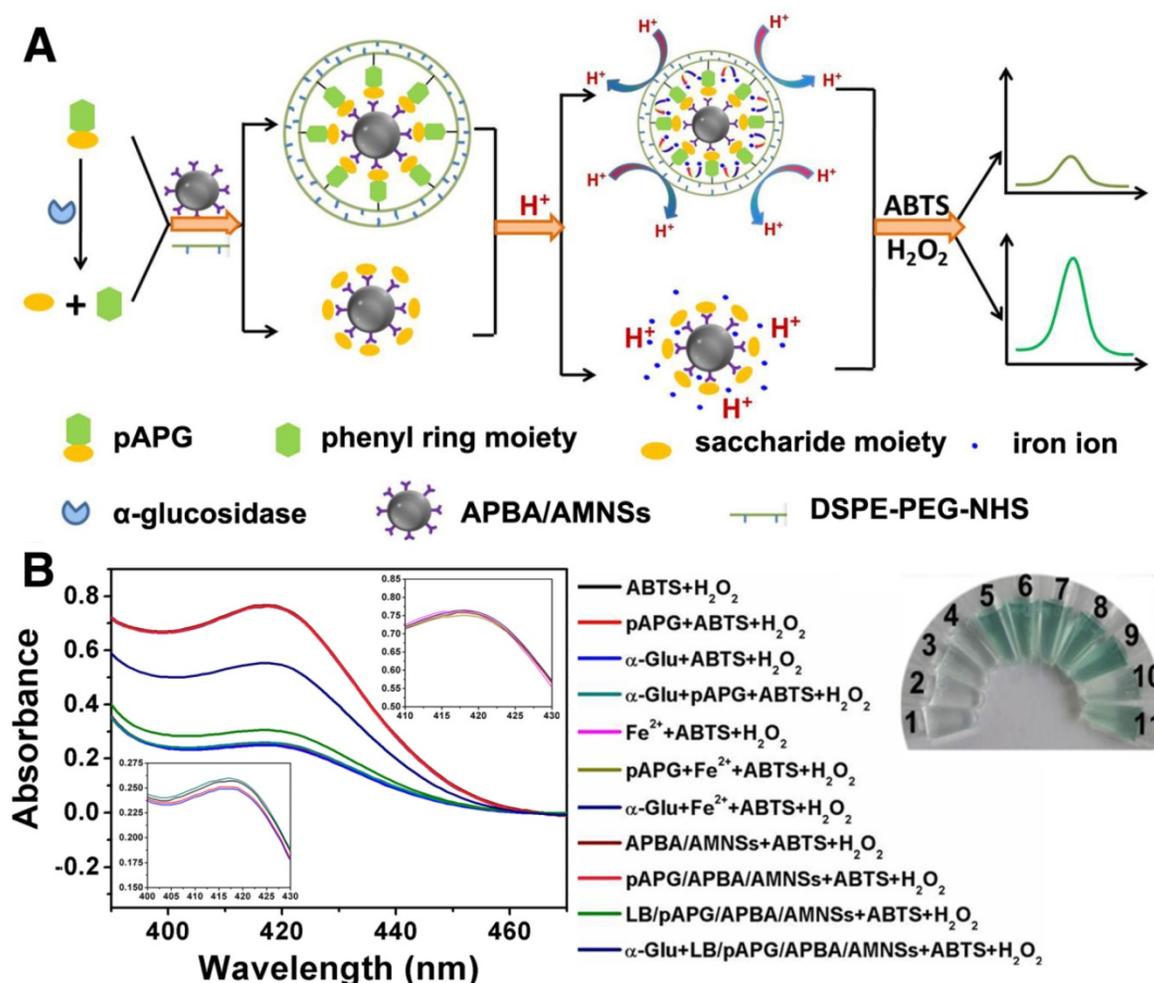


Figure 3. (A) Schematic illustration for the mechanism of α -Glu assay via inhibition of lipid bilayer on iron release. (B) UV-vis spectra of the mixtures prepared by separate addition of ABTS (20 μ L, 5 mM) + H₂O₂ (20 μ L, 4 mM) (vial 1, black curve), pAPG (20 μ L, 20 μ M) + ABTS (20 μ L, 5 mM) + H₂O₂ (20 μ L, 4 mM) (vial 2, red curve), α -Glu (10 μ L, 1 U/mL) + ABTS (20 μ L, 5 mM) + H₂O₂ (20 μ L, 4 mM) (vial 3, blue curve), α -Glu (10 μ L, 1 U/mL) + pAPG (20 μ L, 20 μ M) + ABTS (20 μ L, 5 mM) + H₂O₂ (20 μ L, 4 mM) (vial 4, green curve), Fe²⁺ (5 μ L, 25 mM) + ABTS (20 μ L, 5 mM) + H₂O₂ (20 μ L, 4 mM) (vial 5, pink curve), pAPG (20 μ L, 20 μ M) + Fe²⁺ (5 μ L, 25 mM) + ABTS (20 μ L, 5 mM) + H₂O₂ (20 μ L, 4 mM) (vial 6, yellow curve), α -Glu (10 μ L, 1 U/mL) + Fe²⁺ (5 μ L, 25 mM) + ABTS (20 μ L, 5 mM) + H₂O₂ (20 μ L, 4 mM) (vial 7, navy curve), APBA/AMNSs + ABTS (20 μ L, 5 mM) + H₂O₂ (20 μ L, 4 mM) (vial 8, wine curve), pAPG/APBA/AMNSs + ABTS (20 μ L, 5 mM) + H₂O₂ (20 μ L, 4 mM) (vial 9, Rose red curve), LB/pAPG/APBA/AMNSs + ABTS (20 μ L, 5 mM) + H₂O₂ (20 μ L, 4 mM) (vial 10, palegreen curve), and LB/pAPG/APBA/AMNSs + α -Glu (10 μ L, 1 U/mL) + ABTS (20 μ L, 5 mM) + H₂O₂ (20 μ L, 4 mM) (vial 11, dark blue curve) into a tube with the final volume of 140 μ L by adding 100 mM MES buffer solution. Inset: the corresponding magnified spectra.

On the other hand, the bilayer also hindered the release of iron ions into the solution [27, 28]. α -Glu cleaved pAPG into pAP and α -glucose [39], increasing the absorbance value (Figure 3B, dark blue curve) and yielding a relatively dark solution color (vial 11). Different from pAPG, the hydrolyzed products (pAP and α -glucose) could not be used as linkers to assist the self-assembly of DSPE-PEG-NHS and the eventual formation of lipid bilayer around AMNSs. Conversely, after magnetic separation, α -glucose/APBA/AMNSs obtained could release iron ions in the acid solution (Figure 2B).

As a membrane-bound enzyme in the epithelium of the small intestine, α -Glu catalyzes the final step of carbohydrate hydrolysis to produce absorbable monosaccharides which determine the postprandial blood level [38]. Therefore, the inhibitors can be accurately analyzed using the membrane-bound

enzyme in cell medium. The human intestinal Caco-2 cell line has been extensively used as a model of the absorptive and defensive properties of the intestinal mucosa [55]. We further confirmed that the established method could be applied for the analysis of membrane-bound enzyme and the screening of inhibitors by using Caco-2 cell. Consistent with the observed increase in absorbance values, color of the solution turned from colorless to dark green upon addition of 0 to 0.5×10^8 cells/mL (Figure 4A). The experiment verified the progressive cleavage of pAPG catalyzed by the membrane-bound α -Glu located on Caco-2 cells, decreasing the lipid bilayer around AMNSs and increasing the concentration of released iron ions. The results indicated that the method can be utilized for the analysis of membrane-bound enzyme in cell medium.

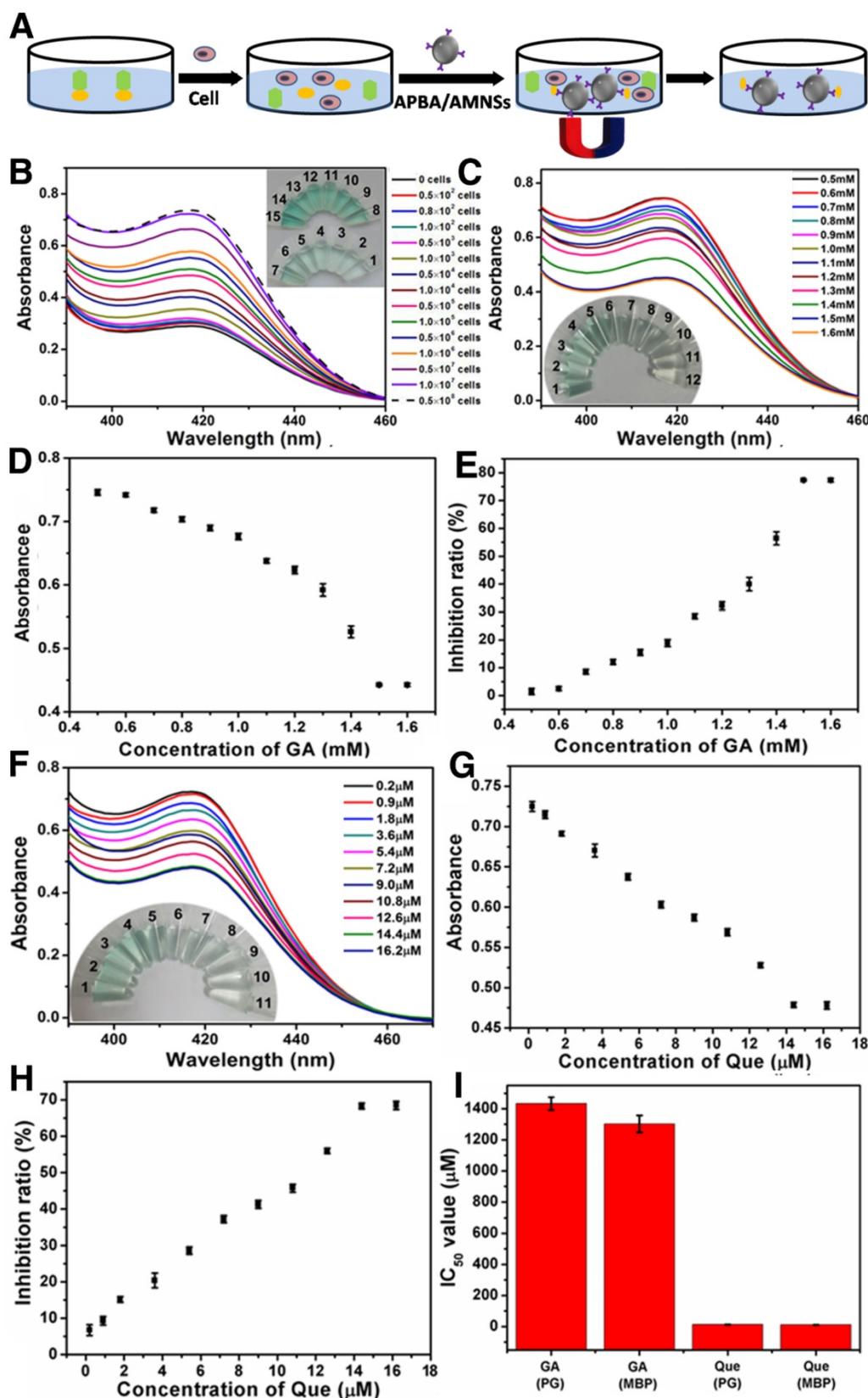


Figure 4. (A) UV-vis spectra and photographs (inset) of the mixtures prepared by the addition of various numbers of Caco-2 cells. Inset, from vial 1 to vial 15: the cell numbers are 0, 0.5×10^2 , 0.8×10^2 , 1.0×10^2 , 0.5×10^3 , 1.0×10^3 , 0.5×10^4 , 1.0×10^4 , 0.5×10^5 , 1.0×10^5 , 0.5×10^6 , 1.0×10^6 , 0.5×10^7 , 1.0×10^7 , and 0.5×10^8 cells/mL. (B) UV-vis spectra and photographs (inset) of mixtures, (C) absorbance values, and (D) inhibition ratios following the addition of various concentrations of GA to Caco-2 cells. Inset (B), vial 1 to vial 9: the GA concentrations are 0.5, 0.6, 0.7, 0.8, 0.9, 1, 1.1, 1.2, 1.3, 1.4, 1.5, and 1.6 mM. (E) UV-vis spectra and photographs (inset) of mixtures, (F) absorbance values, and (G) inhibition ratios following the addition of various concentrations of Que to Caco-2 cells. Inset (E), vial 1 to vial 9: the Que concentrations are 0.2, 0.9, 1.8, 3.6, 5.4, 7.2, 9, 10.8, 12.6, 14.4, and 16.2 μ M. (H) IC_{50} values of GA and Que on purified α -Glu (PG) and membrane-bound α -Glu (MBG). Cell numbers are 5×10^7 cells/mL. Error bars indicate standard deviations ($n = 3$).

We further evaluated the efficiencies of inhibitors in cell medium. Changes in absorbance spectra, absorbance values, and inhibitory ratios with the increasing concentrations of GA (Figure 4B-D) and Que (Figure 4E-G) were obtained and compared with those acquired by using purified α -Glu (Figure S3D-I). For both GA and Que, the inhibitory ratios (80.25% and 70.99%, respectively) and IC_{50} values (1.4 mM and 13.5 μ M, respectively) tested using purified enzyme were in agreement with the corresponding ratios (78.1% and 70.1%, respectively) and values (1.36 mM and 12.4 μ M, respectively) obtained by membrane-bound enzyme. The results confirmed that the method could not only be successfully employed for the evaluation of inhibitors using purified α -Glu but also by the membrane-bound enzyme.

Conclusion

In this study, a new strategy for direct analysis of a membrane-bound enzyme has been proposed based on the inhibition of ion transport by the lipid bilayer. We used membrane-bound α -Glu as an example, since its substrate can mediate the self-assembly of lipid bilayer around AMNSs to inhibit the release and transport of iron ions under acidic pH. A simple and specific method was established for α -Glu. The method could test the activities of both purified α -Glu and membrane-bound α -Glu in cell medium. Furthermore, it could be applied for the evaluation of inhibitors at the molecular as well as cellular levels. Our study provides a new strategy for the development of a method to detect membrane-bound enzymes *in situ*.

Abbreviations

ABTS: 2, 2'-azino-bis(3-ethylbenzothiazoline-6-sulphonic acid) diammonium salt; ALP: alkaline phosphatase; AMNSs: assembled magnetic nanospheres; APBA: 3-aminophenylboronic acid; BSA: bovine serum albumin; DSPE-PEG-NHS: 1, 2-distearoyl-sn-glycero-3-phosphoethanolamine (DSPE) conjugated polyethylene glycol 10000 (PEG) and N-hydroxysuccinimide (NHS); EDC: 1-(3-dimethylaminopropyl)-3-ethylcarbodiimide hydrochloride; GA: gallic acid; GO: glucose oxidase; HEPES: N-2-hydroxyethylpiperazine-N'-2-ethanesulfonic acid; LB: lipid bilayer; LF: lactoferrin; Mal: maltopentaose; MBEs: membrane-bound enzymes; MES: 2-(N-morpholino)ethanesulfonic acid; Ova: ovalbumin; pAPG: *para*-aminophenyl- α -D-glucopyranoside; Que: quercetin.

Supplementary Material

Experimental procedures for the preparation of AMNSs and APBA/AMNSs, characterization of the

lipid bilayer, cell culture, specificity analysis, results and discussion for optimization of experimental conditions, as well as purified α -Glu assay and screening of its inhibitors are provided in the Supplementary Material and figures.

<http://www.thno.org/v08p3275s1.pdf>

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Competing Interests

The authors have declared that no competing interest exists.

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