ORIGINAL ARTICLE

Preparation of nanogel-immobilized porous gel beads for affinity separation of proteins: fusion of nano and micro gel materials

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We describe the preparation and evaluation of nanogel-immobilized porous gel beads (GB) for application as a protein purification medium. Nanogel particles (NP) that bind with the Fc fragment of immunoglobulin G (IgG) were immobilized on the pore surface of macroporous hard GB containing quaternary ammonium cations on the surface via multipoint electrostatic interactions. The amount of NPs that were irreversibly immobilized in 1 ml of GB slurry was determined to be ~ 30 mg using fluorescent-labeled NPs. Images obtained via scanning electron microscopy established that the NPs were uniformly immobilized on the surface of the pores without blocking the macropores. The model target protein (IgG) was reversibly captured by the NPimmobilized GBs through NP-IgG interactions. NP-immobilized GBs have potential applications as novel affinity purification media for proteins, combining inexpensive and stable ligands with high-performance supports. Polymer Journal (2015) 47, 220-225; doi:10.1038/pj.2014.101; published online 26 November 2014

INTRODUCTION

Porous microbeads immobilized with affinity ligands that selectively bind to target proteins are important as the purification media for proteins. For example, antigens and antibodies can be purified via affinity chromatography using antibody or antigen-immobilized beads as an affinity separation medium. The affinity purification process of monoclonal antibodies using protein A-immobilized beads is one of the most important processes in the production of monoclonal antibody drugs.

Synthetic ligands that strongly bind to specific proteins are of great interest as stable and inexpensive alternatives to biomacromolecular ligands.^{1,2} Schrader and colleagues have reported that synthetic polymer hosts that bind to specific domains of target proteins^{3,4} can be prepared by copolymerizing an optimized combination of monomers that interact strongly with side chains of specific amino acids. Haag and colleagues reported that dendritic polymer ligands for specific proteins could be prepared by tuning the type and combination of functional groups available on the surface of the polymers.⁵⁻⁸ Shea and colleagues reported that multifunctional nanogel particles (NPs) polymerized with a combination of functional monomers capture target peptides,9-12 proteins13-15 and specific domains of proteins¹⁶ via a combination of multipoint electrostatic, hydrophobic, aromatic17 and/or hydrogen binding18 interactions. Target affinity can be enhanced by optimizing the feed ratio of functional monomers,¹⁹ by molecular imprinting polymerization^{20,21} and/or by affinity purification procedures.²² It has also been reported that porous

membranes polymerized with functional monomers containing carbohydrates can be used as affinity separation media for lectins.^{23–25}

Solid supports for affinity purification media must be carefully designed to achieve an efficient purification process.²⁶ Porous materials with large surface areas and pore diameters larger than that of the target protein (>30 nm for IgG), enable high binding capacity and fast adsorption kinetics.²⁷ The supports must be hydrophilic (ideally a hydrophilic gel such as agarose) to reduce non-specific adsorption of proteins on the surface.²⁸ However, soft and hydrophilic gel beads (GB) are not suitable for high flow rates and/or large-scale processes because the gel structures would easily collapse under the backpressure and would become stuck in the column.29

Macroporous and hydrophilic GB stabilized by cross-linkers are ideal solid supports for high-capacity, large-scale and fast protein purification processes.³⁰ We have reported that hydrophilic macroporous GB with a high-pressure resistance can be prepared by the suspension polymerization of methacrylate and a cross-linker in the presence of porogens.³¹ The GB have a large pore diameter of ~ 50 nm and a large surface area of 50 m² g⁻¹, which enable 1 ml of gel slurry to adsorb 50 mg of IgG.³² Furthermore, the spherical and rigid beads enable an extremely high flow rate purification process, with a flow rate of 10 m min⁻¹.

In this study, we report a novel affinity separation medium for proteins that consist of macroporous GB as the solid support and multifunctional NPs as the affinity ligands. Poly-N-isopropylacrylamide NPs copolymerized with an optimized feed ratio of N-tert-

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butylacrylamide (TBAm) and acrylic acid (AAc) were synthesized as affinity ligands for the Fc fragment of IgG.¹⁷ Macroporous GBs with a pore diameter of 50 nm and a quarterly ammonium group immobilized on the pore surface were used as solid supports. A single layer of NPs was immobilized on the pore surface using multipoint electrostatic interactions between the quarterly ammonium cations and the carboxylate anions.^{33–35} The amount of NPs immobilized on the GBs was calculated from the concentration of NPs (measured using fluorescently labeled NPs) remaining in the solution after the immobilization process. The morphology of NP-immobilized pore structures was observed using a scanning electron microscope. Finally, the amount of IgG, which can be reversibly captured by the NP-immobilized GBs, was quantified.

EXPERIMENTAL PROCEDURE

Materials

Most of the chemicals were obtained from commercial sources. N,N,N', N'-tetramethylethylenediamine, dansyl chloride, triethylamine and ammonium persulfate were purchased from Sigma-Aldrich Japan (Tokyo, Japan) Nisopropylacrylamide (NIPAm), AAc, TBAm and sodium dodecyl sulfate were obtained from Wako Pure Chemical Industries (Osaka, Japan), Ltd. N,N '-methylenebisacrylamide (BIS), ethylenediamine and methacryloyl chloride were purchased from Tokyo Chemical Industry Co. Ltd., (Tokyo, Japan), Thermo Fisher Scientific K. K. (Waltham, MA, USA) and Alfa Aesar (Ward Hill, MA, USA), respectively. NIPAm was recrystallized from hexanes. The porous polymer beads (MCI gel CQA35P) were provided by Mitsubishi Chemical Corporation (Tokyo, Japan) and were washed with an excess amount of water, phosphate buffered saline (PBS; 10 mM phosphate, 137 mM NaCl and 27 mM KCl, pH 7.4), an aqueous solution of NaOH (0.1 M) and water, two to three times each before use. The water used in this study was purified using a Millipore water purification system (Merck, Darmstadt, Germany). Other chemicals were used as received.

Synthesis of N-(2-Aminoethyl)-5-(dimethylamino)-1-naphthalene sulfonamide



The procedure reported by Shea and colleagues¹⁹ was modified to prepare compound 1. A solution of dansyl chloride (0.94 g, 3.47 mmol) in dry tetrahydrofuran (THF) (75 ml) was added dropwise to a cooled (0 °C) solution of ethylenediamine (2.32 ml, 34.7 mmol) in dry THF (150 ml). The reaction mixture was stirred at 0 °C for 3 h, followed by the addition of aqueous KOH (10 ml, 1 N). The THF was evaporated and the aqueous layer was extracted with CH₂Cl₂ (5×50 ml). The combined organic layers were dried over MgSO₄ and evaporated to leave a pale yellow-green solid. Recrystallization from benzene provided light-green needles (0.333 g, 1.14 mmol, 32%; proton nuclear magnetic resonance (CDCl₃) δ 8.54 (d, *J* = 8.5 Hz, 1H), 8.30 (d, *J* = 8.6 Hz, 1H), 8.24 (d, *J* = 7.3 Hz, 1H), 7.55 (m, 2H), 7.17 (d, *J* = 7.3 Hz, 1H), 2.88 (m, 8H) and 2.69 (m, 2H)).

Synthesis of N-[2-[[[5-(Dimethylamino)-1-naphthalenyl]sulfonyl] amino]ethyl]-2-propenamide (DNS methacrylate)



Methacryloyl chloride (0.112 ml, 1.36 mmol) and triethylamine (0.160 ml, 1.36 mmol) were added to a solution of N-(2-Aminoethyl)-5-(dimethylamino)-1-naphthalene sulfonamide (0.333 g, 1.14 mmol) in THF (33 ml) at room temperature. The reaction mixture was stirred at room temperature for 16 h,

followed by the addition of more methacryloyl chloride (0.056 ml, 0.68 mmol) and triethylamine (0.08 ml, 0.68 mmol). The reaction mixture was further stirred at room temperature for 3 h, and the salts were then filtered and washed with THF. The combined filtrate was evaporated and purified via flash chromatography (1:1 ethyl acetate: ethyl ether), which yielded DNS methacrylate as a light-green solid (0.170 g, 0.470 mmol, 41%; proton nuclear magnetic resonance (CDCl₃) δ 8.52 (d, *J* = 8.5 Hz, 1H), 8.27 (d, *J* = 8.6 Hz, 1H), 8.19 (dd, *J* = 1.0 and 7.3 Hz, 1H), 7.50 (m, 2H), 7.16 (d, *J* = 7.5 Hz, 1H), 6.58 (br t, *J* = 5.2 Hz, 1H), 6.15 (t, *J* = 5.9 Hz, 1H), 5.59 (s, 1H), 5.22 (s, 1H), 3.36 (m, 2H), 3.07 (m, 2H), 2.87 (s, 6H) and 1.81 (s, 3H)).

Preparation of NPs

To improve the stability of NPs in water, NPs with different degrees of crosslinking (X mol%, where X=2, 5, 10 and 20) were prepared. NIPAm (40-X mol%), AAc (35.6 µl, 20 mol%), TBAm (133 mg, 40 mol%), Bis (X mol%), DNS methacrylate (94 µg, 0.01 mol%) and sodium dodecyl sulfate (80 mg) were dissolved in water (400 ml). TBAm and DNS methacrylate were dissolved in ethanol (4 ml) and acetone (5 ml), respectively, prior to being added to the monomer solution, resulting in a final monomer concentration of 6.5 mm. The resulting solution was degassed by bubbling nitrogen through the reaction mixture for 120 min. Following the addition of ammonium persulfate (APS, 240 mg in 4 ml water) and N,N,N',N'-tetramethylethylenediamine (120 µl) to the solution prepared above, the polymerization was carried out at 23-25 °C for 17 h under a nitrogen atmosphere. The polymerized solutions were purified by dialysis (molecular weight cut off-12 000-14 000) against an excess of water (changed over three times a day) for >4 days. The yield of the NPs was determined by measuring the weight of the NPs obtained by lyophilization of 40 ml of the dialyzed solution. The dilution in dialysis was corrected. The hydrodynamic diameter of the NPs in aqueous solutions at 25 °C was determined via dynamic light scattering (Zetasizer Nano ZS, Malvern Instruments Ltd., Worcestershire, UK). The solution temperature was preequilibrated for 2 min at 25 °C. Subsequently, the measurements were taken in triplicate at each solution temperature at 2 min intervals. All dynamic light scattering data met the quality criteria set by Malvern. The NP solutions were stored in a refrigerator at 5 °C.

Adsorption of NPs on beads

A slurry of porous polymer beads (1001) was added to a solution of NPs (3.4 mg) in water (7.1 ml) at room temperature. The mixture was stirred at 70 r.p.m. at room temperature using a shaker. The concentration of NPs remaining in the supernatant was determined by the fluorescence intensity of the dansyl group measured using a fluorescent microplate reader (Thermo Fisher Scientific K.K., Varioskan Flash) or a fluorescent spectrometer (JASCO, Tokyo, Japan). The unbound NPs were removed from the solution by replacing the solution with water four times.

Washing of NP-adsorbed beads

A slurry of the NP-adsorbed porous polymer beads (\sim 100 µl) was incubated with PBS (1 ml) once for 5 min, seven times for 1 h, and three times for 3 h. The slurry was further washed with an acidic buffer (100 mM glycine-HCl, 1 ml) three times for 1 h.

Adsorption of IgG on NP-immobilized porous beads

NP-immobilized porous beads were equilibrated with PBS and then washed with an excess amount of water. A slurry of the NP-immobilized porous polymer beads (100 μ l) was incubated with an aqueous solution of IgG (0.5 mg ml⁻¹, 1 ml) at room temperature. The mixture was incubated at room temperature using a shaker for 3 h. The concentration of IgG remaining in the supernatant was quantified by measuring the ultraviolet (UV) absorbance at 280 nm using a UV–visible (UV–Vis) spectrometer (JASCO). The adsorption of IgG on the porous beads without NPs was also tested as a control experiment.

Desorption of IgG from NP-immobilized porous beads

After incubation with IgG, the slurry of NP-immobilized porous polymer beads $({\sim}100\,\mu l)$ was washed four times with water to remove weakly bound IgG from

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the beads. The beads were then incubated with an acidic buffer (100 mM glycine-HCl, 1 ml) at room temperature. The mixture was incubated at room temperature using a shaker for 3 h. The concentration of IgG released into the supernatant was quantified by measuring the UV absorbance at 280 nm using a UV–Vis spectrometer (JASCO). The desorption of IgG from the porous beads without NPs was also tested as a control experiment.

SEM imaging

GBs were deposited by sputtering with Pt/Pd (6/4) targets using Ion Sputter E-1030 (Hitachi High-Technologies Co., Tokyo, Japan). After the deposition of the GBs, they were imaged using a scanning electron microscope (SEM ULTRA55, Carl Zeiss, Oberkochen, Germany, 2 kV).

RESULTS AND DISCUSSION

Preparation of NPs

NPs that interact with the Fc fragment of IgG with a dissociation constant of ~ 10 nM were used as affinity ligands in this study.¹⁷ It has been reported that certain NPs are capable of interacting with the Fc fragment via a combination of hydrophobic and electrostatic interactions.¹⁷ The NPs were prepared in an aqueous solution by pseudo-precipitation polymerization of NIPAm (38 mol%), TBAm (40 mol%), AAc (20 mol%) and BIS (2 mol%) using sodium dodecyl sulfate as the stabilizer. To quantify the amount of NPs immobilized on GBs, we labeled NPs with dansyl groups by copolymerizing dansylated methacrylamide. Dansyl methacrylamide was synthesized using the method reported.²² When the NPs were polymerized with greater than 0.1 mol% of dansyl methacrylamide, a mono-modal size distribution of NPs was not obtained because of the tendency of the NPs to aggregate during the preparation process. When the NPs were polymerized with 0.01 mol% of dansyl methacrylamide, NPs with a mono-modal size distribution were obtained. However, some aggregates were observed when the NPs were incubated at 25 °C for approximately 10 h (Figure 1). To improve the stability, NPs with a larger feed ratio of cross-linker (5, 10 and 20 mol% BIS) were prepared. The NPs polymerized with greater than 5 mol% of BIS showed a mono-modal size distribution at 25 °C for more than 20 h (Table 1, Figure 1). The detection limits of 0.01 mol% dansylated NPs in aqueous solution were 10 and 0.5 µg ml⁻¹ when measured using a fluorescent plate reader and a fluorescent spectrometer, respectively.



Figure 1 Time profile of the hydrodynamic diameter of 2, 5, 10 and 20 mol % cross-linked nanogel particles, represented by black circles, gray squares, black triangles and gray diamonds, respectively, in water at 25 °C.

Adsorption of NPs on GBs

Lyon and colleagues have reported that a monolayer of NPs consisting of NIPAm and AAc can be formed by the electrostatic deposition of negatively charged NPs on a positively charged glass surface modified with quaternary ammonium cations (99–101). Utilizing this idea in Lyon's report, we attempted to immobilize NPs on the pore surface of GBs via electrostatic deposition. Macroporous poly methacrylate-based GBs ($30 \,\mu$ m) with a pore diameter of 50 nm and containing a quaternary ammonium group immobilized on the pore surface were used as the solid support.

Figure 2 shows the time profile of the concentration of NPs measured with the fluorescence intensity of the supernatant when an aqueous solution of NPs (3.4 mg, 7.1 ml) was incubated with 100 µl of the GB slurry at room temperature. The concentration of all NPs, regardless of the degree of cross-linking, decreased dramatically after incubating for several hours, indicating the adsorption of NPs on the GBs. The adsorption kinetics of slightly aggregated and relatively larger NPs polymerized with 2 mol% BIS was slower than that of stable and smaller NPs (polymerized with 5, 10, 20 mol% BIS). This result indicates that the NPs must be small enough to diffuse immediately into the macropores of the GBs.

Scanning electron microscope images of the GBs before and after adsorption of nanogel particles are shown in Figure 3. Figures 3b and d show little aggregation of NPs on or around the GBs. From this result, combined with the quantitative adsorption

Table 1 Composition, yield and hydrodynamic diameter of NPs

NP#	Feed ratio of functional monomers (mol%)						
	NIPAm	TBAm	AAc	Dansyl	Bis	Yield (%)	Diameter (nm)
NP1	37	40	20	0.01	2	41	57.3±7.0
NP2	35	40	20	0.01	5	100	42.0 ± 0.1
NP3	30	40	20	0.01	10	100	36.9 ± 0.3
NP4	20	40	20	0.01	20	100	28.6 ± 0.4

Abbreviations: AAc, acrlamide; NIPAm, *N*-isopropylacrylamide; NPs, nanogel particle; TBAm, *N*-tert-butylacrylamide.



Figure 2 Time profile of 2, 5, 10 and 20 mol% cross-linked non (represented by black circles, gray squares, black triangles and gray diamonds, respectively) quantified by measuring the fluorescence intensity in the supernatant when an aqueous solution of the nanogel particles (3.4 mg, 7.1 ml) was incubated with $100 \,\mu$ l of the gel beads slurry at room temperature.



Figure 3 Scanning electron microscope images of gel beads before (a, c and e) and after (b, d and f) adsorption of NP1.

results from fluorescent measurement, it was confirmed that NPs were adsorbed on the GBs without forming aggregation products. Figure 3e and f indicate that the surface morphology of the pore structures after adsorption of the NPs were less angulated than the surface morphology before adsorption, although none of the pores were filled by the NPs. The apparent surface roughness of the NPs-immobilized pore surface seemed to be comparable to that of the mono-layered dried NPs films (<10 nm),^{10,19,20} suggesting that the surface of GBs were coated with NPs. From those results, it was established that NP-adsorbed macroporous GBs could be prepared simply by incubating negatively charged NPs and positively charged GBs in water.

Desorption of NPs

In the industrial IgG purification process, IgG is captured by protein A-immobilized beads at a neutral pH value of \sim 7 and eluted with an acidic buffer with a pH in the range of 2–3. During the capture and elution process, it is important to ensure that bleeding of the affinity

ligand is minimized. To prevent bleeding of the NPs during the IgG capture and elution process, the NP-adsorbed beads were carefully pre-washed with buffers with pH of 7.4 and 2.3, respectively, to remove weakly adsorbed NPs.

The amount of NPs remaining on the surface of the GBs after the prewashing process is plotted in Figure 4. When the GBs were initially washed with PBS for 1 h, a significant amount of NPs were desorbed from the GBs. However, little desorption was observed after washing five times for 3 h, even when the GBs were washed with PBS and an acidic buffer (pH 2.3). This result indicates that most of the weakly adsorbed NPs can be removed by washing with PBS approximately five times. The amount of NPs that were irreversibly immobilized on the porous GBs was determined to be ~ 30 mg ml⁻¹. Although this amount is comparable to the amount of IgG that can be captured by the protein A-immobilized porous GBs, it is three times smaller than the amount of protein that can be captured by the porous ion-exchange GBs.^{31,32}

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Figure 4 Cross-linked nanogel particles in 2, 5, 10 and 20 mol% (represented by black circles, gray squares, black triangles and gray diamonds, respectively) remaining on the surface of gel beads after the prewashing process.



Figure 5 (a) Ultraviolet–visible absorption spectra of an aqueous solution of immunoglobulin before (circle) and after incubation with 2, 5, 10 and 20 mol% cross-linked nanogelparticle-immobilized gel beads (represented by black circles, gray squares, black triangles and gray crosses, respectively) and gel beads without nanoparticles (represented by gray triangles). (b) Amount of immunoglobulin reversibly adsorbed and desorbed by 2, 5, 10 and 20 mol% cross-linked nanogelparticle-immobilized gel beads (represented by black circles, gray squares, black triangles and gray diamonds, respectively).

Adsorption of IgG on NP-immobilized porous beads

A solution of NP-immobilized porous GBs (100 μ l) was incubated with an aqueous solution of IgG (0.5 mg ml⁻¹, 1 ml) at room temperature, and the amount of IgG remaining in the supernatant was measured using UV–Vis spectroscopy (Figure 5a). The GBs without NPs were also tested to estimate the extent of non-specific interactions of IgG with the GBs. Whereas the concentration of IgG in the supernatant decreased significantly after incubation with the NP-immobilized GBs, little difference was observed after incubation of the GBs without NPs, indicating that the IgG was captured by the NPs on the pore surface of the GBs. The amount of IgG adsorbed with 1 ml of the GBs slurry was ~ 3 mg.

Reversible adsorption of IgG on NP-immobilized porous beads

To elute IgG that is weakly bound to the surface of NP-immobilized GBs, the GBs were washed four times with water. Then, to elute strongly bound IgG, the beads were incubated with an acidic buffer (100 mM glycine buffer, pH 2.3). Further, the GBs were regenerated with PBS and incubated with an aqueous solution of IgG. Figure 5b

shows the amount of IgG adsorbed and eluted by the NP-immobilized GBs. Most of the IgG adsorbed on the GBs can be eluted with an acidic buffer. The GBs can be reused at least three times. Furthermore, it was confirmed using the fluorescent measurement of the eluted solutions that bleeding of NPs was not observed at all throughout the whole reversible IgG adsorption process. These results established that NP-immobilized GBs could be used as novel affinity purification media for proteins.

CONCLUSION

In this paper, NP that bind with the Fc fragment of IgG were successfully immobilized on the pore surface of macroporous hard GB via multipoint electrostatic interactions. Scanning electron microscope established that the NPs were uniformly immobilized on the surface of the pores without blocking the macropores. IgG (3 mg), was reversibly captured by the NP-immobilized GBs through NP–IgG interactions. Macroporous hard GB with nanogel-based affinity ligands on the pore surface can be used as a novel affinity purification medium for proteins, combining an inexpensive and stable ligand and a high-performance support.

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