



Enrichment and Detection of Rare Proteins with Aptamer-Conjugated Gold Nanorods

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S Supporting Information

ABSTRACT: Rare protein enrichment and sensitive detection hold great potential in biomedical studies and clinical practice. This work describes the use of aptamer-conjugated gold nanorods for the efficient enrichment of rare proteins from buffer solutions and human plasma. Gold nanorod (AuNR) surfaces were modified with a long PEG chain and a 15-mer thrombin aptamer for protein enrichment and



detection. Studies of the effect of surface modification on enrichment efficiency of thrombin showed that a change of only one EG₆ linker unit, i.e., from 2EG₆ to 3EG₆, could increase thrombin protein capture efficiency by up to 47%. Furthermore, a 1 ppm sample of thrombin in buffer could be enriched with around 90% efficiency using a low concentration (0.19 nM) of gold nanorod probe modified with 3EG₆ spacer, and with the same probe, effective capture was achieved down to 10 ppb (1 ng) thrombin in plasma samples. In addition to α -thrombin enrichment, prothrombin was also efficiently captured from plasma samples via gold nanorods conjugated with 15-mer thrombin aptamer. Our work demonstrates efficient enrichment of rare proteins using aptamer-modified nanomaterials, which can be used in biomarker discovery studies.

Rare protein enrichment has great significance in clinical applications, biomedical studies, proteomic studies, and signal transduction. Because changes in rare protein abundances or structures are often associated with disease pathogenesis, recognition of these indications may lead to early diagnosis.¹⁻⁵ Previous attempts to capture and detect low-abundance proteins from biological media have met with limited success, generally from the lack of robust, sensitive, and specific capturing probes.² To achieve these designer features, surface modification of nanoparticles, as capture scaffolds, plays an essential role.

Specific capturing can be accomplished with affinity tags such as proteins,⁶ antibodies,⁷ and aptamers⁵ that can be conjugated to nanomaterials. Affinity tag proteins can be used as bait for direct detection of their target proteins. However, this type of proteinprotein interaction is limited by weak binding, which is further decreased when the bait protein is used outside of the context of the native protein. This weak binding affinity can also lead to nonspecific binding with more abundant proteins in the medium.⁶ Antibodies are used as affinity tags in immunoprecipitation (IP) experiments, which are usually performed with gelconjugated antibodies that can specifically bind to their target proteins.⁷ However, because these agarose or sepharose beads have highly porous structures, large target protein molecules may not be able to diffuse adequately into the pores. In addition, the porous structure can also increase the nonspecific binding of smaller molecules that can easily diffuse into the pores. Another drawback of gel beads is rapid precipitation caused by their large sizes (>1 μ m), which can cause incomplete interactions with the proteins of interest.⁸ As an alternative to gel beads, monodispersed, superparamagnetic beads with diameters varying from nanometer to micrometer are used to minimize sample loss and facilitate the isolation process via magnetic separation.^{9–11} However, to fabricate magnetic nanoparticles (MNPs) with good size control and narrow size distribution, they are generally synthesized in the presence of hydrophobic surfactants in organic solvents by thermal decomposition of the precursors. For further biological applications in aqueous solutions, hydrophilic polymers should be tethered to the surface of hydrophobic MNPs, which is fairly challenging because of the inert MNP surface.¹²

Gold nanoparticles (AuNPs) have become attractive choices for biomedical applications because of their unique physical and chemical properties. Gold nanoparticles have been used for the detection of specific DNAs^{13–15} and proteins,^{3–5,16} because they can be easily functionalized via simple thiolate chemistry, and they show significant size-dependent color differences. In these "self-assembled monolayer" modifications, the dispersibility of AuNPs in solutions can be increased. For example, after PEG

Received: March 22, 2012 Accepted: June 14, 2012 Published: June 14, 2012 (polyethylene glycol) modification, AuNPs can remain suspended for months.⁵ Thus, despite the facile isolation of MNPs, AuNPs are preferred over MNPs because of their easy surface functionalization. Moreover, surface-only binding can avoid limitations in the sizes of the target proteins and eliminate the necessity of pore penetration, thereby decreasing nonspecific binding.

Aptamers represent another type of affinity protein tag, consisting of single-stranded oligonucleotides that bind to target molecules, such as small biomolecules and proteins, with affinities equal to those of antibodies. In contrast to other affinity tags, aptamers have many advantages, including small size, nontoxicity, relatively easy preparation, and functionalization with no batch-to-batch variations and easy surface immobilization via their functional groups.^{9–11,17,18}

Proteins captured via affinity tag-conjugated nanoparticles can generally be analyzed using 2-D gels, SDS-PAGE, or mass spectrometry (MS) techniques.⁴ Besides the challenges described above for protein capturing probes, even detection of captured proteins is limited, to some extent, by the sensitivity of these analytical tools. In SDS-PAGE, the proteins can be stained with Coomassie Brilliant Blue or silver staining methods, with detection limits of 50 ng and 1 ng, respectively.^{19,20} On the other hand, MS can achieve detection limits in the femtomole range.^{21,22} Regardless of the analytical method, existing affinity enrichment techniques for capture of low-abundance proteins from complex biological fluids still need improvement.

Therefore, in this work, we demonstrate the use of surfacemodified, aptamer-conjugated gold nanorods (AuNRs) for efficient capture and isolation of target proteins. As a model capturing platform, 15-mer thrombin aptamers are immobilized on the surface of gold nanorods. To increase the binding efficiency, thrombin aptamers are modified with different lengths of hexaethyloxy glycol (EG₆) linker prior to immobilization on the gold nanorod surface.

EXPERIMENTAL SECTION

Materials. Cetyltrimetylammonium bromide (CTAB) \geq 96.0% was obtained from Fluka (52370), and sodium tetrachloroaurate (III) dihydrate (99%), sodium borohydride (98%), and tris(2-carboxyethyl)phosphine (TCEP) were obtained from Sigma-Aldrich. Ascorbic acid, glycine, Tris, and silver nitrate (99%) were obtained from Fisher Scientific. Thiolterminated methoxypoly(ethylene glycol) (mPEG-SH, MW 5000) was purchased from Nanocs. Deoxyribonucleotides, spacer phosphoramidite 18, and S'-thiol modifiers were purchased from Glen Research. The pH value of the solution containing glycine (0.5 M) was adjusted with 2.0 M NaOH to 8.0. 20X of NuPAGE MOPS SDS running buffer, 4X of NuPAGE LDS sample buffer, and 4–12% Bis-Tris SDS-polyacrylamide gel with 10 wells were purchased from Invitrogen. Deionized water (18.2 M Ω ·cm) was used to prepare all of the aqueous solutions.

Preparation of AuNR Seed Solution. Five milliliters of 0.2 M CTAB solution and 5 mL of 0.5 mM NaAuCl₄ solution were mixed in a 50 mL plastic tube, and then 0.6 mL of 0.01 M freshly prepared NaBH₄ was added. Following sonication for 3 min, the mixture turned brownish-yellow. This reaction mixture was protected from the light with aluminum foil and placed in a water bath at 25 °C for 1 h.

Preparation of AuNR Growth Solution. Fifty milliliters of 0.2 M CTAB solution and 50 mL of 1 mM NaAuCl₄ solution were mixed in a 100 mL glass bottle. Then 3.25 mL of 0.004 M AgNO₃ was added to this mixture, followed by 0.7 mL of 0.0788

M ascorbic acid. After addition of ascorbic acid, the color of the solution changed from dark yellow to colorless. Then 0.12 mL of the seed solution was added, and the overall mixture was protected from the light with aluminum foil. Finally, the reaction mixture was placed in a water bath at 25 °C for 1 h. The solution changed to dark pink within 30 min.

Synthesis of AuNRs. Forty milliliters of 0.5 M glycine (adjusted pH 8.0) was prepared in a 200 mL glass bottle, and 60 mL of DI water was added, followed by 100 mL of the growth solution. The reaction mixture was protected from light with aluminum foil and was placed in a water bath at 25 °C for 12 h.

Removing Excess CTAB from AuNRs. Aliquots (25 mL) of nanorod solution were added to centrifuge tubes and were centrifuged at 14000 rpm for 8 min. For each fraction, 22.5 mL of the supernatant was removed and replaced with 22.5 mL of DI water. The solutions were again centrifuged at 14000 rpm at 25 °C for 5 min. Then 22.5 mL of the supernatant was removed from each fraction and replaced with 22.5 mL of 0.5 mM CTAB solution. The centrifugation step was repeated with the CTAB solution twice at the same speed and temperature for each fraction, and, finally, 2.5 mL of the solution was collected from each fraction.

Characterization of AuNRs. After removal of excess CTAB, all fractions were combined to have approximately 10 times concentrated gold nanorod solution. Then this nanorod solution was diluted 1:20, and the UV–vis spectrum was taken using a Cary Bio-300 (Varian, Walnut Creek, CA) to find its concentration. Two absorbance peaks were obtained at ca. 530 nm (transverse band) and ca. 860 nm (longitudinal band). TEM images of gold nanorods were obtained with a JEOL TEM 2010F transmission electron microscope on a copper grid. The sizes of gold nanorods were calculated with FemtoScan software as 77.3 \pm 5.6 nm and 17.3 \pm 1.1 nm in length and width, respectively.

Synthesis of Thrombin Aptamer. The 15-mer thrombin aptamer was selected as a model capture agent: (SH-2EG₆-Apt-FAM): 5' HS-C₆-2(EG)₆-GGT TGG TGT GGT TGG-FAM 3'; (SH-3EG₆-Apt-FAM): 5' HS-C₆-3(EG)₆-GGT TGG TGT GGT TGG-FAM 3'. All aptamers were coupled with 5'-thiol modifier for conjugation to the gold nanorod surface and were labeled with fluorescein at the 3'-end using 3'-(6-fluorescein) CPG to detect and quantify the aptamer modification on the gold nanorod surface. Also, either two or three hexaethyloxy glycol units (EG_6) were added between the thiol and thrombin aptamer sequence as spacers to observe the length effect on thrombin capture. All oligonucleotides were synthesized by solid-state phosphoramidite chemistry at a $1-\mu$ mol scale using an ABI3400 DNA/RNA synthesizer (Applied Biosystems, Foster City, CA). The completed sequences were then deprotected in AMA (ammonium hydroxide/40% aqueous methylamine 1:1) at 65 °C for 20 min and further purified using a ProStar HPLC (Varian, Walnut Creek, CA) with a C18 column (Econosil, 5 μ m, 250×4.6 mm) from Alltech (Deerfield, IL). A Cary Bio-300 UV spectrometer (Varian, Walnut Creek, CA) was used to measure absorbance to quantify the concentrations of the product sequences.

Aptamer Immobilization to the Surface of AuNRs. Before aptamer immobilization, 0.1 mM 5' S-S tagged thrombin aptamers were incubated with 5 mM TCEP in 50 mM Tris/HCl (pH = 7.5) buffer for 1 h at room temperature to reduce S-S to SH groups. The 5' SH-thrombin aptamers were then collected in small portions by eluting the TCEP mixture through a NAP-5 column. Concentrations of these portions were calculated by measuring their absorbance using a Cary Bio-300 UV

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Figure 1. (a) TEM image of the gold nanorods with the dimensions 77.3 ± 5.6 nm and 17.3 ± 1.1 nm. (b) Absorption spectrum of gold nanorods with two absorption maxima at 530 and 860 nm.



Figure 2. (A) Surface modification design of gold nanorods, where x = 2, 3. (B) Scheme of the α -thrombin capturing protocol: 1, 2, and 3 represent 0.19, 0.48, and 0.96 nM gold nanorods conjugated with $3EG_6$ - or $2EG_6$ -modified thrombin aptamers, respectively.

spectrometer (Varian, Walnut Creek, CA). Then a 100 μ L sample of 0.96 nM gold nanorod solution was centrifuged at 14000 rpm at 25 °C for 3 min. The supernatant was removed, and the precipitate was resuspended in 100 μ L of 2 mM CTAB. Then 850 μ L of DNA grade water, 50 μ L of 2 mM thiol-PEG (MW 5000), and 0.25 μ L of 100 μ M (25 nM as final concentration) thrombin aptamer were added to the gold nanorod solution. The solution was incubated for 12 h at room temperature. Then the reaction solution was centrifuged at 14000 rpm at 25 °C for 5 min to remove the unbound aptamers and SH-PEG as a supernatant. The precipitate was resuspended in 100 μ L of DI or DNA grade water. Gold nanorods conjugated

with thiol-PEG only (MW 5000) were prepared in the same manner.

Thrombin Capture with Aptamer-Conjugated AuNRs. Three samples, including 20, 50, and 100 μ L of 0.96 nM gold nanorod—thrombin aptamer (2EG₆- or 3EG₆-modified), and 50 μ L of 0.96 nM of only SH-PEG-modified gold nanorod were centrifuged at 14000 rpm for 5 min, and the supernatants were removed. Then 100 μ L of thrombin activation buffer and human plasma samples containing 338 ng, 100 ng, or 1 ng human α -thrombin were prepared separately for each precipitated gold nanorod solution. The activation buffer solutions also contained 10 mg/mL BSA (3000 times more concentrated than α -

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Figure 3. The fluorescence intensities of (A) $2EG_6$ -modified aptamers, (B) $3EG_6$ -modified aptamers, (upper panels) aptamer-conjugated AuNRs, (lower panels) unbound aptamers, while varying the concentrations of SH-PEG (MW 5000) in the surface modification of AuNRs. (Aptamer is abbreviated as Apt).

thrombin in the 338 ng α -thrombin-spiked buffer solutions) to mimic multiplex biological fluids. Thrombin-spiked activation buffer (20 mM Tris-HCl pH = 7.4, 140 mM NaCl, 5 mM KCl, 1 mM CaCl₂, 1 mM MgCl₂) and plasma samples were added to the precipitates of gold nanorod solutions separately, and the mixtures were vortexed, or briefly sonicated, to mix the contents. The mixtures were incubated for 20 min at room temperature and were then centrifuged at 14000 rpm for 3 min. The precipitates were washed three times with 200 μ L of washing buffer and centrifuged at 14000 rpm for 3 min after each buffer addition; the precipitates were then dispersed in 10 μ L of DI water.

One-Dimensional Gel Electrophoresis and Gel Staining. A volume of 10 μ L of the prepared α -thrombin protein standards and the gold nanorod solutions with α -thrombin captured from human plasma or buffer were mixed with 10 μ L of gel loading buffer (NuPAGE LDS sample buffer; Invitrogen). The mixtures and 2 μ L of the prestained protein marker solution were heated at 95 °C for 5 min and loaded into the gel (4-12% Bis-Tris SDS-polyacrylamide (Invitrogen)) with a thickness of 1.0 mm. Gel separation in running buffer proceeded at a constant voltage of 200 V for 1 h. After separation, the gels that were loaded with 338 ng or 100 ng thrombin samples were washed with DI water three times at 10 min increments, and the gels were stained with Coomassie Blue dye (GelCode Blue Stain Reagent, Thermo Scientific) for 1 h. The gels that were loaded with 1 ng thrombin samples were stained by the silver staining method using a SilverQuest Staining Kit (Invitrogen). Then the stained

gels were scanned with a scanner, and the bands in the gels were analyzed by ImageJ software. To determine the thrombin capture efficiency, the peak areas of the protein standard bands were considered to represent 100% yield.

RESULTS AND DISCUSSION

To synthesize gold nanorods, a seed-mediated method was used.^{23–26} The synthesized gold nanorods were characterized by their UV–vis absorption and by their TEM images, as shown in Figure 1. The prepared gold nanorods showed good uniformity, as indicated by their TEM image. The characteristic transverse and longitudinal bands²³ were observed at 530 and 860 nm, respectively, in the UV–vis absorption spectrum (Figure 1).

Surface modification of gold nanorods utilizing thiolate reaction consisted of PEGylation of the surface using SH-PEG (MW 5000) and immobilization of thiol-labeled 15-mer thrombin aptamers that were previously modified with different hexaethyloxy glycol (EG₆) lengths (Figure 2). Gold nanorods were functionalized with thiol-PEG (MW 5000) to avoid agglomeration and decrease nonspecific binding.^{5,27–31}

The immobilization of thiol-PEG was optimized by monitoring the fluorescence intensities of the FAM-labeled aptamerconjugated gold nanorods and the unbound aptamers for different concentrations of SH-PEG. As shown in Figure 3, the highest fluorescence intensity of aptamer-conjugated gold nanorods and the least fluorescence intensity of the unbound aptamers were obtained when 100 μ M SH-PEG was used, and that concentration was subsequently used for AuNRs with either



Figure 4. Upper panels: Gel electrophoresis of enriched α -thrombin (A) from 338 ng of α -thrombin-spiked activation buffer: lane 1, α -thrombin standard; lanes 2, 3, and 4, α -thrombin captured, respectively, via 0.19, 0.48, and 0.96 nM aptamer–2EG₆–AuNRs; lanes 5, 6, and 7, α -thrombin captured, respectively, via 0.19, 0.48, and 0.96 nM aptamer–3EG₆–AuNRs; lane 8, α -thrombin captured via 0.48 nM of PEGylated-only gold nanorods and (B) from 100 ng of α -thrombin-spiked activation buffer: lane 1, α -thrombin standard; lanes 2, 3, and 4, α -thrombin captured, respectively, via 0.96, 0.48, and 0.19 nM aptamer–2EG₆–AuNRs; lanes 5, 6, and 7, α -thrombin captured, respectively, via 0.19, 0.48, and 0.96 nM aptamer–3EG₆–AuNRs; lanes 8, α -thrombin captured, respectively, via 0.96, 0.48, and 0.19 nM aptamer–2EG₆–AuNRs; lanes 5, 6, and 7, α -thrombin captured, respectively, via 0.19, 0.48, and 0.96 nM aptamer–3EG₆–AuNRs; lane 8, α -thrombin captured via 0.48 nM of PEGylated-only gold nanorods. Figures on the right: Pixel intensity peaks of each band drawn by ImageJ software. Lower panels: Relative band intensities calculated by ImageJ software for the captured α -thrombin compared to α -thrombin standards to demonstrate the thrombin capture percentage.

two or three EG₆ units. This amount of SH-PEG also led to welldispersed AuNR solutions (Figure S-1, Supporting Information). The average number of aptamers bound on the surface per nanorod was found by dividing the concentration of aptamers bound on the surface by the concentration of the gold nanorods used in conjugation. To find the concentration of the bound aptamers, the concentration of the unbound aptamers was subtracted from the overall aptamer concentration that was used in the modification. The concentrations of the aptamers were evaluated using a fluorescence standard calibration curve for standard solutions of the thrombin aptamer.^{32,33} According to these calculations, approximately 162 and 157 of 2EG6- and 3EG₆-modified 15-mer thrombin aptamers were immobilized on each gold nanorod surface, respectively, while using 100 μ M SH-PEG (MW 5000) (Figures S-2 and S-3, Supporting Information). Because the aptamer immobilization did not differ significantly for either of these modifications, the thrombin capture efficiency can be compared with the change in the linker length.

The aptamer accessibility toward its target can be increased by using linkers, such as thymidine³⁴ and PEG.³⁵ In previous reports, arbitrarily chosen lengths of these linkers were used.^{31,34} Balamurugan et al. showed that there are two factors affecting the target capture efficiency, surface density, and the distance from the immobilization surface of the aptamers. In their study, thymidine linkers in four different increments (T₀, T₅, T₁₀, T₂₀) decreased the surface density of thrombin aptamers immobilized on gold slides, resulting in decreased target capturing efficiency in T₅ \rightarrow T₁₀ and T₁₀ \rightarrow T₂₀ increment change; however in the T₀ \rightarrow T₅ modification, the distance effect suppressed the surface density decrease, which increased the accessibility of aptamers

toward their target.³¹ In this study, 15-mer thrombin binding aptamers were modified with either two or three units of hexaethyloxy glycol to determine which length resulted in more efficient thrombin capture. Instead of a thymidine, hexaethyloxy glycol was chosen as a linker because it does not affect the surface density of the aptamers.³¹ On the other hand, one unit of hexaethyloxy glycol is approximately as long as a T₅ linker.

Gold surfaces have a high tendency to quench fluorescence of nearby fluorophores.^{36,37} Consequently, before starting the thrombin capturing efficiency comparison experiments, it was necessary to estimate the distance between aptamer and AuNR surface. Therefore, the fluorescence quenching percentages of $3EG_{6^-}$ and $2EG_{6^-}$ modified thrombin aptamers having FAM dye on their 3' ends were compared. The fluorescence quenching percentage of $3EG_6$ -modified aptamers was found to be 10% less than that for aptamers with $2EG_6$ units, indicating that the $3EG_6$ modified aptamers were farther away from the AuNR surface (Figure S-4, Supporting Information).

Thrombin capturing efficiency comparison experiments were then conducted by an SDS-PAGE technique. Different concentrations (0.19, 0.48, and 0.96 nM) of gold nanorods conjugated with $3EG_{6^-}$ or $2EG_{6^-}$ modified thrombin aptamers were incubated with 338 ng, 100 ng, or 1 ng human α -thrombinspiked buffer or human plasma samples. The enriched α thrombin on these gold nanorods and a standard human α thrombin sample were loaded in a 4–12% Bis-Tris SDSpolyacrylamide gel for electrophoresis. Also, gold nanorods that were conjugated only to thiol-PEG (MW 5000) were incubated with the samples containing human α -thrombin protein to determine any contribution from nonspecific enrichment.

As shown in Figure 4, an increase in concentration of the AuNRs conjugated with 2EG₆-modified thrombin aptamers from 0.19 nM to 0.96 nM resulted in an increase in thrombin capturing efficiency from 48% to 84% for the buffer sample containing 338 ng (~84 nM, 3.38 ppm) of human α -thrombin (Figure 4A, lanes 2-4). However, when the concentration of the AuNRs conjugated with 3EG₆-modified thrombin aptamers increased from 0.19 nM to 0.96 nM, the thrombin capturing efficiency decreased from 90% to 28% for the same thrombin standard (Figure 4A, lanes 5-7). Therefore, the most efficient capturing occurred with the 0.19 nM 3EG₆-modified aptamer-conjugated AuNRs. A similar thrombin capturing trend occurred for the buffer sample containing 100 ng (~25 nM, 1.00 ppm) of α thrombin. In Figure 4B, the thrombin capture efficiency difference between the 0.19 nM 3EG6- and 2EG6-modified thrombin aptamer-conjugated AuNRs is even more evident (lane 5 and lane 4, respectively). Thus, even a one unit change of hexaethyloxy glycol increased the thrombin capturing efficiency by 47%.

As reported by Balamurugan et al.,³¹ the thrombin binding capacity was controlled by two different factors, surface density and the distance from the immobilization surface of the aptamers. In this paper, because the numbers of $2EG_{6}$ and 3EG₆-modified thrombin aptamers immobilized on each gold nanorod surface were close to each other, the distance of aptamers from the gold surface became the major effect on thrombin capture efficiency. For AuNRs conjugated with 2EG₆modified thrombin aptamers, which had shorter linkers, the thrombin capture efficiency was greatly affected by the steric effect. Thus, larger concentrations of AuNRs were needed to increase the chance that α -thrombin would come close to the aptamers immobilized on the gold surface. That is the reason for the increase in the thrombin capture efficiency with the increase of the concentration of AuNRs conjugated with 2EG₆-modified thrombin aptamers. However, AuNRs conjugated with 3EG₆modified thrombin aptamers, which had longer linkers, showed slight steric effects on thrombin binding to aptamer. Thus, even low concentrations of AuNRs (0.19 nM) conjugated with 3EG₆modified thrombin aptamers could reach up to 90% of thrombin capture efficiency. On the other hand, with the increase of the concentration of AuNRs, the 3EG₆-modified thrombin aptamers on the adjacent AuNRs may have twisted around each other because of their flexibility and longer distance from the gold surface. Therefore, the effective number of aptamers that can bind to thrombin was reduced. This may be the reason why thrombin capture efficiency was decreased with the concentration increase of AuNRs conjugated with 3EG₆-modified thrombin aptamers.

In Figure 4A and 4B, lane 8 shows that the nonspecifically captured α -thrombin from buffer solutions via gold nanorods conjugated only with thiol-PEG (MW 5000).⁵ Because the band intensities for this nonspecifically captured α -thrombin were so low, it can be concluded that the nonspecific capture of α -thrombin in the absence of the affinity tag conjugation was negligible.

These buffer solutions contained α -thrombin, as well as high concentrations of BSA (10 mg/mL), to mimic complex biological fluids. As shown in Figure S-5 in Supporting Information, thrombin aptamer-conjugated gold nanorods showed some nonspecific binding to BSA (66 kDa), but the band intensities were very low compared to the bands for captured α -thrombin (37 kDa), indicating that the nonspecific binding was negligible. The gel in Figure S-5 also shows that the

thrombin capture efficiency increased with an increase in the concentration of $2EG_6$ -modified aptamer-conjugated AuNRs, but it decreased with an increase in the concentration of $3EG_6$ -modified aptamer-conjugated AuNRs. Moreover, 0.96 nM aptamer $-2EG_6$ -AuNRs and 0.19 nM aptamer $-3EG_6$ -AuNRs again showed similar band intensities for captured α -thrombin, which is an indication of the efficacy of aptamer $-3EG_6$ -AuNRs, despite their low concentration.

Thrombin capturing efficiency comparison experiments were also conducted with 1 ng (\sim 250 pM, 10 ppb) human α thrombin-spiked buffer and human plasma samples. This concentration was chosen since 1 ng is close to the detection limit of the silver staining method for proteins after gel electrophoresis.^{19,20} Significant differences in thrombin capture efficiency for different probes from 1 ng α -thrombin-spiked buffer solutions were not observed, as shown in Figure S-6, Supporting Information.

Similarly, the α -thrombin capture efficiency trend observed with thrombin aptamer activation buffer solutions was not observed with the 1 ng α -thrombin-spiked human plasma sample (Figure 5). The band intensities of the captured α -thrombin (red dashed lines, Figure 5) via different concentrations of $2EG_{6^-}$ or $3EG_{6^-}$ modified thrombin aptamer-conjugated AuNRs did not differ significantly from each other (except in lane 7, which could have resulted from a staining error), and they even showed slightly higher intensities than thrombin standard (lane 1) by the presence of some α -thrombin in the original plasma sample. Even though one hexaethyloxy glycol unit change of the spacer in the surface modification of gold nanorods did not affect the thrombin capturing efficiency significantly in plasma samples, 1 ng (10 ppb) of α -thrombin was successfully enriched from



Figure 5. Left panel: Gel electrophoresis of enriched α -thrombin from 1 ng α -thrombin-spiked human plasma: L, ladder; lane 1, α -thrombin standard; lanes 2, 3, and 4, α -thrombin captured, respectively, via 0.19, 0.48, and 0.96 nM aptamer–2EG₆–AuNRs; lanes 5, 6, and 7, α -thrombin captured, respectively, via 0.19, 0.48, and 0.96 nM aptamer–3EG₆–AuNRs. Right panel: Pixel intensity peaks of each lane drawn by ImageJ software; blue dashed lines show the upper bands (72 kDa), and red dashed lines show the lower bands (37 kDa).

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human plasma with the aptamer-conjugated gold nanorods. An unexpected band could also be seen in the gel electrophoresis of the α -thrombin standard (blue dashed lines, lane 1, Figure 5). To identify this band, the MALDI-MS spectrum of the α -thrombin standard was taken. Besides the expected mass-to-charge ratio peak at around 36000 m/z, another peak was obtained at around 72000 m/z (Figure S-7, Supporting Information). This was most likely caused by the prothrombin content of the α -thrombin standard solution, because α -thrombin (~37 kDa) is generated by proteolysis of two peptide bonds in prothrombin (\sim 72 kDa) via prothrombinase enzyme.^{38,39} The same band (\sim 72 kDa) occurred for the lanes of the captured α -thrombin in Figure 5. This result indicates that 15-mer thrombin aptamer can also bind to prothrombin. This correlates with some previous studies, which found that a 15-mer thrombin aptamer can block the clotting activity of thrombin either by binding to exosite I or prothrombin to inhibit its activation by prothrombinase enzyme, which also leads to the inhibition of thrombin generation.⁴⁰ The band intensity for prothrombin in the standard (lane 1, Figure 5) did not substantially differ from band intensities recorded for captured prothrombin via aptamer-conjugated gold nanorods modified with different linker lengths (lanes 2-6). Thus, it may be concluded that effective capturing was achieved at ppb levels for prothrombin using 15-mer thrombin aptamer-conjugated gold nanorods.

CONCLUSION

The overall results in this study demonstrate the applicability of the surface-modified, aptamer-conjugated gold nanorods for the enrichment of low-abundance proteins (in ppb range) from plasma samples and the vital role of surface modification to generate robust, sensitive, specific, and efficient protein capturing platforms. This new method can enrich proteins at ppb levels using very low amounts of the capturing platform (0.19 nM gold nanorods) and with negligible nonspecificity. The dramatic increase in thrombin capturing efficiency with only one unit change of hexaethyloxy glycol linker proves the significance of surface modification in protein enrichment techniques. This study also outscores similar, but more complex, methods only capable of enrichment down to 6 ng of captured thrombin.⁴¹ In addition to the detection of proteins using gel electrophoresis, this work can also open new avenues for aptamer-conjugated gold nanorods for use as an LDI-MS matrix⁴² where NIR lasers are used. In this way, aptamer-conjugated gold nanorods can be used as a dual platform for enrichment and detection of rare proteins utilizing MS, as shown in our previous work with aptamer-conjugated graphene oxide.43 It may be concluded that aptamer-conjugated gold nanorods are promising protein capturing platform candidates for biomarker discovery studies involving capture and detection of low-abundance proteins from biological fluids.

ASSOCIATED CONTENT

S Supporting Information

Additional information as noted in the text. This material is available free of charge via the Internet at http://pubs.acs.org.

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Notes

The authors declare no competing financial interest.

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REFERENCES

(1) Anderson, N. L.; Anderson, N. G. Mol. Cell. Proteomics 2002, 1, 845–867.

(2) Darmanis, S.; Nong, R. Y.; Hammond, M.; Gu, J.; Alderborn, A.; Vanelid, J.; Siegbahn, A.; Gustafsdottir, S.; Ericsson, O.; Landegren, U.; Kamali-Moghaddam, M. *Mol. Cell. Proteomics* **2010**, *9*, 327–335.

(3) Kirk, J. S.; Bohn, P. W. J. Am. Chem. Soc. 2004, 126, 5920–5926.
(4) Wang, A.; Wu, C. J.; Chen, S. H. J. Proteome Res. 2006, 5, 1488–1492.

(5) Cheng, P. C.; Chang, H. K.; Chen, S. H. Mol. Cell. Proteomics 2010, 9, 209-224.

(6) Schulze, W. X.; Mann, M. J. Biol. Chem. 2004, 279, 10756–10764.
(7) Phizicky, E. M.; Fields, S. Microbiol. Rev. 1995, 59, 94–123.

(8) Alber, F.; Dokudovskaya, S.; Veenhoff, L. M.; Zhang, W.; Kipper, J.;

Devos, D.; Suprapto, A.; Karni-Schmidt, O.; Williams, R.; Chait, B. T.; Sali, A.; Rout, M. P. *Nature* **200***7*, *450*, 695–701.

(9) Sefah, K.; Phillips, J. A.; Xiong, X.; Meng, L.; Van Simaeys, D.; Chen, H.; Martin, J.; Tan, W. Analyst **2009**, 134, 1765–1775.

(10) Meir, A.; Marks, R. S.; Stojanovic, M. N. Aptameric Biosensors. In *Handbook of Biosensors and Biochips*; Cullen, D. C., Karube, I., Lowe, C. R., Weetal, H. H., Eds.; John Wiley & Sons, Ltd: New York, 2008.

(11) Shangguan, D.; Tang, Z.; Mallikaratchy, P.; Xiao, Z.; Tan, W. ChemBioChem 2007, 8, 603–606.

(12) Tong, S.; Hou, S.; Ren, B.; Zheng, Z.; Bao, G. *Nano Lett.* **2011**, *11*, 3720–3726.

(13) Elghanian, R.; Storhoff, J. J.; Mucic, R. C.; Letsinger, R. L.; Mirkin, C. A. *Science* **1997**, 277, 1078–1081.

(14) Storhoff, J. J.; Elghanian, R.; Mucic, R. C.; Mirkin, C. A.; Letsinger, R. L. J. Am. Chem. Soc. **1998**, 120, 1959–1964.

(15) Storhoff, J. J.; Lazarides, A. A.; Mucic, R. C.; Mirkin, C. A.; Letsinger, R. L.; Schatz, G. C. J. Am. Chem. Soc. 2000, 122, 4640–4650.
(16) Teng, C. H.; Ho, K. C.; Lin, Y. S.; Chen, Y. C. Anal. Chem. 2004, 76, 4337–4342.

(17) Mallikaratchy, P.; Tang, Z.; Kwame, S.; Meng, L.; Shangguan, D.; Tan, W. *Mol. Cell. Proteomics* **2007**, *6*, 2230–2238.

(18) Shangguan, D.; Cao, Z.; Meng, L.; Mallikaratchy, P.; Sefah, K.; Wang, H.; Li, Y.; Tan, W. J. Proteome. Res. **2008**, *7*, 2133–2139.

(19) Williams, L. Biotech. Histochem. 2001, 76, 127-132.

(20) Roegener, J.; Lutter, P.; Reinhardt, R.; Blüggel, M.; Meyer, H. E.; Anselmetti, D. Anal. Chem. 2003, 75, 157–159.

(21) Aebersold, R.; Goodlett, D. R. Chem. Rev. 2001, 101, 269-295.

(22) Aebersold, R.; Mann, M. Nature 2003, 422, 198-207.

(23) Nikoobakht, B.; El-Sayed, M. A. Chem. Mater. 2003, 15, 1957–1962.

(24) Johnson, C. J.; Dujardin, E.; Davis, S. A.; Murphy, C. J.; Mann, S. J. Mater. Chem. **2002**, *12*, 1765–1770.

(25) Gole, A.; Murphy, C. J. Chem. Mater. 2004, 16, 3633-3640.

(26) Gao, J.; Bender, C. M.; Murphy, C. J. *Langmuir* **2003**, *19*, 9065–9070.

(27) Karakoti, A. S.; Das, S.; Thevuthasan, S.; Seal, S. Angew. Chem., Int. Ed. 2011, 50, 1980–1994.

(28) Liu, Z.; Robinson, J. T.; Sun, X.; Dai, H. J. Am. Chem. Soc. 2008, 130, 10876–10877.

(29) Jokerst, J. V.; Lobovkina, T.; Zare, R. N.; Gambhir, S. S. Nanomedicine (London, U. K.) **2011**, *6*, 715–728.

(30) Suh, J.; Choy, K. L.; Lai, S. K.; Suk, J. S.; Tang, B. C.; Prabhu, S.; Hanes, J. Int. J. Nanomed. **2007**, *2*, 735–741.

(31) Balamurugan, S.; Obubuafo, A.; McCarley, R. L.; Soper, S. A.; Spivak, D. A. *Anal. Chem.* **2008**, *80*, 9630–9634.

(32) Huang, Y. F.; Sefah, K.; Bamrungsap, S.; Chang, H. T.; Tan, W. Langmuir 2008, 24, 11860–11865.

Analytical Chemistry

(33) Huang, Y. F.; Chang, H. T.; Tan, W. Anal. Chem. 2008, 80, 567–572.

(34) Balamurugan, S.; Obubuafo, A.; Soper, S. A.; Spivak, D. A. Anal. Bioanal. Chem. 2008, 390, 1009–1021.

(35) Balamurugan, S.; Obubuafo, A.; Soper, S. A.; McCarley, R. L.; Spivak, D. A. *Langmuir* **2006**, *22*, 6446–6453.

(36) Lakowicz, J. R. Anal. Biochem. 2004, 324, 153-169.

(37) Swierczewska, M.; Lee, S.; Chen, X. Phys. Chem. Chem. Phys. 2011, 13, 9929–9941.

(38) Lundblad, R. L.; Kingdon, H. S.; Mann, K. G. Thrombin. In *Methods in Enzymology*; Laszlo, L., Eds.; Academic Press: New York,1976; Vol. 45, pp 156–176.

(39) Michael, E.; Nesheim, K. G. M. J. Biol. Chem. 1983, 258, 5386-5391.

(40) Kretz, C. A.; Cuddy, K. K.; Stafford, A. R.; Fredenburgh, J. C.; Roberts, R.; Weitz, J. I. *Thromb. Haemostasis* **2010**, *103*, 83–93.

(41) Li, F.; Li, J.; Wang, C.; Zhang, J.; Li, X. F.; Le, X. C. Anal. Chem. **2011**, 83, 6464–6467.

(42) Castellana, E. T.; Gamez, R. C.; Gomez, M. E.; Russell, D. H. Langmuir 2010, 26, 6066-6070.

(43) Gulbakan, B.; Yasun, E.; Shukoor, M. I.; Zhu, Z.; You, M.; Tan, X.; Sanchez, H.; Powell, D. H.; Dai, H.; Tan, W. J. Am. Chem. Soc. **2010**, *132*, 17408–17410.