Ultrasensitive protein detection in blood serum using gold nanoparticle probes by single molecule spectroscopy

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Abstract. A one-step rapid and ultrasensitive immunoassay capable of detecting proteins in blood serum is developed using gold nanoparticles and fluorescence correlation spectroscopy (FCS). In this approach we take advantage of the inherent photoluminescence property of gold nanoparticles (GNPs) to develop a fluorophore-free assay to observe binding events by monitoring the diffusion of bound versus unbound molecules in a limited confocal volume. 40-nm GNPs conjugated separately with rabbit anti-IgG (Fc) and goat anti-IgG (Fab) when incubated in blood serum containing IgG forms a sandwich structure constituting dimers and oligomers that can be differentiated by detecting IgG in blood serum at a limit of detection (LOD) of 5 pg/ml. The novelty of integrating GNPs with FCS to develop a sensitive blood immunoassay brings single molecule methods one step closer to the clinic.© 2009 Society of Photo-Optical Instrumentation Engineers. [DOI: 10.1117/1.3181789]

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All measurements were performed using a scanning confocal time-resolved microscope with a 465-nm (40-MHz pulsing) picosecond laser as an excitation source (Picoquant, Germany). The details of the instrumentation can be found elsewhere.10 The autocorrelation curve of GNP diffusion was fitted to a 3-D diffusion model by Origin Lab as Eq. (1).

\[
G(\tau) = \frac{1}{\langle N \rangle} \left(1 + \frac{\tau}{\tau_D}\right)^{-1} \left(1 + \frac{\tau}{\tau_D \cdot \kappa^2}\right)^{-1/2}.
\] (1)

Calibration of the confocal volume for autocorrelation studies was performed with rhodamine 123 before actual measurements. The autocorrelation traces were fitted using the known diffusion coefficient (D) of rhodamine (rho) 123 as 300 mm^2/sec. The radial and axial beam were calculated by the expression \(D = \omega^2 / 4 r_p\), where \(\omega\) is the radial beam size and \(r_p\) is the diffusion time. Next, 40-nm gold nanoparticles (GNPs) were synthesized by a citrate reduction of HAuCl4 \(^{11}\) and functionalized with the respective biomolecules. Its diffusion characteristics in
20% dog blood serum were investigated. As shown in the count rate trace of GNPs from FCS, GNPs did not display any bleaching of their photoluminescence [Fig. 2(a)], which is a major advantage compared to conventional fluorescence dyes. The autocorrelation curve could be fitted well by a single component, indicating isotropic diffusion. The diffusion time of GNP obtained from Eq. (1) was 1.8 ms. Viscosity of 20% blood serum was determined by evaluating the change in diffusion coefficient of rho 123 from water to 20% blood serum. In addition to diffusion, the size of the particles was also back-calculated to confirm the estimations according to the Stokes-Einstein equation \( D = kT/(6\pi \eta d) \), where \( k \) is the Boltzmann constant, \( T \) is the temperature, and \( \eta \) is the viscosity of liquid. From the calculations, the average hydrodynamic diameter of the diffusing GNP was estimated to be 42 nm, which is in agreement with the size of the as-prepared 40-nm-diam GNP.

Before proceeding with immunoassay experiments, two batches of the citrate-coated GNP were separately conjugated with either the rabbit or goat anti-human IgG antibody fragment. To ensure attachment, biofunctionalization was assessed by the red shift of the surface plasmon resonance band for the GNP and GNP-antibody conjugates from the vis-NIR spectra (Jasco, Easton, Maryland) (data not shown). The antibody conjugation resulted in an increase in the diffusion time of GNP from 1.8 to 2.4 ms, indicating an increase in the mean hydrodynamic size from 42 to 56 nm according to the Einstein-Stokes equation. The concentration of 40-nm GNP was estimated using a molar extinction coefficient of 7.666 \( \times 10^9 \) M\(^{-1}\)cm\(^{-1}\) at a plasmon resonance wavelength maximum \( \lambda_{\text{max}} \) of 528 nm.\(^3\)

Since the single-component diffusion mode [Eq. (1)] is not accurate enough for estimating diffusion of antigen-induced GNP aggregations in a heterogeneous system, a multiple-component model with a maximum entropy method called MEMFCS\(^5\) was used to obtain an optimal fit to further assess the autocorrelation. By introducing Eq. (1), \( G(\tau) \) can be related to a continuous distribution of diffusion times in MEMFCS as follows:

\[
G(\tau) = \int a_i \left( 1 + \frac{\tau}{\tau_D} \right)^{-1} \left( 1 + \kappa^2 \frac{\tau}{\tau_D} \right)^{-0.5} d\tau_D. \tag{2}
\]

The algorithm attempts to minimize the normalized chi-square value \( (\chi^2) \) while maximizing the entropy (S) term defined as \( S = \sum_i p_i \ln p_i \). Unlike the conventional fitting method where fitting components (i.e., 1, 2, 3, etc.) need to be set \textit{a priori}, MEMFCS instead yields a size distribution that is discrete and particularly suitable for analysis of heterogeneous systems. The detection of human IgG in 20% dog serum was then demonstrated using the antibody-conjugated GNP based on FCS in the following experiments. The two gold nanoparticle probes (20 \( \mu \)l of 0.2-nM concentration each) were added to 20 \( \mu \)l of 20% dog serum, and the diffusion time was observed for varying concentrations (0, 10 ng/mL, 1 ng/mL, 0.1 ng/mL, 10 pg/mL, 5 pg/mL, and 1 pg/mL) of human IgG. 30-\( \mu \)L solutions were used for measurement each time after incubating the GNP probes for 20 min at 37 °C. As expected, the half-value decay time of the normalized autocorrelation curves, which allows for a crude estimation of the diffusion time of the fluorescing particles, varies significantly with increased antigen concentration [Fig. 2(a)]. The diffusion time distribution obtained using the MEMFCS fitting routine is shown in Fig. 2(b). The GNP probes show two predominant peaks with a higher IgG concentration (10 ng/mL), one at 2.5 ms and the other at...

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**Fig. 2** (a) Autocorrelation spectra of 40 nm GNP (●) and antigen-induced GNP oligomers in the presence of different concentrations of human IgG (10 ng/mL (○), 5 pg/mL (▲)) in 20% blood serum and the corresponding fitting curves (solid line). Arrow indicates increased diffusion time. (b) The diffusion time distribution of GNP with respect to antigen concentration and the resulting GNP oligomers from the MEMFCS fit. (c) Count-rate trace of GNPs with average size of 40 nm during FCS measurements.
around 21 ms, corresponding to free GNP and oligomeric GNP, respectively. However, bare GNP showed a single peak around 1.9 ms, which is in a good agreement with the one-component fit (1.8 ms). GNP probes with 5 pg/ml IgG concentration showed a peak at 7 ms, which is in the range between 2 to 10 ms, indicating the presence of both free and oligomeric GNP oligomers. Figure 3 shows the diffusion time and hydrodynamic diameter of GNP aggregates in the presence of different concentrations of human IgG in 20% blood serum. As the antigen concentration increased from 0 to 10 ng/ml, the size of oligomers increased from 56 to 470 nm, and the diffusion time of GNP aggregates increased accordingly with higher antigen concentration (2.5 to 21 ms), as shown in Fig. 3. To further confirm the sandwich configuration resulting from the binding of the two GNP antibody conjugates (Fa and Fc probes), specifically to the target (human IgG), nonspecific binding control experiments were carried out with the two GNP probes simultaneously suspended in 20% serum without human IgG (0 ng/mL). In the control experiment, the diffusion time of the GNP probes was similar to the free GNP, demonstrating that the probes were stable with respect to nonspecific aggregation in blood serum, and their antigen-induced aggregation can be detected via the FCS at the single-particle level. Each measurement was replicated three times with three different sample preparations to demonstrate reproducibility to provide an assay at a LOD of 5 pg/ml. To the best of our knowledge, this is the first blood serum immunoassay to report sensitivities in this order within 30 min at the single-particle level. It is also important to mention that the detection limit of our assay can be improved by orders of magnitude in buffers, because protein-rich environments of blood serum could hinder the assay performance, in addition to contributing to background scattering noise.

In conclusion, by taking advantage of the superior optical property of GNP s and single-molecule spectroscopy, a label-free immunoassay that is simple, rapid, and ultrasensitive was shown to detect antigens at a LOD of 5 pg/ml in blood serum. Compared to traditional plate-based methods, our strategy is wash-free and can detect very low concentrations of targets directly in blood serum within minutes without.

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References