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# Quantitative and sensitive detection of SARS coronavirus nucleocapsid protein using quantum dots-conjugated RNA aptamer on chip

Changhyun Roh\* and Sung Kee Jo

# Abstract

BACKGROUND: Globally, severe acute respiratory syndrome coronavirus (SARS-CoV) is a newly emerging virus that causes SARS with high mortality rate in infected people. The nucleocapsid (N) protein of the severe acute respiratory syndrome (SARS)-associated coronavirus (SARS-CoV) is an important antigen for the early diagnosis of SARS and the detection of diseases. Here, a new quantum dots (QDs)-conjugated RNA aptamer with high sensitivity and rapidity is proposed for the detection of SARS-CoV N protein using an on chip system.

RESULTS: A QDs-conjugated RNA aptamer can specifically hybridize on the immobilized SARS-CoV N protein on the surface of a glass chip. Detection is based on the optical signal variation of a QDs-supported RNA aptamer interacting on an immobilized protein chip. Using an optical QDs-based RNA aptamer chip, SARS N protein was detected at concentrations as low as 0.1 pg mL<sup>-1</sup>.

CONCLUSIONS: It was demonstrated that the QDs-conjugated RNA aptamer could interact on a designed chip specifically and sensitively. This device could form a QDs-conjugated biosensor prototype chip for SARS-CoV N protein diagnosis. The proposed visual SARS-CoV N protein detection technique may avoid the limitations of other reported methods because of its high sensitivity, good specificity, ease of use, and the ability to perform one-spot monitoring. (C) 2011 Society of Chemical Industry

Keywords: SARS coronavirus (CoV); nucleocapsid (N) protein; aptamer; quantum dots (QDs)

# **INTRODUCTION**

Severe acute respiratory syndrome (SARS) is an infectious disease that began in Guandong, China in November 2002, and has caused serious infection in Asia, Europe and Canada.<sup>1-5</sup> The SARS coronavirus (CoV), which is a pathogenic virus of SARS, is a single-strand RNA virus. It is known that this virus infects animals other than humans.<sup>6-8</sup> According to the World Health Organization (WHO), mortality for patients afflicted with SARS is about 15% on average, and can be 50% or higher in the case of patients aged 65 and over.<sup>9</sup> SARS-CoV is an enveloped and positively single-stranded RNA virus with a typical genome size 29.7 kb, encoding a RNA directed RNA polymerase and structural proteins including the spike (S), envelope (E), membrane (M) and nucleocapsid (N) proteins.<sup>10–12</sup> The N protein is a 422-amino-acid alkaline protein with a short lysine-rich region suggested to be the nuclear localization signal. It plays an important role in the process of virus particle assembly by enveloping the entire genomic RNA.<sup>13</sup> The nucleocapsid (N) protein of the SARS-associated coronavirus (SARS-CoV) is an important antigen for the early diagnosis of SARS and the detection of diseases.<sup>14</sup> Since SARS broke out in 2003, researchers have made great efforts in the development of fast and accurate analytical methods for its early diagnosis.<sup>15–19</sup>

The Nucleocapsid (N) protein of SARS-CoV is a major pathological determinant in the host that may cause host cell apoptosis, up-regulate pro-inflammatory cytokine production, and block innate immune responses. Therefore, SARS-CoV N protein has long been thought an ideal target for the detection of SARS. The monitoring of SARS-CoV N protein has been of considerable interest in the development of simple and reliable methods for the detection of SARS for applications in diagnostic medicine. At present, the most widely used method of diagnosing SARS is the detection of anti-SARS antibodies using a screening enzyme-linked immunosorbent assay (ELISA), based on recombinant proteins from the genome of SARS.<sup>18,19</sup> Although it is highly specific, this assay has limitations. For example, it cannot detect viruses during the early stage of an infection, at a time when antibodies against SARS antigens are not produced. In addition, the ELISA method sometimes generates false-positive or false-negative results. In addition, using antibodies creates problem such as being

Radiation Research Division for Biotechnology, Advanced Radiation Technology Institute (ARTI), Korea Atomic Energy Research Institute (KAERI), 1266, Sinjeong-dong, Jeongeup, Jeonbuk 580-185, Republic of Korea

<sup>\*</sup> Correspondence to: Changhyun Roh, Radiation Research Division for Biotechnology, Advanced Radiation Technology Institute (ARTI), Korea Atomic Energy Research Institute (KAERI), 1266, Sinjeong-dong, Jeongeup, Jeonbuk 580-185, Republic of Korea. E-mail: chroh@kaeri.re.kr

temperature-sensitive, specific reactions conditions and requiring secondary antibody conjugated with enzyme and fluorescent dyes.

Aptamers have become increasingly important molecular tools for diagnostics and therapeutics. In particular, Aptamers exhibit advantages as recognition elements in biosensing when compared with antibodies and enzymes, thus making it possible to develop a wide range of aptamer-based biosensors. Chemically stable and cost-effective aptamers offer great flexibility in the design of novel biosensors with high detection sensitivity and selectivity.<sup>20–24</sup>

Quantum dots (QDs), which are colloidal nanoparticles of semiconductor materials, have attracted much attention in the fields of nanotechnology and biotechnology, especially in biological imaging applications because of their remarkable optical characteristics compared with conventional organic fluorophores in terms of being bright, tunable and having narrow fluorescence emission, and broad absorption spectra.<sup>25–27</sup>

In this study, it was established that optical QDs-conjugated aptamer could be used as a probe for the detection of SARS-CoV N protein using a chip system. Using an optical QDs-based RNA aptamer chip, we could detect the SARS-CoV N protein at concentrations as low as 0.1 pg mL<sup>-1</sup>. A biosensor assay for SARS-CoV N protein detection based on an optical QDs-based aptamer has attracted significant attention. It is a promising method for the specific detection of oligonucleotide because of its high sensitivity, low cost, rapid response, compatibility for miniaturization and low labor requirements.

# MATERIALS AND METHODS

## Chemicals

EDC (*N*-(3-dimethylaminopropyl)-*N*'-ethylcarbodiimide hydrochloride), bovine serum albumin (BSA), ampicillin and kanamycin were purchased from Sigma–Aldrich Chemical Co. (St Louis, MO, USA). Quantum dots (QDs 605) were purchased from Invitrogen Corporation (Carlsbad, CA, USA). Prolinker<sup>™</sup> terminated glass slides from Proteogen (Seoul, Republic of Korea) were used. All other chemicals were of the highest grade.

#### **Preparation of aptamer**

#### Subcloning, expression and purification of SARS-CoVN protein

The gene was amplified by a PCR with the primer set, sense: 5'- agt*ggatcc*atgtctgataatggaccca-3', antisense: 5'- gcc*gtcgac*ttatgcctgagttgaatcagc-3', containing restriction enzyme sites of *BamHI/Sall*. PCR was run with the following conditions on a thermal cycler: denaturation at 94 °C for 1 min, annealing at 60 °C for 30 s, and an extension step at 72 °C for 2 min 30 s. The sequence was repeated 35 times followed by a 7 min final extension step at 72 °C. The PCR product was digested with

BamHI/Sall, and then ligated into BamHI/Sall digested expression vector pET 24a+ (Novagen, Madison, WI), and transformed into E. *coli* DH5 $\alpha$  (Stratagene, La Jolla, CA). The transformed colony with inserted gene was transformed into E.coli BL21 (DE3) (Stratagene, La Jolla, CA). Then, it was plated on LB agar containing 50  $\mu$ g mL<sup>-1</sup> kanamycin. GroES/EL expressing plasmid from E.coli and SARS-CoV N-expressing plasmid, which possessed ampicillin and kanamycin resistant markers, respectively, were co-transformed into E. coli BL21 (DE3) according to the biotransformation procedures. The transformant was grown in a 250 mL flask containing 50 mL Luria-Bertani (LB) medium supplemented with 50  $\mu$ g mL<sup>-1</sup> of kanamycin and ampicillin at 37 °C until the cell concentration reached an  $OD_{600nm}$  of 0.6, and isopropyl-thio- $\beta$ -D-galactopyranoside (IPTG) of a final concentration of 0.1 mmol  $L^{-1}$ , followed by growing for an additional overnight at 25 °C with shaking at 180 rpm. The cells were harvested by centrifugation at 4000 rpm for 30 min at  $4^{\circ}C$ and resuspended in 100 mmol L<sup>-1</sup> potassium phosphate buffer (pH 7.5) containing 1 mmol  $L^{-1}$  phenylmethylsulfonyl fluoride (PMSF). The cells were lysed by Sonicator (F60 Sonic Dismembrator, Fisher Scientific, USA). The cell debris was removed by centrifugation at 13 000 rpm for 30 min. The supernatant was collected and the recombinant SARS-CoV N protein was purified with Ni-nitrilotriacetic acid (Ni-NTA) affinity chromatography column (Qiagen, Germany). The supernatant was equilibrated with buffer A (10 mmol  $L^{-1}$  Tri-HCl, 500 mmol  $L^{-1}$  NaCl, 50 mmol  $L^{-1}$ imidazole, 1 mmol L<sup>-1</sup> PMSF, pH 8.0). The bound protein was eluted with buffer B (10 mmol  $L^{-1}$  Tris-HCl, 500 mmol  $L^{-1}$  NaCl, 250 mmol L<sup>-1</sup> imidazole, 1 mmol L<sup>-1</sup> PMSF, pH 8.0) at 4 °C. The purity of the purified protein was estimated by SDS-PAGE in the eluted fractions, using 12% polyacrylamide running gels.<sup>29</sup> The purity of the enzyme was estimated by SDS-PAGE. The protein concentration was determined using the Bradford method.<sup>30</sup> The purified sample was supplemented with 50% glycerol and stored at -20 °C until use.

#### Conjugation of quantum dots and aptamer

The amine group of RNA aptamer was first covalently conjugated onto the surface of the carboxyl terminated QDs605 (10 pmol  $L^{-1}).$  That is, 10 pmol  $L^{-1}$  of QDs605 were conjugated with 400 pmol  $L^{-1}$  of aptamer with the coupling reagent EDC (N-(3-dimethylaminopropyl)-N-ethyl-carbodiimide hydrochloride, 40 nmol  $L^{-1}$ ), which was used to activate amide bond formation to produce QDs-conjugated aptamer (QDs based SARS-CoV N aptamer) at a QDs: aptamer molar ratio of 1:40 for 1 h at room temperature. Then, QDs-aptamer conjugate was collected using centrifugal filtration at 15 000 rpm for 30 min, followed by several washing steps with a Tris buffer (50 mmol  $L^{-1}$  Tris-HCl (pH 7.4), 5 mmol L<sup>-1</sup> KCl, 100 mmol L<sup>-1</sup> NaCl, 1 mmol L<sup>-1</sup> MgCl<sub>2</sub>, and 0.1% NaN<sub>3</sub>). After centrifugal filtration and washing, the pellet of QDsaptamer was dispersed by brief sonication (22 kHz, amplitude 12 µm, and sonication time 120 s) using a Sonic Dismembrator Model F60 (Fisher Scientific, Fair Lawn, NJ, USA).

#### Confocal laser scanning microscopy

The recombinant SARS-CoV N protein was directly immobilized onto the functional ProLinker<sup>™</sup>-terminated surface. For the binding of the specific aptamer, the conjugated QDs-conjugated aptamer was facilitated by spotting on immobilized SARS-CoV N protein chip. After incubation for 1 h at 25 °C, the chip was then washed three times with phosphate buffer (pH 7.2) for 1 min. The chip was analyzed using a confocal laser scanning microscope



Figure 1. A representative scheme for detection of SARS-CoV N protein on chip: (A) scheme of SARS-CoV N protein assay on chip; (B) total procedures of SARS-CoV N protein assay.

LSM510 META (Carl Zeiss, Jena, Germany). The signal intensity was determined by software for the LSM510 (LSM Image Browser). A histogram of the intensity was obtained for the region of the spotted chip. The value of signal intensity was recorded calculating and expressing it as the mean intensity.

# **RESULTS AND DISCUSSION**

## A schematic outline of aptamer chip assay

To ensure the feasibility of the targeting and imaging, we designed QDs605 conjugates having RNA aptamer for specific SARS-CoV N protein. We used a highly sensitive chip using a ProteoChip coated with ProLinker<sup>™</sup>, novel calixcrown derivatives with a bi-functional coupling property that permits the efficient immobilization of captured SARS-CoV N protein on solid matrixes and makes simple analysis of protein – RNA interactions possible.<sup>31</sup> Specific detection of the SARS-CoV N protein was demonstrated by the QDsconjugated RNA aptamer on chip. The total procedure for the detection on chip was as follows. First, the SARS-CoV N protein was spotted on a glass chip with carboxyl as the functional group. Second, the QDs-conjugated RNA oligonucleotide was bound on an immobilized chip. Third, the bound chip was washed to remove unspecific binding. Fourth, it was analyzed to recognize the specific detection of SARS-CoV N on chip. The design of targeting for effectively monitoring the SARS-CoV N protein on chip is illustrated in Fig. 1.

#### Quantum dots (QDs)-conjugated aptamer

The QDs-supported RNA aptamer was conjugated in the reaction for the amide formation from the coupling of 5'-end-aminemodified RNA aptamer at the surface of QDs displaying carboxyl groups via standard EDC coupling. The QDs-conjugated RNA aptamer was confirmed on a 2.5% agarose gel at 100 V in TAE buffer. Agarose gel electrophoresis showed a band with mobility, showing a different band pattern between the free QDs and the QDs-conjugated RNA aptamer, confirming the formation of QDsconjugated RNA oligonucleotide. The mobility shifts are compared in Fig. 2. On agarose gel, the QDs-conjugated RNA aptamer showed less mobility shift than free QDs, thus demonstrating the amide formation between QDs and RNA aptamer for conjugation.



**Figure 2.** The conjugation pattern on 2.5% agarose gel electrophoresis was examined at UV excitation wavelength 345 nm, showing a different band pattern between free QDs (lane 1) and QDs-conjugated aptamer (lane 2).

#### Sensitive detection of SARS-CoV N protein on chip

To improve the solubility of bacterially overexpressed recombinant N protein, the co-expression of N protein with E. coli molecular chaperone GroES/EL was performed. The SARS-CoV N protein was purified by a single chromatography step on a Ni<sup>2+</sup> affinity column. The C-terminally his-tagged SARS-CoV N protein was visualized with a molecular mass of approximately 48 kDa on a SDS-PAGE (Fig. 3(A)). Figure 3(B) shows the sequence and secondary structure of RNA aptamer that binds to SARS-CoV N protein. The RNA secondary structure of the used aptamer was analyzed using the Mfold program.<sup>32</sup>

We demonstrated the specific interaction between QDsconjugated RNA aptamer and immobilized SARS-CoV N protein on a chip. Preferential binding of QDs-conjugated RNA aptamer for the detection of SARS-CoV N protein was determined by analyzing the fluorescence intensity measured by confocal microscopy. As illustrated in Fig. 4(B), the QDs-supported RNA aptamer conjugates showed high fluorescence signals on the chip. No fluorescence signal was detected with BSA protein, due to its lack of affinity (Fig. 4(A)). The signal of QDs-conjugated RNA oligonucleotide for BSA protein binding affinity was very similar to the background signal. Figure 4(C) and 4(D) shows that QDs-conjugated RNA aptamer is selective against SARS-CoV N protein due to protein concentrations. The binding affinity was examined by confocal assay with varying concentrations of the SARS-CoV N protein. It was observed that the signal intensity of the QDs-conjugated



**Figure 3.** Purification of SARS-CoV N protein: (A) 12% SDS-PAGE gel showing SARS-CoV N protein with his-tag. lane 1, protein marker; lane 2, his-tag form of SARS-CoV N protein; (B) sequence and secondary structure of RNA aptamer that binds to N protein.

RNA aptamer could be detected even at the lowest concentration  $(0.1 \text{ pg mL}^{-1})$ . The signal intensity was found to increase gradually by up to  $0.1 \text{ pg mL}^{-1}$  of SARS-CoV N protein concentration. These results suggest the QDs-conjugated RNA aptamer could specifically target the SARS-CoV N protein on an immobilized chip. The detection limit was improved at least 10-fold compared with the aptamer-based chemiluminescence immunosorbent assay.<sup>28</sup> Furthermore, this system may be useful for rapid detection of SARS-CoV N protein within 1 h. These results suggest that the QDs-conjugated RNA oligonucleotide could be effective for the

quantification of SARS-CoVN protein. The immobilized BSA protein chip showed no binding to RNA aptamer, thus demonstrating selectivity for SARS-CoVN protein. The remarkable imaging by semiconductor nanocrystals QDs allows for monitoring with the intense light emission and accuracy.

In summary, we showed that QDs-conjugated RNA oligonucleotide can specifically recognize the SARS-CoV N protein. We demonstrated that this QDs-conjugated RNA aptamer could interact on a designed chip specifically and sensitively. This device could form a QDs-conjugated biosensor prototype chip for SARS-CoV N protein diagnosis. The proposed visual SARS-CoV N protein detection technique may avoid the limitations of other reported methods because of its high sensitivity, good specificity, simplicity, and speed.

# CONCLUSIONS

The results of this study suggest that the QDs-conjugated RNA aptamer can recognize the SARS-CoV N protein with a detection limit of 0.1 pg mL<sup>-1</sup> on a designed chip. Thus, it could be developed as a sensitive diagnostic tool. Furthermore, this QDs-conjugated aptamer chip might have application potential as a simple and rapid diagnosis tool. This technique has potential applications in many fields, especially in multiple diseases detection.

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Figure 4. Detection of SARS-CoV N protein by QDs-conjugated aptamer assay. Difference in fluorescent emission was measured: (A) BSA; (B) SARS-CoV N protein. Effect of SARS-CoV N protein concentrations on binding assay: (C) spotted chip; (D) normalized signal intensity.

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