# Quantum dots-based multiplexed immunohistochemistry of protein expression in human prostate cancer cells

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Semiconductor quantum dots (ODs) are bright fluorescent nanoparticles that have been successfully used for the detection of biomarker expression in cells. The objective of the present study is to use this technology in a multiplexing manner to determine at a single cell level the expression of a cell-specific bio-marker, prostate-specific antigen (PSA) expressed by human prostate cancer LNCaP and ARCaP cell lines. Here we compared the sensitivity of immunohistochemistry (IHC) and OD-based detection of AR and PSA expression in these cell lines. Further, we conducted multiplexing QD-based detection of PSA and androgen receptor (AR) expression in LNCaP cells subjecting to androgen (R1881) stimulation. The involvement of AR in PSA regulation in LNCaP cells, at a single cell level, was confirmed by the co-incubation of LNCaP cells in the presence of both R1881 and its receptor antagonist, bicalutamide (Casodex). We showed here the superior quality of QDs, in comparison to IHC, for the detection of AR and PSA in cultured LNCaP and ARCaP cells. Multiplexing QDs technique can be used to detect simultaneously AR and PSA expression induced by R1881 which promoted AR translocation from its cytosolic to the nuclear compartment. We observed AR antagonist, bicalutamide, inhibited AR nuclear translocation and PSA, but not AR expression in LNCaP cells.

Key words: Multiplexed imaging, quantum dots, prostate cancer.

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mmunohistochemical detection of gene expression has been a widely used technique in cellular and molecular biologic studies. Recent advances in semiconductor quantum dot (QD) chemistry provided an alternative approach to detect gene expression in cells with high degree of sensitivity and specificity (Alivisatos et al., 2005; Pinaud et al., 2006). QD nanoparticles tagged with antibody (Chan and Nie, 1998), proteins or peptides (Zhou et al., 2007), enzyme substrates and organic molecules ((Wu et al., 2003) have been successfully used in the detection of gene expression (Alivisatos et al., 2005) and movement of cells (Michalet et al., 2005) and the delivery of drugs (Yezhelyev et al., 2006) in biologic systems. Because of the sensitivity of fluorescent QD, their photostability and tunable nature (a broad-spectrum of light excitation of QD can result in the emission of multiple sharp and narrow spectra representative of QD of different sizes and chemical composition (Smith et al., 2006) for the detection of biomolecules and cells in an organism, QD have become the preferred substrates over organic dyes for tagging cells or cellular components in biologic studies (Michalet et al., 2005; West et al., 2003). In several recent reports (Yezhelyev et al., 2006; Chan et al., 2005), QD have been employed in IHC for the detection of gene expression and localization of key regulatory molecules in cells and tissues. The present study focused on employing the distinct advantages of QD, including its sensitivity and multiplexing capability, to simultaneously detect androgen receptor (AR) and its target gene, prostate-specific antigen (PSA) expression in human prostate cancer LNCaP and ARCaP cells. We compared IHC and QD-based methods in the detection of AR and PSA in cells with either high (LNCaP) or low (ARCaP) levels of expression of AR and PSA. We also employed a multiplexing QD-based technique to detect and quantify AR and PSA simultaneously with cells

either stimulated by androgen (R1881) or in the presence of both R1881 and its antagonist, bicalutamide (Casodex) to assess the involvement of AR in PSA expression. In sum, we validated that QD-based gene expression methods are superior to IHC and can be used to detect gene expression in a single cell in a multiplexing manner.

#### **Materials and Methods**

#### **Antibodies**

Quantum dot secondary antibody conjugates were from Invitrogen, (Carlsbad, CA) including QD-655nm, QD-605nm, and QD-565nm anti-mouse IgGs. Monoclonal antibodies were anti-human prostate specific antigen (PSA) and anti-human androgen receptor (AR) from Santa Cruz Biotechnology (Santa Cruz, CA) and anti-human elongation factor-1 alpha (EF-1 $\alpha$ ) from Upstate (Billerica, MA). Androgen analog R1881 and fetal bovine serum (FBS) were from Sigma (St. Louis, MO). Androgen antagonist, bicalutamide (Casodex) was obtained from AstraZeneca (Los Angeles, CA). Tissue culture T medium was purchased from Life Technologies (Rockville, MD).

#### **Cell lines and cell culture**

Human prostate cancer cell lines LNCaP and ARCaP were cultured in 8-well chamber slides (Nalge Nunc International, Rochester, NY) in T-medium supplemented with 5% FBS and 1% penicillin/streptomycin as described previously (Gleave, et al., 1991). The cells were maintained at 37°C in 5% CO<sub>2</sub> and 95% air. After culture in T-medium with 5% FBS for 24 hours, the culture medium was replaced with phenol red free RPMI 1640 with or without 5 nM R1881 in the presence or absence of its antagonist 10 uM Casodex for 48 hours. The cells were rinsed briefly with cold PBS and fixed in -20°C acetone for 10 min and were used for IHC or QD staining.

### Immunostaining with quantum dot probes

Acetone fixed cells were incubated with a blocking buffer containing 2% BSA/5% goat serum/PBS for 60 min at room temperature and were incubated with primary antibodies for one hour at room temperature. The antibody dilutions were PSA (1:50), AR (1:100), and EF-1 $\alpha$  (1:100). After rinsing 3 times with PBS, QD-IgGs (20 nM) were

applied to the slides and incubated overnight at 4°C or 2 hrs at room temperature. For multiplexed imaging, the three proteins were stained separately and concurrently. After the QD staining, cells were counter-stained with DAPI (20 ng/mL) for cell nuclei. For conventional IHC staining, the secondary antibodies were replaced by HRP labeled antimouse IgG using hydrogen peroxide as substrate and diaminobenzadine as chromogen, followed by hematoxylin nuclear counterstain.

### **Imaging and spectroscopy**

True-color fluorescence images were obtained with an inverted Olympus IX-70 microscope and a digital Nikon D70 color camera. Broad-band excitation in the near-UV range (330-385 nm) was provided by a 100 W mercury lamp. A longpass dichroic filter (DM 400, Chroma Technologies, Brattleboro, VT) was used to reject the scattered light and to pass the Stokes-shifted fluorescence signals. All images were captured using a 100X oil objective lens unless otherwise specified. Fluorescence spectra of the QD labeled single cells were collected using a spectrometer (Princeton Instruments, Princeton, NJ) coupled with WinSpec 32 software. For each sample, 40-50 cells were randomly picked for spectral analyses.

#### Western blot

Protein was extracted from cell pellets with a lysis buffer [50 mmol/L Tris (pH 8.0), 150 mmol/L NaCl, 0.02% NaN3, 0.1% SDS, 1% NP40, 0.5% sodium deoxycholate, and 1 mmol/L phenylmethylsulfonyl fluoride] in the presence of protease inhibitor cocktail (Roche Applied Science, Indianapolis, IN). Samples containing equal amounts of protein (30 µg) were electrophoresed on Tris-Glycine gels (Invitrogen) and transferred to nitrocellulose membranes. After blocking with Tween-tris buffered saline (T-TBS) containing 5% nonfat milk powder, membranes were incubated separately and sequentially with mouse monoclonal antibodies against either PSA or AR at 4°C overnight. After washing with T-TBS, the membranes were incubated with corresponding secondary antibodies, which were conjugated with horseradish peroxidase (Santa Cruz Biotechnology). The blots were stripped by a standard protocol used in our laboratory. In brief, the membranes were placed in a heated (50°C) stripping buffer (2% SDS, 100 mM β- mercaptoethanol and 50 mM Tris, pH 6.8)

for 15-30 min with gentle shaking. The membranes were rinsed with TBS several times and the blots were ready for re-blot with anti- EF-1 $\alpha$  antibody (1:1000 dilution). Immunoreactive bands were visualized with enhanced chemiluminescence (Amersham Pharmacia Biotech, Little Chalfont, United Kingdom) using previously established protocols (Huang *et al.*, 2006; Nomura *et al.*, 2006; Wu *et al.*, 2007).

### **Deconvolution of composite spectra and quantitative analysis**

The photostability and the narrow and Gaussianshaped emission spectra of QDs make it possible to use QDs for absolute quantification of targets and it has been documented that there is a linear relationship between the measured fluorescence intensity and the number of stained QDs (Arya et al., 2005). Based on this assumption, the Gaussian function is chosen to represent each peak in the spectra. The Gaussian function is represented as:  $f(x)=\alpha e^{-(x-b)^2/c^2}$  where a, b, and c are real constants and a > 0, c > 0; the improper integrals over the whole real line can be evaluated exactly:  $\int_{-\infty}^{+\infty} F(x) =$  $\sqrt{\pi}$ ac. Each spectrum can be represented as a sum of *n* peaks: spectrum =  $\sum_{l=1}^{n}$  peak. A linear combination of QD spectra was used to fit the composite photoluminescence signal for each sample. Thus, the composite signal can be calculated as:  $I_{total}(\lambda) = \sum_{l=1}^{n} I_{l}(\lambda) = \sqrt{\pi} \sum_{l=1}^{n} a_{i}$ ,  $c_{i}$  where  $a_{i}$ ,  $c_{i}$  are the parameters in the Gaussian function obtained from fitting the ith peak in the spectrum,  $I_{total}(\lambda)$  is the composite signal (corrected for background signal). The area ratio between ith and ith peaks can be easily obtained by  $r = \frac{\partial_i C_i}{\partial_i C_i}$ . In this work, we will concentrate on the QD samples (565, 605, and 655 nm emission maxima). The 565 nm peak will be used as the reference. A software package was developed in Matlab that can fit QD spectra with a minimum of 1 to a maximum of 8 peaks for a single spectrum or multiple spectra in a set. The areas under each fitted peak and the ratios of peaks at 605 nm to 565 nm and 655 nm to 565 nm are also given. For quantitative analysis, the relative expression level of the target gene was calculated as the ratio of the fluorescence intensity of the target gene to the intensity of the control gene. Statistical analysis was performed by Student's t test. The significant level  $\alpha$ =0.05 was used in the study.

#### **Results**

# Optical imaging of low-abundant genes in prostate cancer cells using quantum dots

We determined the sensitivity of QDs-antibody conjugates in the detection of low abundant AR and PSA in ARCaP cells. As shown in Figure 1A, clear fluorescence signals for both AR and PSA proteins were detected in ARCaP cells. PSA was found to be predominately cytoplasmic. AR was found to be present in both cytoplasmic and nuclear compartments. In a parallel study, however, the presence of AR and PSA in ARCaP cells was less apparent when stained by IHC using the same primary antibodies (Figure 1A). One other advantage of QDbased IHC is that under these assay conditions, the fluorescent signal intensity was stable for several hours. These results indicate that the QDs-combined IHC method can detect the expression of genes of low abundance which cannot be clearly discerned using the conventional IHC methods.

# Imaging of ligand-induced androgen receptor translocation and PSA expression in LNCaP cells

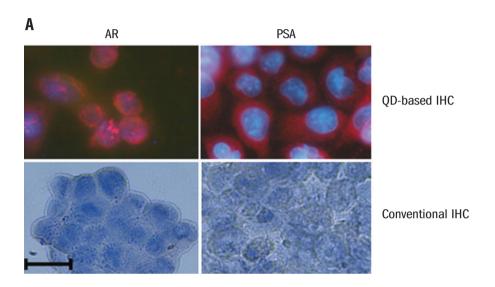
Since QD staining proved to be more sensitive than conventional IHC in detecting low-abundant genes, we next analyzed the relationship between AR nuclear translocation and PSA expression in androgen-responsive LNCaP cells. After 24 h attachment to the plastic dishes, replaced with phenol red free RPMI 1640 medium in the absence of serum, cells were exposed to an androgen analog, R1881, as previously described (Mizokami et al., 2000). As shown in Figure 1B, AR nuclear translocation was clearly visible in LNCaP cells after treatment with R1881 for 48 hours. In control cells without R1881 treatment, AR staining was mostly in the cell cytoplasm. Since PSA expression is known to be under regulation by liganded AR, we next examined PSA expression in LNCaP cells under stimulation by R1881 in the presence of an androgen antagonist, Casodex. Results showed that up-regulation of PSA expression in LNCaP cells by R1881 was blocked by Casodex (Figure 1B). These results were further confirmed by Western blot analysis (Figure 1C), suggesting QD-antibody conjugates are suitable for the detection of gene expression in prostate cancer cells. These observations are in agreement with our previously published reports where antiandrogen such as Casodex or flutamide antagonized PSA expression in

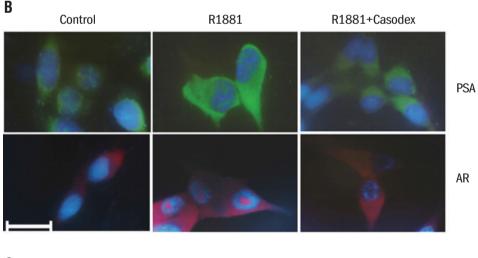
LNCaP cells stimulated by androgen (Thalmann et al., 1996; Kiyama et al., 2003; Yeung et al., 2000).

# Multiplexed quantum dots imaging of gene expression in a single cell

Since LNCaP cells express both AR and PSA, we explored the possibility of using this cell line as target for androgen-induced responses by monitoring

AR and PSA simultaneously. Based on the fluorescent characteristics of quantum dots, this makes it an ideal candidate for multiplexing AR and PSA in LNCaP cells. We performed multiplexed staining of AR, PSA and EF-1 $\alpha$ , an internal standard, in LNCaP cells subjected to R1881 treatment either in the presence or absence of an androgen antagonist, Casodex. Figure 2A showed that multiplexed LNCaP





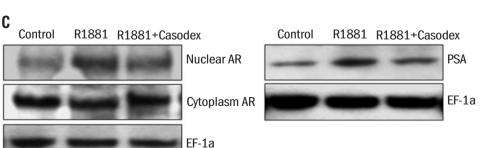


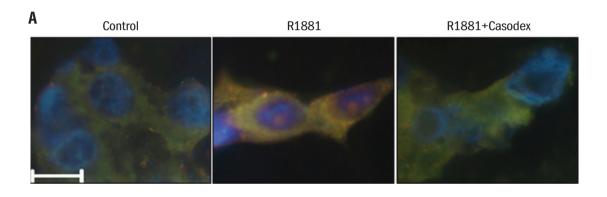
Figure 1. Optical imaging of protein markers of prostate cancer cells using quantum dots combined with immunohistochemical staining.

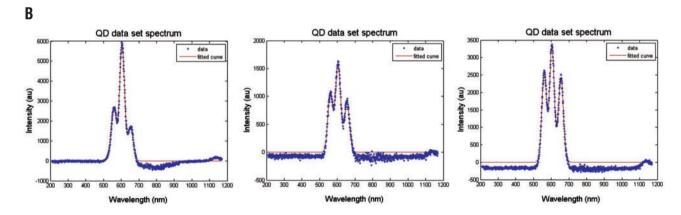
(A) Comparative study of

- the detection sensitivity of AR and PSA in ARCaP cells between conventional IHC and QD-based IHC. OD605 and OD655 were used to label PSA and AR, respectively. Bar, 20 um. (B) Optical imaging using OD-based IHC showed that AR nuclear translocation and PSA expression in LNCaP cells was stimulated by androgen analog, R1881. Casodex, androgen antagonist, blocked the stimulation of PSA expression induced by R1881. QD605 was used to label PSA and QD655 was used to label AR. Bar, 20 um. (C) Western blot analysis
- confirmed the AR nuclear translocation and PSA expression in R1881-treated LNCaP cells.

cells can be imaged quantitatively, at a single cell level, for the expression of all three genes, AR, PSA and EF- $1\alpha$ , using QDs with different emission spectra. We observed AR nuclear translocation in LNCaP cells after R1881 treatment. In order to perform the quantitative analysis, a software program was developed to monitor the gene expression based on fluorescence intensity of the spectrum for the corresponding marker gene. The spectrum files for single cells were obtained using the spectrometer under the fluorescence microscope and the data were processed using the WinSpec 32 program (see

Materials and Methods). As shown in Figure 2B, the fitting spectrum curve generated from the software program matched very well the spectrum curve generated from the real data. We next performed quantitative analysis based upon the ratios of each gene expression to that of the control gene. As shown in Figure 2C, R1881 treatment increased PSA expression in LNCaP cells, but there was no significant increase in the expression of total AR. In the presence of an androgen antagonist, Casodex, it blocked the ability of AR to induce PSA expression without affecting AR expression at the single cell level.





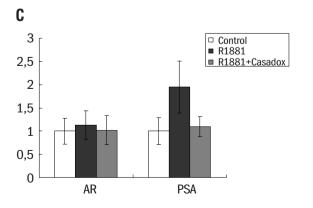


Figure 2. Multiplexed imaging and profiling of protein markers in LNCaP cells. (A) Triple stain of AR, PSA and EF-1 $\alpha$  in single LNCaP cells using QD with QD655, QD605 and QD565, respectively. QD655 (red) was used to label AR, QD605 (orange) was used to label PSA and QD565 (green) was used to label EF-1 $\alpha$ . Bar, 20 um. (B) Spectra analysis of the gene expression by the spectrometer. A software program was developed in Matlab that can fit the QD spectra with different peaks from the spectrometer. (C) Quantitative analysis of target gene expression was based on the ratio of the fluorescence intensity of target gene to control gene, EF-1 $\alpha$ . indicates the significant difference of PSA expression between R1881 treatment group and control group.

#### **Discussion**

The present study demonstrated for the first time the use of QD multiplexing technology for the detection of the expression of an androgen target gene, PSA, in a human prostate epithelial cell line, LNCaP, under the influence of androgen and its antagonist, Casodex. We monitored AR and PSA simultaneously with a house-keeping gene, EF-1 $\alpha$ , as an internal standard. We found that QD immunostaining of gene expression is superior to conventional IHC with respect to its sensitivity. Though ARCaP cells express low levels of AR and PSA, these markers can be detected readily by QDbased but not chromogen-based assays using identical monoclonal antibodies (Figure 1). We found that the multiplexing QD-based gene expression study yielded the expected results at the single cell level where PSA is regulated transiently by the addition of R1881, and this effect can be blocked by the simultaneous presence of an anti-androgen Casodex (Figure 2).

There are several reasons why QD-based multiplexing technology could have unique roles in biology: 1) QD are bright and more resistant to photobleaching than the conventional organic fluorescent dyes; thus they are more efficient in detecting particularly low abundant target molecules in cells (Wu et al., 2003). 2) QD can be excited at a broad range of wavelengths but their emission spectra are specific with sharp and narrow peaks, dependent on their sizes and chemical compositions, with minimal overlap; thus they can be readily multiplexed and quantified (Gao et al., 2007). 3) QD can be modified chemically and linked to various targeting moieties for the detection and delivery of desirable molecules to cells and tissues (Gao et al., 2003, 2004). In our study, we confirmed that QD-based detection of AR and PSA is superior to conventional IHC and that they can be multiplexed for simultaneous detection of both AR and PSA in LNCaP cells. Ness et al. (2003) have shown that QD-based IHC staining for certain intracellular antigens in tissue sections was more sensitive than conventional IHC approaches (Ness et al., 2003). For example, ARCaP cells express only a low level of AR and PSA (Zhau et al., 1996) which can be readily detected by QD-based IHC but not conventional IHC. While androgen-mediated PSA expression through AR has been studied extensively in LNCaP

cells, none of the previous studies used multiplexing methods to detect the activation of AR, as evaluated by its nuclear translocation on a single cell basis, coupled with PSA expression. This multiplexed imaging of target genes allows investigators to identify multiple target genes of interest for: 1) elucidating complex cell-cell or protein-protein interactions; 2) clarifying spatial arrangements of cells within a tissue; 3) localizing multiple proteins within a cell; and 4) identifying complex protein expression patterns in single cells. Since the introduction of biocompatible QD in 1998, the potential of QD for multiplexing was quickly realized (Pinaud et al., 2006). However, actual demonstrations of this capability were only recently begun. Examples include preliminary two-color fluoroimmuno assay, cellular labeling and microbead and microsphere QD encoded bar codes used in DNA hybridization. In our work, we stained AR, PSA and EF1- $\alpha$  gene in cultured LNCaP cells using quantum dots with three colors (655, 605, 565 nm). The staining of all the markers was confirmed by spectra analysis. Although multiplexing can be obtained using quantum dots with different emitting wavelengths in cells or tissue specimens, the key future application of this assay is the quantification of different gene expressions. In this work, we developed a software program that can separate the different spectra from the composite data. We used the total area of each spectrum to reflect the gene expression level using a housekeeping gene EF-1 $\alpha$  as an internal control. The ratio between the target gene and internal control gene was generated and used for statistical analysis. Using this protocol, we confirmed the relationship between AR and PSA in single prostate cancer cells. QD-based multiplexing technology, once adopted by the scientific community, could address many important regulatory biologic questions in cells.

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