

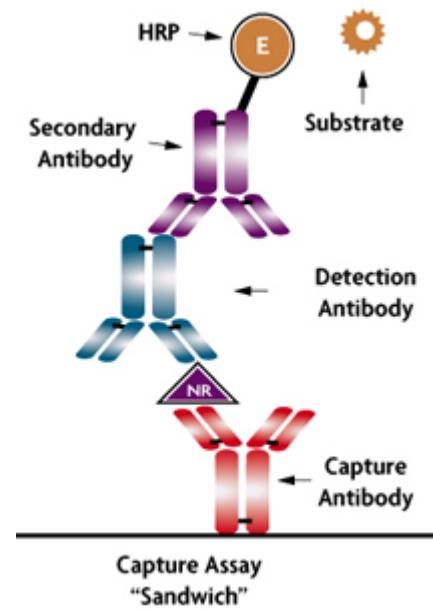
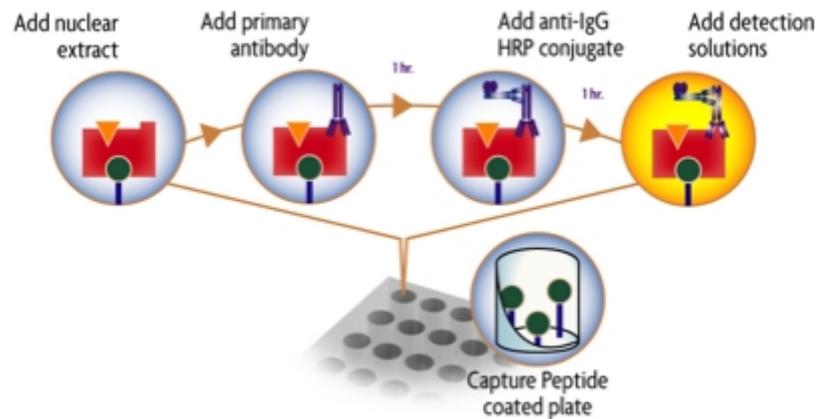
Nanomaterials for Biodiagnostic

- Nucleic Acid
 - Genetic information for identification
 - Diseases, bacterium, virus, pathogen
 - PCR with molecular fluorophore, State of the Art
 - Expansive, Non-portable, Non-multiplexing
- Proteins
 - Cancers and diseases, unusual high concentration of marker
 - ELISA (~pM) with molecular fluorophore
 - No PCR version



ELISA (Enzyme-Linked Immunosorbent Assay)

is a biochemical technique used mainly in immunology to detect the presence of an antibody or an antigen in a sample. It utilizes two antibodies, one of which is specific to the antigen and the other of which is coupled to an enzyme. This second antibody gives the assay its "enzyme-linked" name, and will cause a chromogenic or fluorogenic substrate to produce a signal.



Why Nanomaterials?

- Molecular fluorophores
 - Limited spectral response
 - photostability
- Nanomaterials
 - Small size (1-100 nm)
 - Chemically tailorabile physical properties
 - Unusual target binding properties
 - Structure robustness



Nanomaterial Detection

- Optical
- Electrical and electrochemical
- Magnetic
- Nanowire and Nanotubes
- Nanofabrication



Colorimetric Detection of DNA

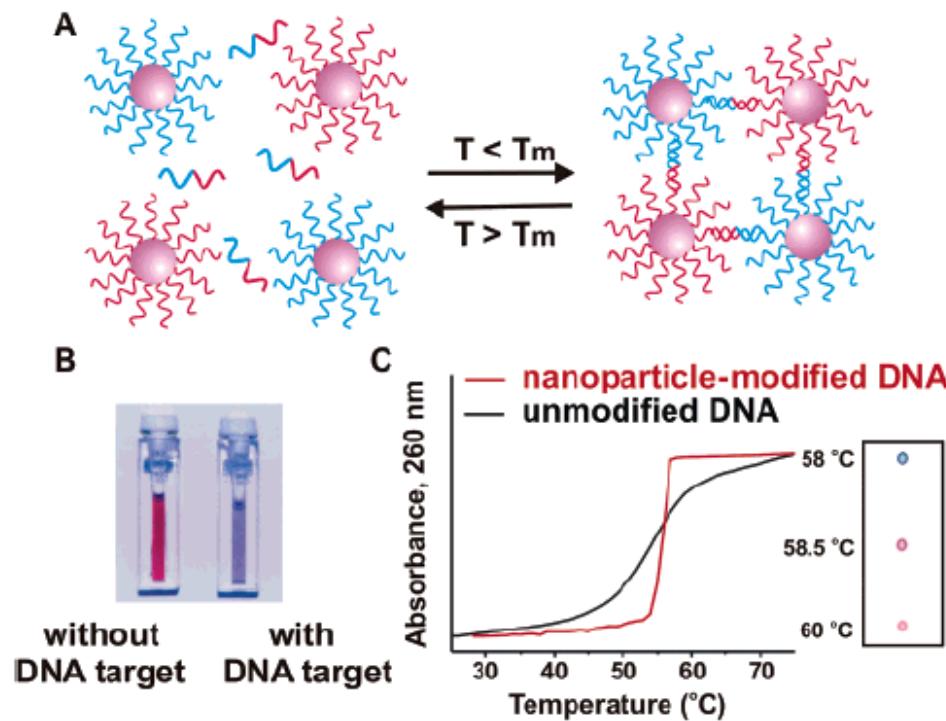


Figure 2. In the presence of complementary target DNA, oligonucleotide-functionalized gold nanoparticles will aggregate (A), resulting in a change of solution color from red to blue (B). The aggregation process can be monitored using UV-vis spectroscopy or simply by spotting the solution on a silica support (C). (Reprinted with permission from *Science* (<http://www.aaas.org>), ref 29. Copyright 1997 American Association for the Advancement of Science.)



A DNA-based method for rationally assembling nanoparticles into macroscopic materials

Chad A. Mirkin, Robert L. Letsinger, Robert C. Mucic
& James J. Storhoff

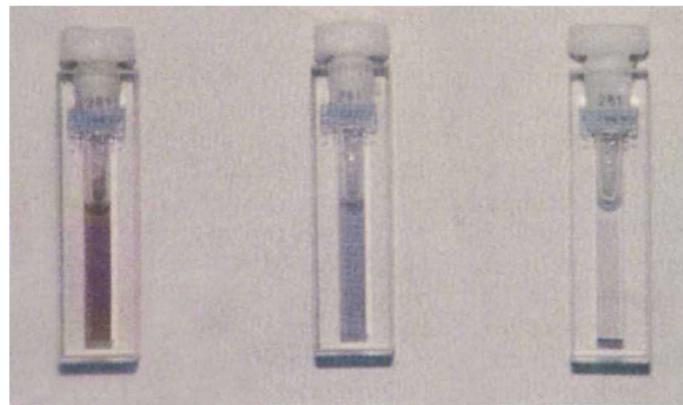
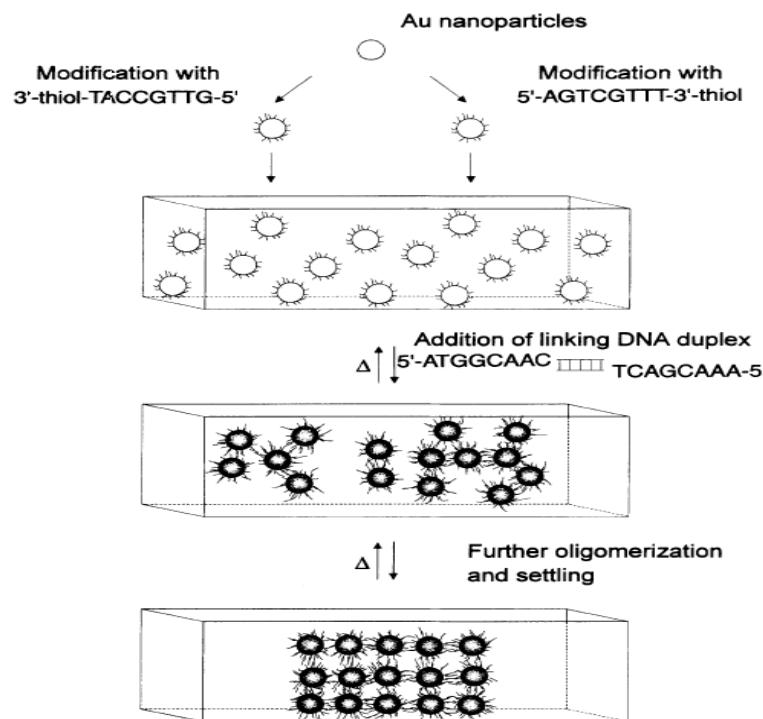
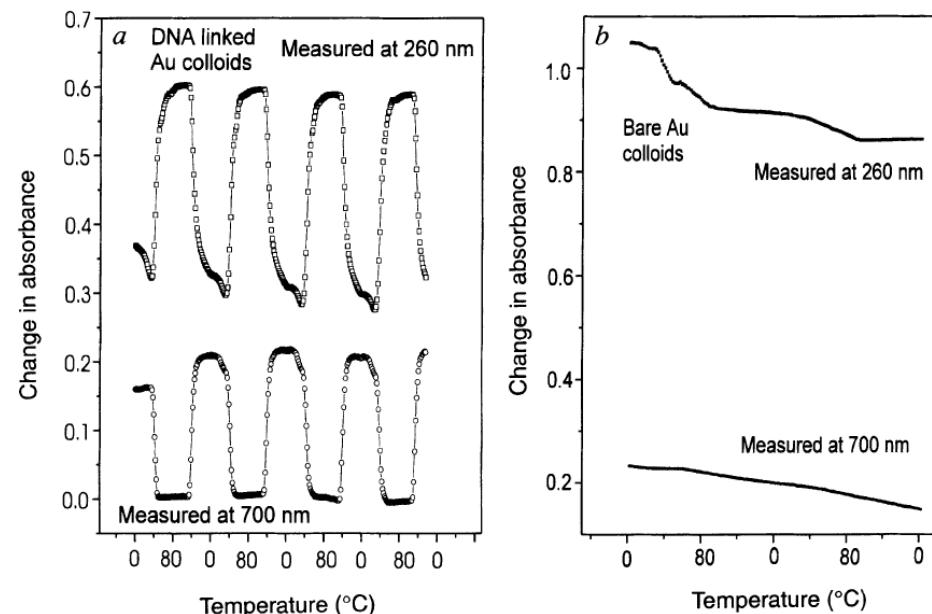


FIG. 2 Cuvettes with the Au colloids and the four DNA strands responsible for the assembly process. Left cuvette, at 80 °C with DNA-modified colloids in the unhybridized state; centre, after cooling to room temperature but before the precipitate settles; and right, after the polymeric precipitate settles to the bottom of the cuvette. Heating either of these cool solutions results in the reformation of the DNA-modified colloids in the unhybridized state (shown in the left cuvette).



Selective Colorimetric Detection of Polynucleotides Based on the Distance-Dependent Optical Properties of Gold Nanoparticles

Robert Elghanian, James J. Storhoff, Robert C. Mucic,
Robert L. Letsinger,* Chad A. Mirkin*

SCIENCE • VOL. 277 • 22 AUGUST 1997

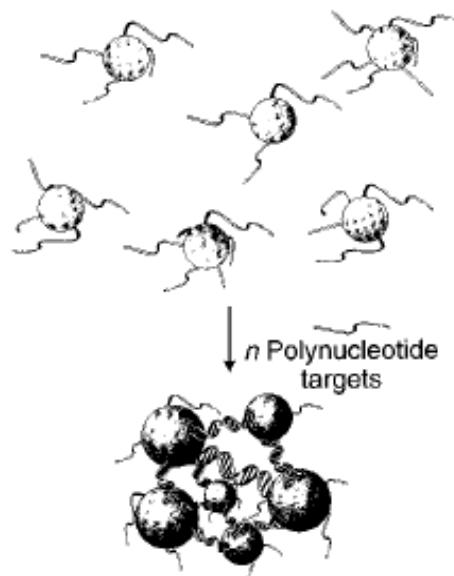


Fig. 1. Schematic representation of the concept for generating aggregates signaling hybridization of nanoparticle-oligonucleotide conjugates with oligonucleotide target molecules. The nanoparticles and the oligonucleotide interconnects are not drawn to scale, and the number of oligomers per particle is believed to be much larger than depicted.

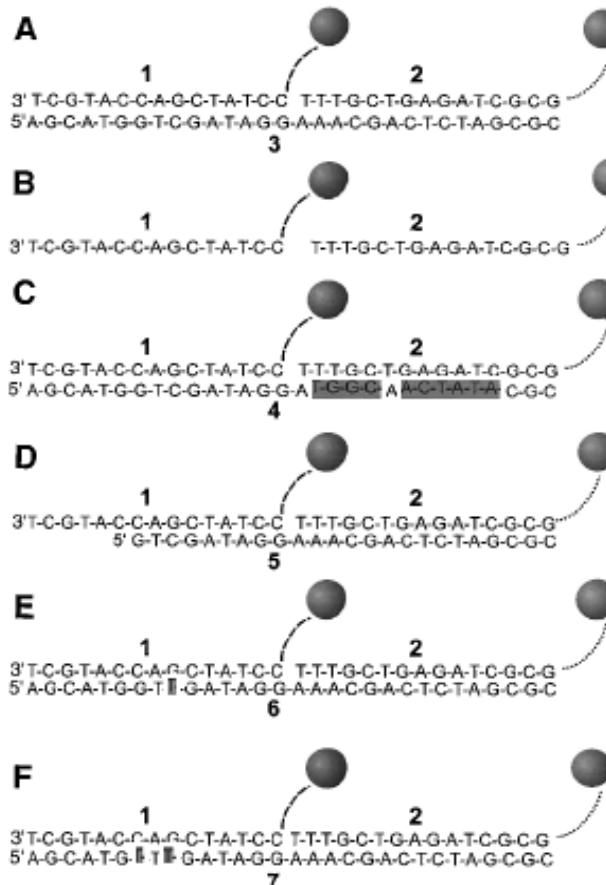


Fig. 2. Mercaptoalkyloligonucleotide-modified 13-nm Au particles and polynucleotide targets used for examining the selectivity of the nanoparticle-based colorimetric polynucleotide detection system. (A) Complementary target; (B) probes without the target; (C) a half-complementary target; (D) a 6-bp deletion; (E) a 1-bp mismatch; and (F) a 2-bp mismatch. For the sake of clarity, only two particles are shown; in reality a polymeric aggregate with many particles is formed. Dashed lines represent flexible spacer portions of the mercaptoalkyloligonucleotide strands bound to the nanoparticles; note that these spacers, because of their noncomplementary nature, do not participate in hybridization. The full sequences for the two probes, 1 and 2, which bind to targets 3 through 7, are

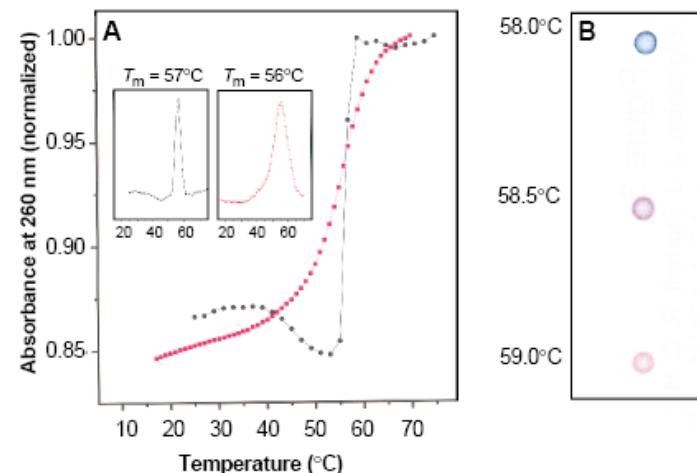
1-5'SH-(CH₂)₆-[CTA-ATC-CGC-ACA-G]
[CC-TAT-CGA-CCA-TGC-T]
probe

2-5'SH-(CH₂)₆-[ATG-GCA-ACT-ATA-C]
[GC-GCT-AGA-GTC-GTT-T]
probe



Fig. 3. (A) Comparison of the thermal dissociation curves for complexes of mercaptoalkyloligonucleotide-modified Au nanoparticles (black circles) and mercaptoalkyloligonucleotides without Au nanoparticles (red squares) with the complementary target, **3**, in hybridization buffer (0.1 M NaCl, 10 mM phosphate buffer, pH 7.0). For the first set (black circles), a mixture of 150 μ l of each colloid conjugate and 3 μ l of the target oligonucleotide in hybridization buffer (0.1 M

NaCl, 10 mM phosphate, pH 7.0) was frozen at the temperature of dry ice, kept for 5 min, thawed over a period of 15 min, and diluted to 1.0 ml with buffer (final target concentration, 0.02 μ M). The absorbance was measured at 1-min intervals with a temperature increase of 1°C per minute. The increase in absorption at 260 nm (A_{260}) was \sim 0.3 absorption units (AU). In the absence of the oligonucleotide targets, the absorbance of the nanoparticles did not increase with increasing temperature. For the second set, the mercaptoalkyloligonucleotides and complementary target (each 0.33 μ M) were equilibrated at room temperature in 1 ml of buffer, and the changes in absorbance with temperature were monitored as before. The increase in A_{260} was 0.08 AU. (**Insets**) Derivative curves for each set (15). (**B**) Spot test showing T_c (thermal transition associated with the color change) for the Au nanoparticle probes hybridized with complementary target. A solution prepared from 150 μ l of each probe and 3 μ l of the target (0.06 μ M final target concentration) was frozen for 5 min, allowed to thaw for 10 min, transferred to a 1-ml cuvette, and warmed at 58°C for 5 min in the thermally regulated cuvette chamber of the spectrophotometer. Samples (3 μ l) were transferred to a C₁₈ reverse phase plate with an Eppendorf pipette as the temperature of the solution was increased incrementally 0.5°C at 5-min intervals.



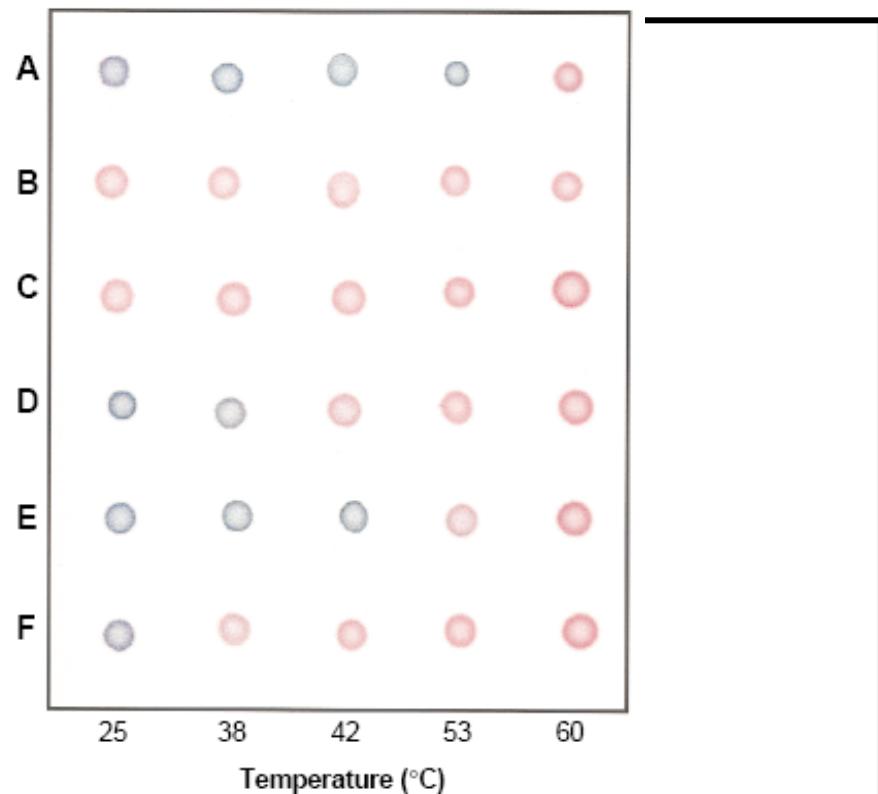
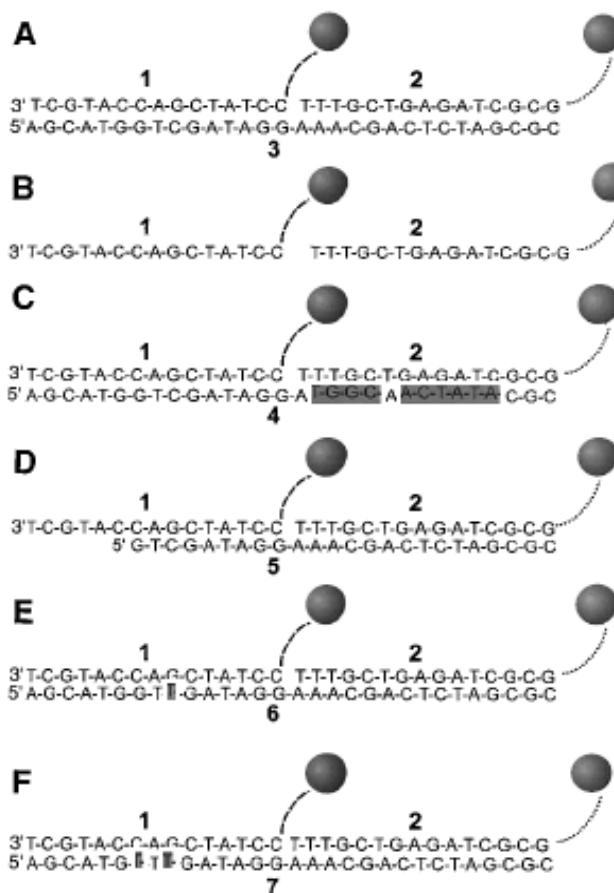


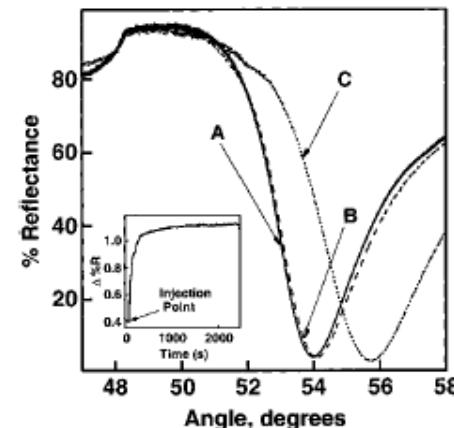
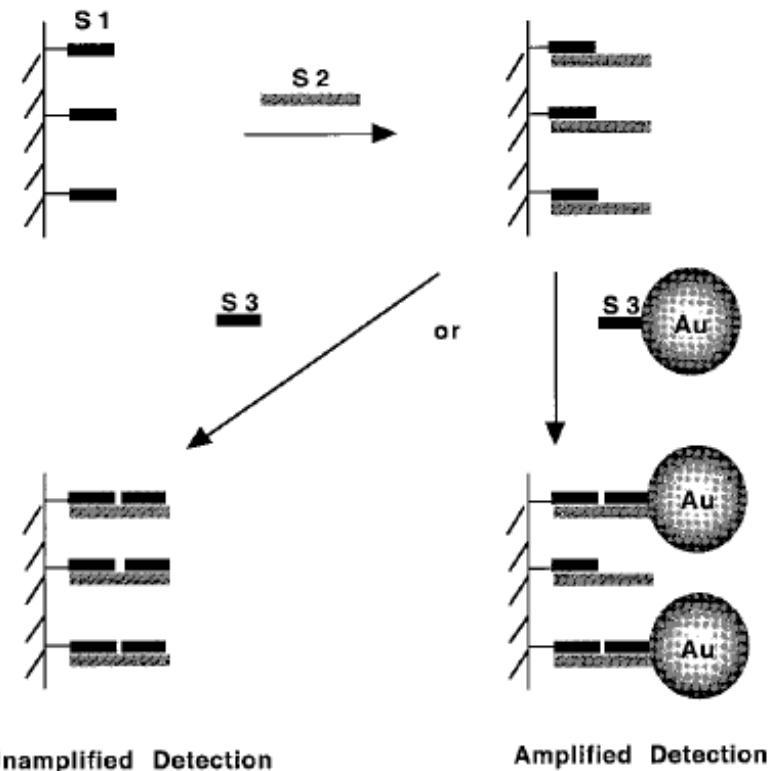
Fig. 4. Selective polynucleotide detection for the target probes shown in Fig. 2: (A) complementary target; (B) no target; (C) complementary to one probe; (D) a 6-bp deletion; (E) a 1-bp mismatch; and (F) a 2-bp mismatch. Nanoparticle aggregates were prepared in a 600- μ l thin-walled Eppendorf tube by addition of 1 μ l of a 6.6 μ M oligonucleotide target to a mixture containing 50 μ l of each probe (0.06 μ M final target concentration). The mixture was frozen (5 min) in a bath of dry ice and isopropyl alcohol and allowed to warm to room temperature. Samples were then transferred to a temperature-controlled water bath, and 3- μ l aliquots were removed at the indicated temperatures and spotted on a C₁₈ reverse phase plate.



Colloidal Au-Enhanced Surface Plasmon Resonance for Ultrasensitive Detection of DNA Hybridization

Lin He, Michael D. Musick, Sheila R. Nicewarner, Frank G. Salinas, Stephen J. Benkovic, Michael J. Natan, and Christine D. Keating*

Scheme 1. SPR Surface Assembly



Scheme 2. SPR Surface Assembly in the Digestion Experiment

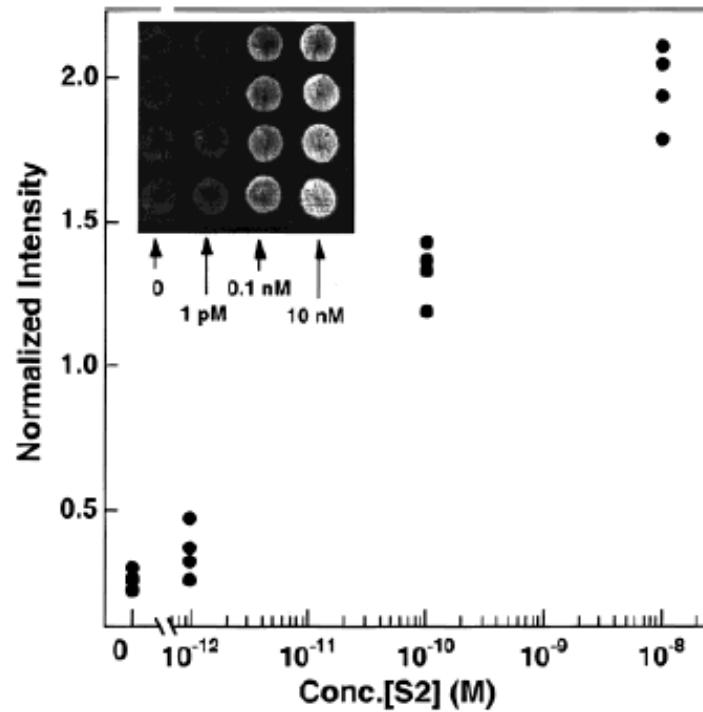
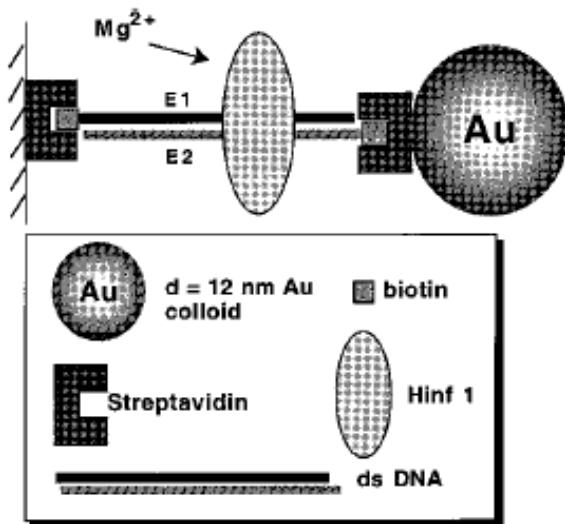


Figure 5. Plot of normalized intensity of SPR reflectance as a function of logarithmic concentration of the analyte 24-mer oligo (S2). Each spot represents one data point at the corresponding concentration. CCD parameters: exposure time = 0.3 s, 16 bit resolution, spot size = 4.5 mm in diameter. Inset: a 2-D SPR image of a Au surface derivatized with 20 μ L of buffer blank, 1 pM, 0.1 nM, and 10 nM S2 oligos (from left to right, respectively).



Self-Assembled Nanoparticle Probes for Recognition and Detection of Biomolecules

Dustin J. Maxwell, Jason R. Taylor, and Shuming Nie*,†

9606 ■ J. AM. CHEM. SOC. 2002, 124, 9606–9612

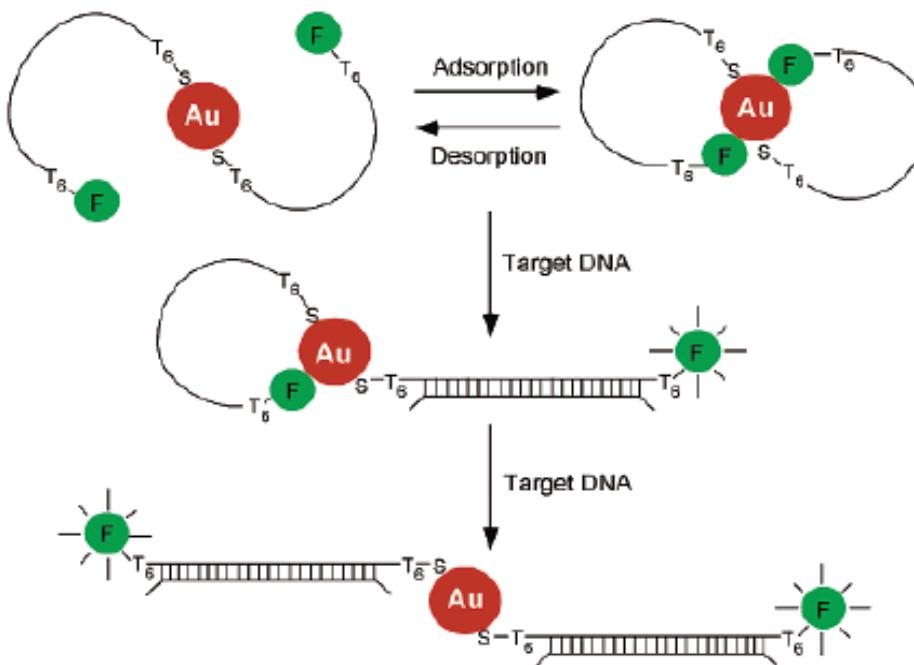


Figure 1. Nanoparticle-based probes and their operating principles. Two oligonucleotide molecules (oligos) are shown to self-assemble into a constrained conformation on each gold particle (2.5 nm diameter). A T₆ spacer (six thymines) is inserted at both the 3'- and 5'-ends to reduce steric hindrance. Single-stranded DNA is represented by a single line and double-stranded DNA by a cross-linked double line. In the assembled (closed) state, the fluorophore is quenched by the nanoparticle. Upon target binding, the constrained conformation opens, the fluorophore leaves the surface because of the structural rigidity of the hybridized DNA (double-stranded), and fluorescence is restored. In the open state, the fluorophore is separated from the particle surface by about 10 nm. See text for detailed explanation. Au, gold particle; F, fluorophore; S, sulfur atom.



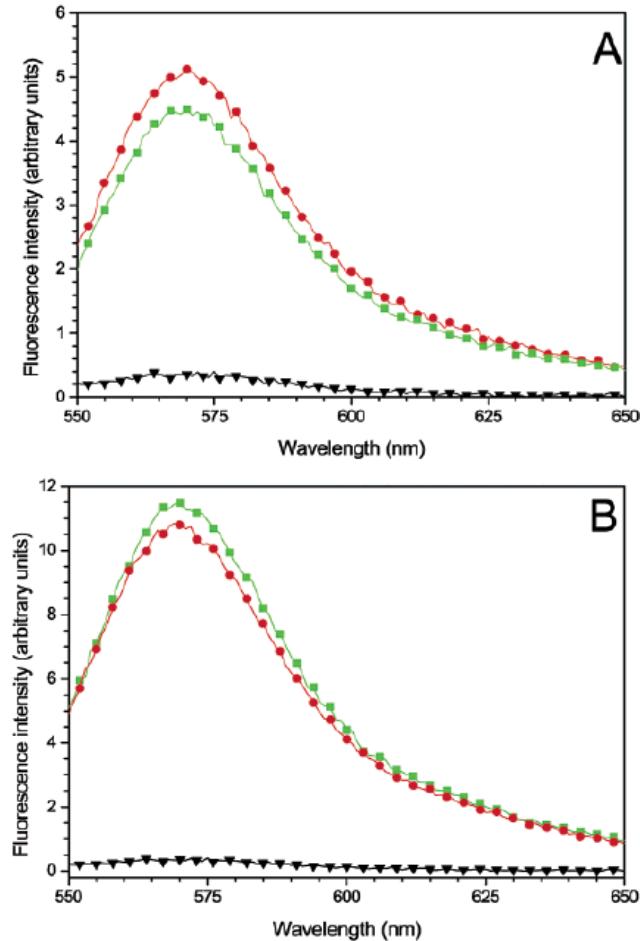


Figure 5. Fluorescence responses and the lack of sequence recognition abilities observed for nonthiolated nanoparticle probes. (A) Fluorescence spectra of nonthiolated probes generated by a complementary target (red curve), a noncomplementary target (green curve), and no target (black curve). These probes are considered nonfunctional because they do not recognize specific DNA sequences. (B) Fluorescence signals obtained from the supernatant solution when the probes were treated with a complementary target (red curve) or a noncomplementary target (green curve). The result revealed that the oligos were released into solution by nonspecific adsorption of the target on the particle surface. With a thiol group, this release was not observed (little or no signal in solution, black curve in B). The nonfunctional probes were prepared in the same way as the functional probes, except that the 3'-end thiol group was deleted. The intensity differences for the red and green curves were within experimental errors and had no particular significance.

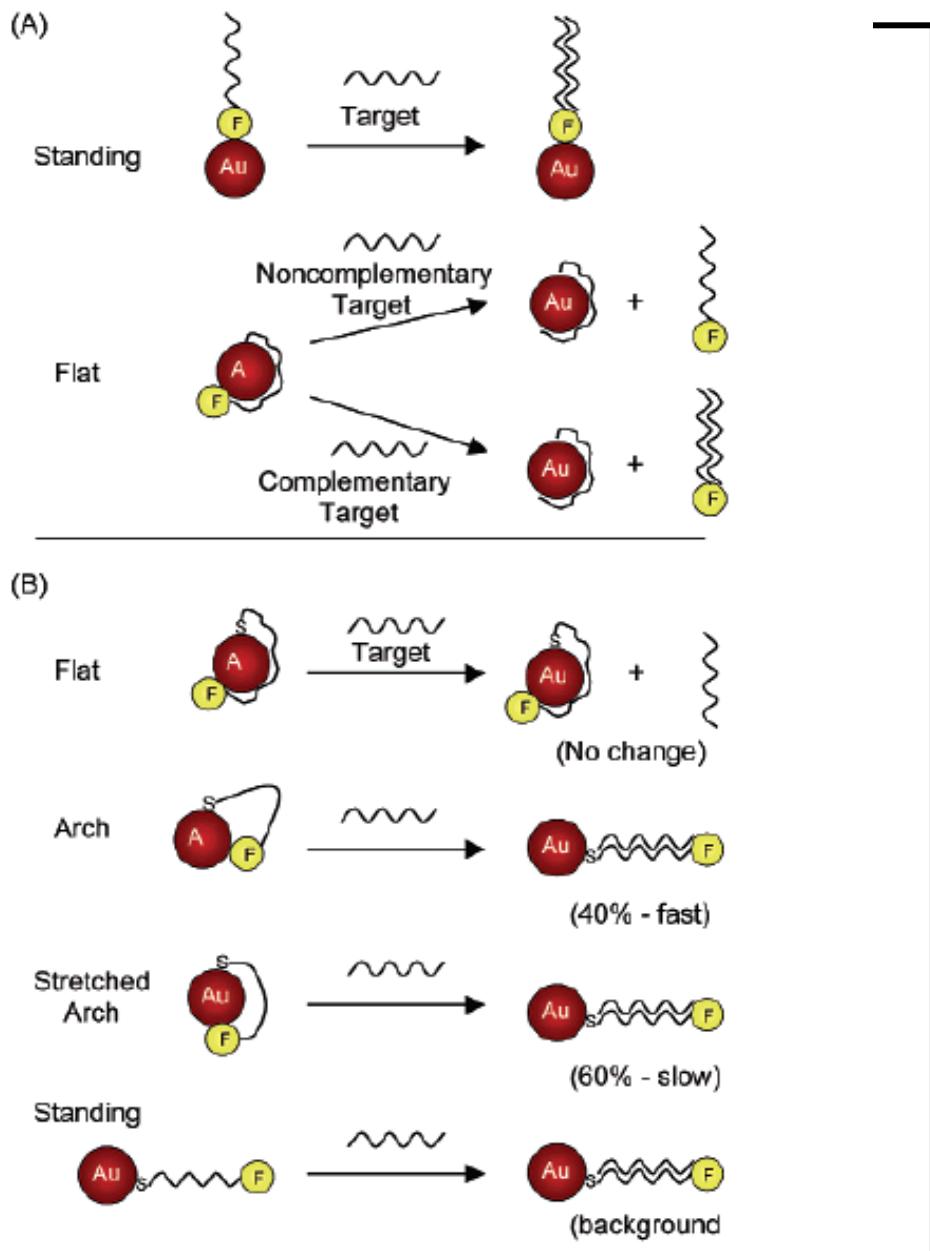


Figure 6. Schematic illustration of possible configurations for (a) nonthiolated and (b) thiolated oligonucleotides adsorbed on colloidal gold nanocrystals. Detailed discussion in text.



Silver Amplification

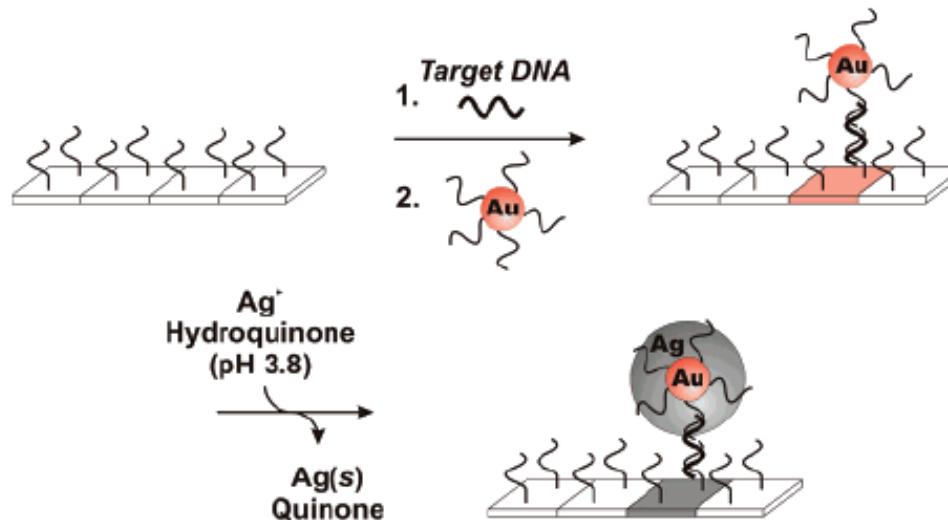


Figure 4. Scanometric DNA assay. In this assay a surface-bound capture oligonucleotide binds one-half of the target of interest, and an oligonucleotide-functionalized gold nanoparticle probe binds to the other half. Catalytic reduction of silver onto the capture/target/probe sandwich results in a signal that can be detected scanometrically. (Reprinted with permission from *Science* (<http://www.aaas.org>), ref 66. Copyright 2000 American Association for the Advancement of Science.)

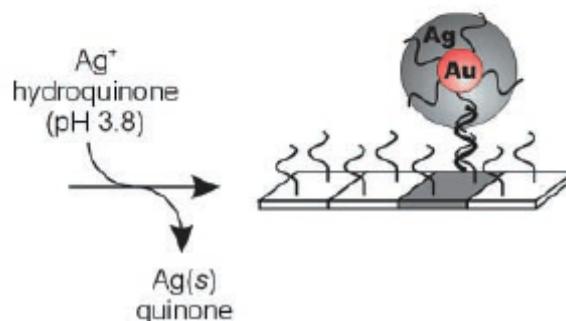
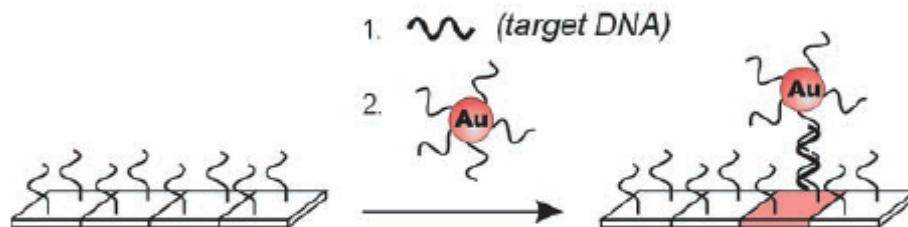
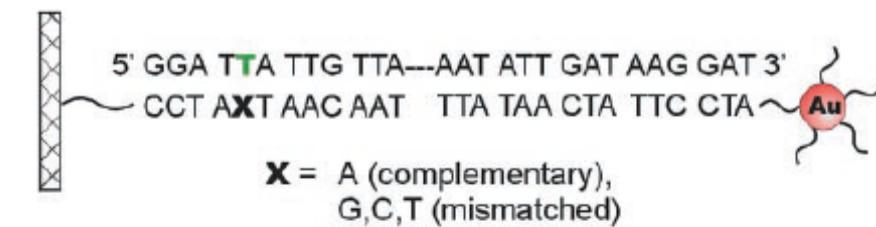
Catalytic reduction of Ag on Au



Scanometric DNA Array Detection with Nanoparticle Probes

SCIENCE VOL 289 8 SEPTEMBER 2000

T. Andrew Taton,^{1,2} Chad A. Mirkin,^{1,2*} Robert L. Letsinger^{1*}



50 fM => 0.2 fM



Fig. 1. Images of 7 mm by 13 mm, oligonucleotide-functionalized, float glass slides, obtained with a flatbed scanner. (A) Slide before hybridization of target and nanoparticle probe. (B) A slide identical to (A) after hybridization with oligonucleotide target (10 nM) and then nanoparticle probes (5 nM in particles). The pink color derives from the Au nanoparticle probes. (C) A slide identical to (B) after exposure to silver amplification solution for 5 min. (D) Slide before hybridization of target and nanoparticle probe. (E) A slide identical to (D) after hybridization with target (100 pM) and then nanoparticle probe (5 nM). The extinction of the sub-monolayer of nanoparticles is too low to be observed visually or with a flatbed scanner. (F) A slide identical to (E) after exposure to silver amplification solution for 5 min. Slide (F) is lighter than slide (C), indicating a lower concentration of target. (G) A control slide exposed to 5 nM nanoparticle probe and then exposed to silver amplification solution for 5 min. No darkening of the slide is observed. (H) Graph of 8-bit gray scale values as a function of target concentration. The gray scale values were taken from flatbed scanner images of oligonucleotide-functionalized glass surfaces that had been exposed to varying concentrations of oligonucleotide target, labeled with 5 nM oligonucleotide probe and immersed in silver amplification solution. For any given amplification time, the grayscale range is limited by surface saturation at high grayscale values and the sensitivity of the scanner at low values. Therefore, the dynamic range of this system can be adjusted by means of hybridization and amplification conditions (that is, lower target concentrations require longer amplification periods). Squares: 18-base capture-target overlap (5), 8× PBS hybridization buffer [1.2 M NaCl and 10 mM $\text{NaH}_2\text{PO}_4/\text{Na}_2\text{HPO}_4$ buffer (pH 7)], 15 min amplification time. Circles: 12-base capture-target overlap, 8× PBS hybridization buffer, 10 min amplification time. Triangles: 12-base capture-target overlap, 2× PBS hybridization buffer [0.3 M NaCl, 10 mM $\text{NaH}_2\text{PO}_4/\text{Na}_2\text{HPO}_4$ buffer (pH 7)], 5 min amplification time. The lowest target concentration that can be effectively distinguished from the background baseline is 50 fM.

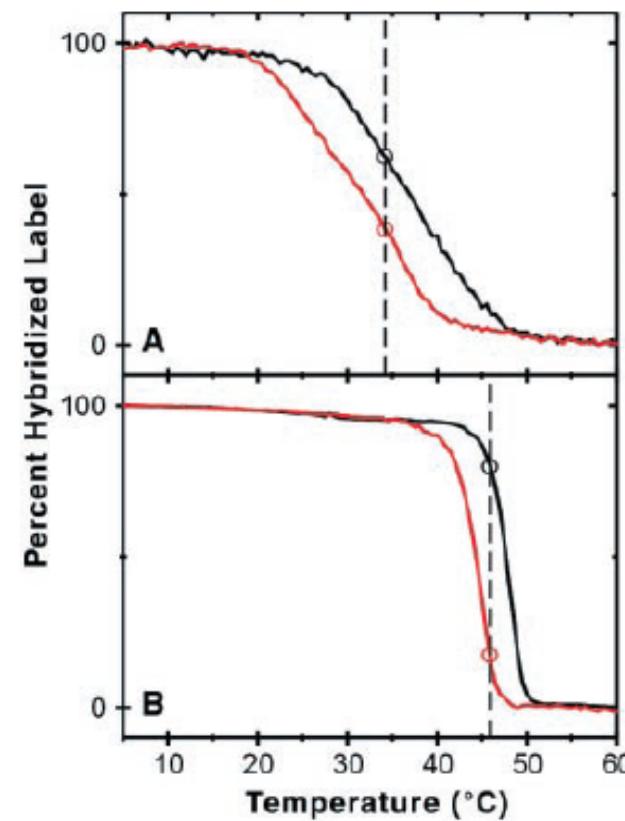
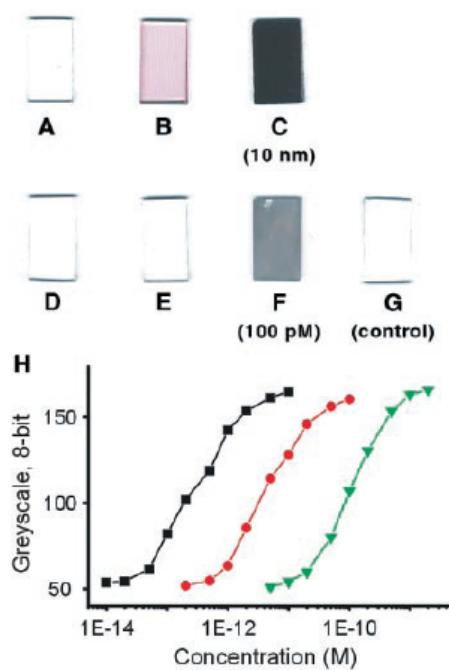
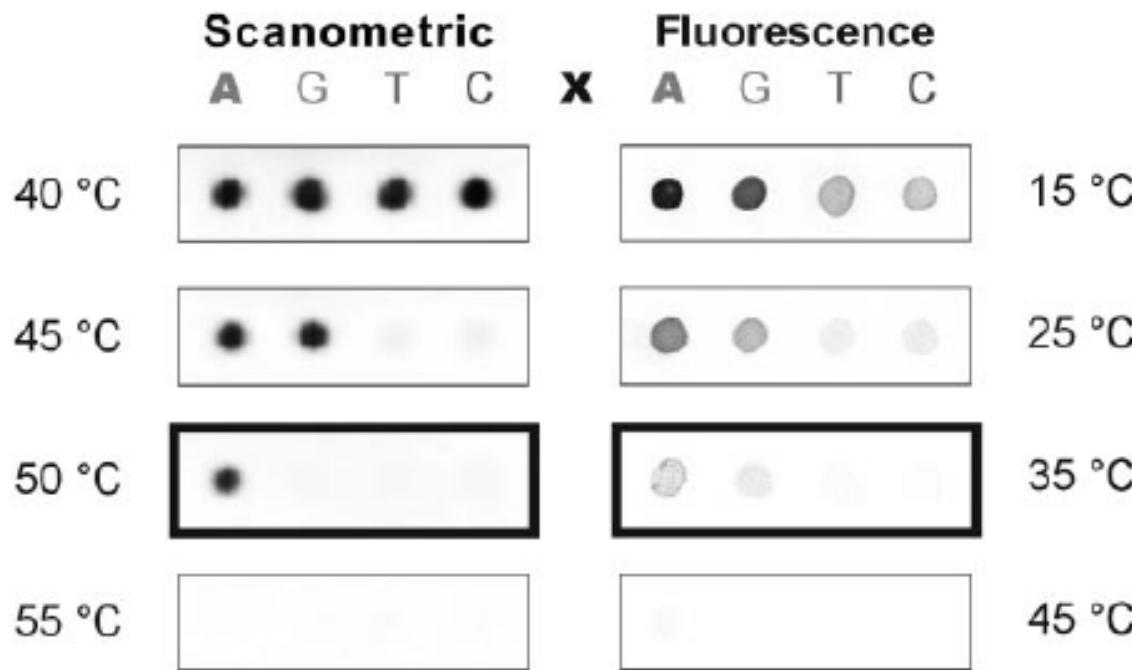


Fig. 3. (Left) Nano-particle-labeled arrays developed at different stringency temperatures. Model oligonucleotide arrays (with the capture sequences shown in Scheme 1) were treated with oligonucleotide target and nanoparticle probes, followed by a 2-min buffer wash at the temperatures shown and subsequent silver amplification (13). Images were obtained



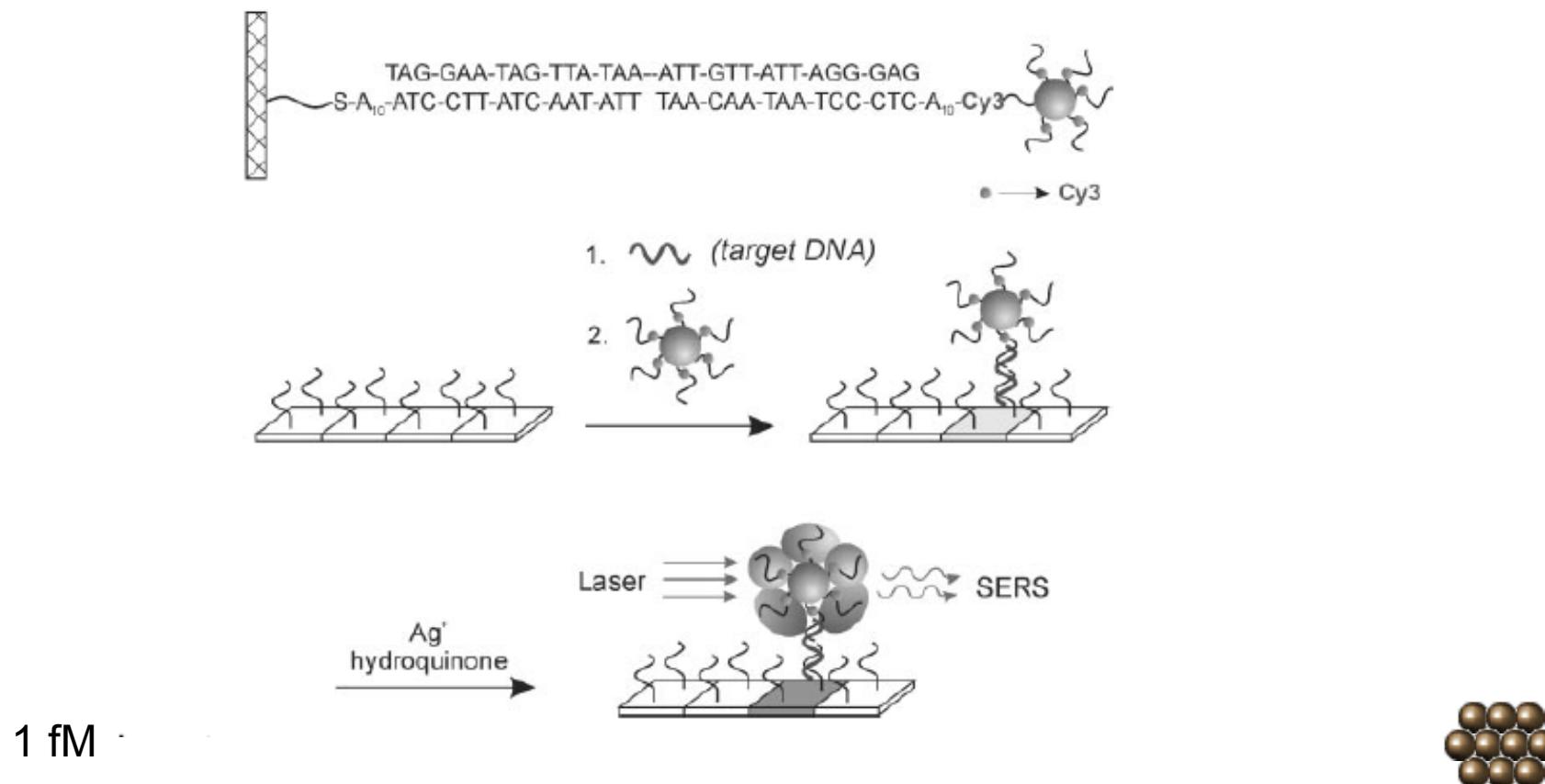
with an Epson Expression 636 (600 dots per inch) flatbed scanner (Epson America, Long Beach, California). The darkened border indicates the array that showed optimum selectivity for the perfectly complementary target; at this temperature, the ratio of background-subtracted, 8-bit gray scale values for elements A:G:T:C, obtained from histogram averages in Adobe Photoshop (Adobe Systems, San Jose, California), is 96:9:7:6. **(Right)** Fluorophore-labeled arrays washed at different stringency temperatures. Model oligonucleotide arrays identical to those shown at left were treated with oligonucleotide target and Cy3-labeled oligonucleotide probes, followed by a 2-min buffer wash at the temperatures shown. Images were obtained with a ScanArray Confocal Microarray Scanner (GSI Lumonics, Billerica, Massachusetts). The darkened border indicates the array that showed the highest selectivity for the perfectly complementary target, as calculated by the QuantArray Analysis software package (GSI Lumonics); at this temperature, the intensity ratio (in percent, with the intensity of the X = A element at 15°C set to 100%) for elements A:G:T:C is 18:7:1:1.



Nanoparticles with Raman Spectroscopic Fingerprints for DNA and RNA Detection

YunWei Charles Cao, Rongchao Jin, Chad A. Mirkin*

30 AUGUST 2002 VOL 297 SCIENCE



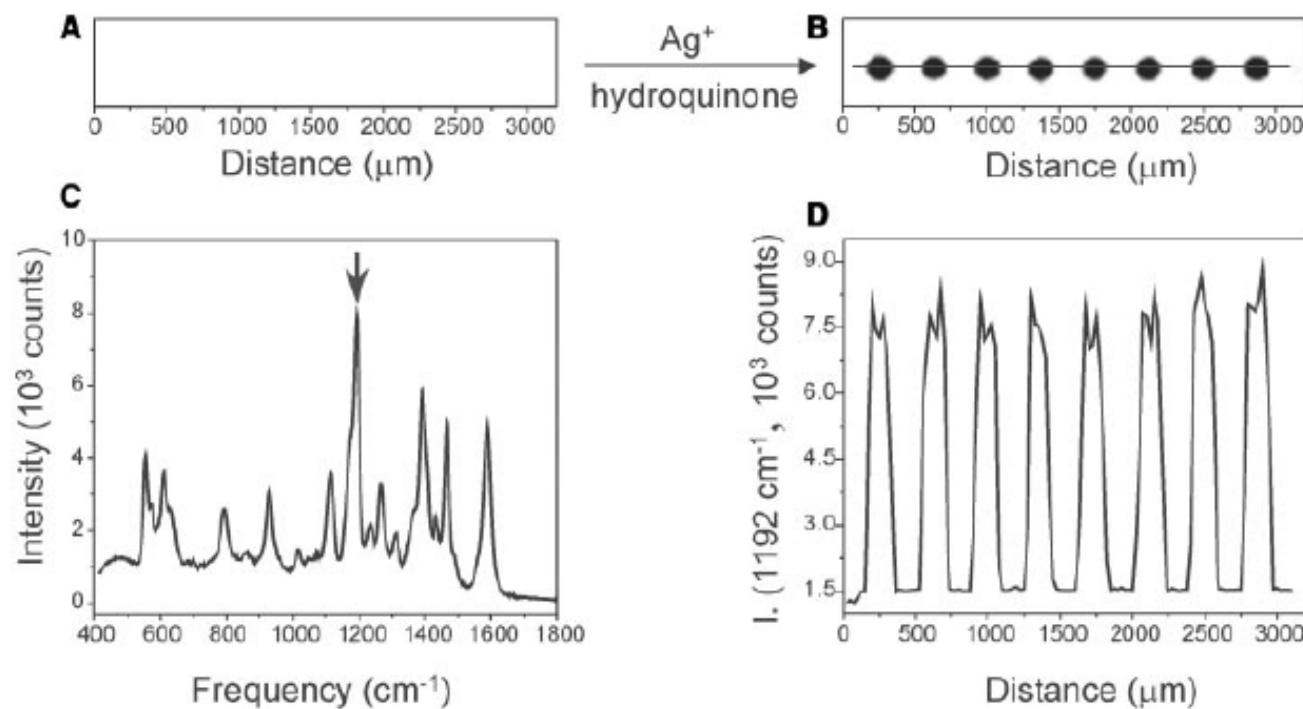


Fig. 1. Flatbed scanner images of microarrays hybridized with nanoparticles (A) before and (B) after Ag enhancing. (C) A typical Raman spectrum acquired from one of the Ag spots. (D) A profile of Raman intensity at 1192 cm^{-1} as a function of position on the chip; the laser beam from the Raman instrument is moved over the chip from left to right as defined by the line in (B).



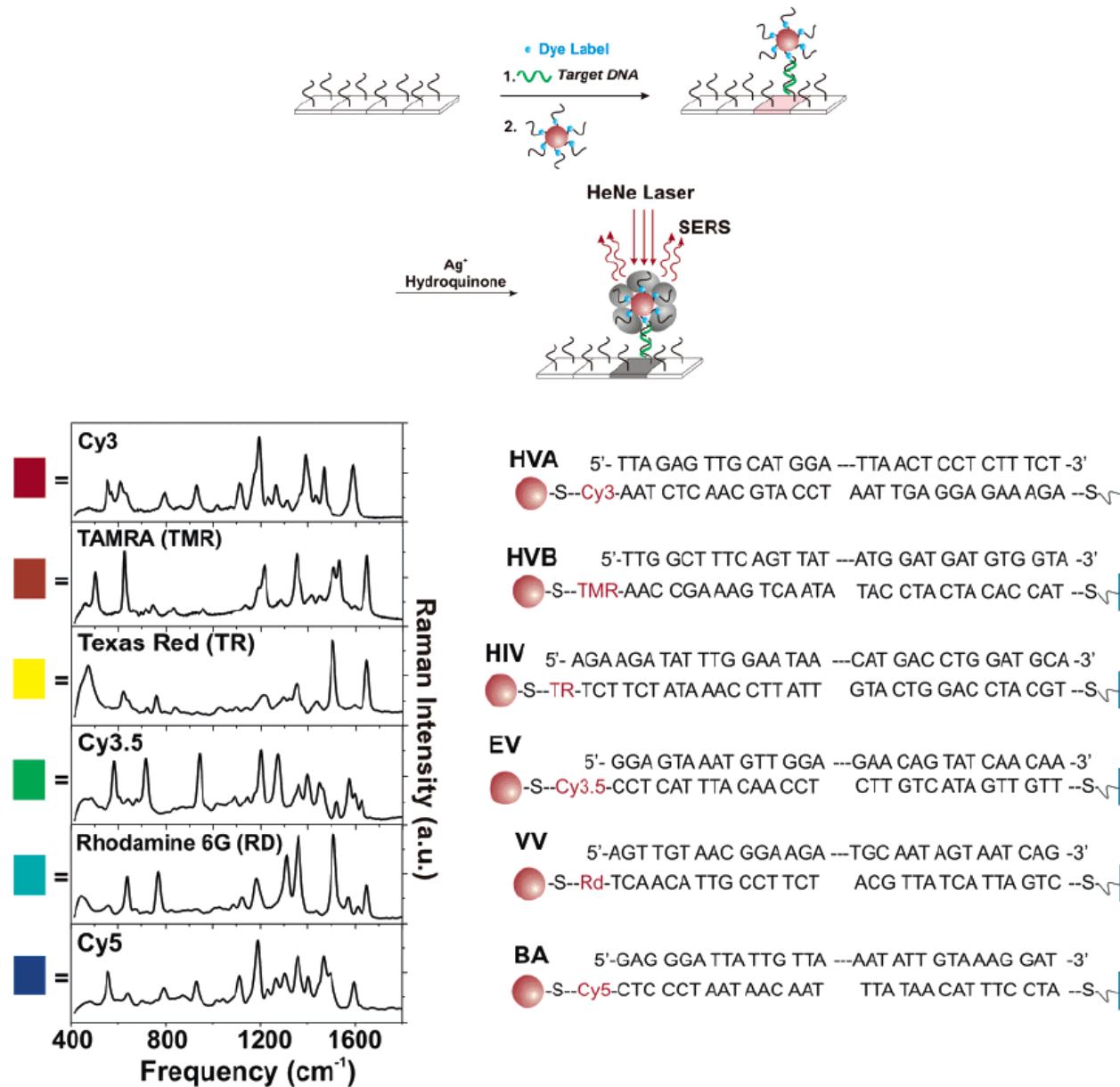


Figure 5. If Raman dyes (blue spheres) are attached to the labeling probe in the scanometric assay, the targets can be encoded and detected via the Raman signal of their labels. (Reprinted with permission from *Science* (<http://www.aaas.org>), ref 68. Copyright 2002 American Association for the Advancement of Science.)



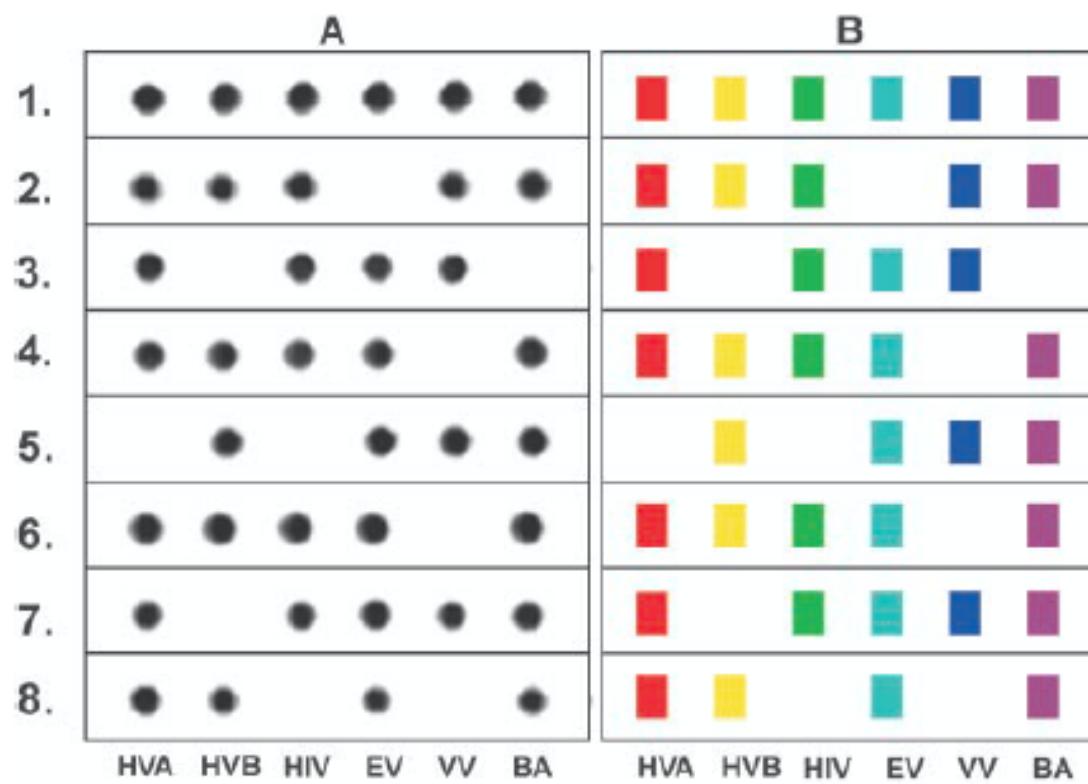


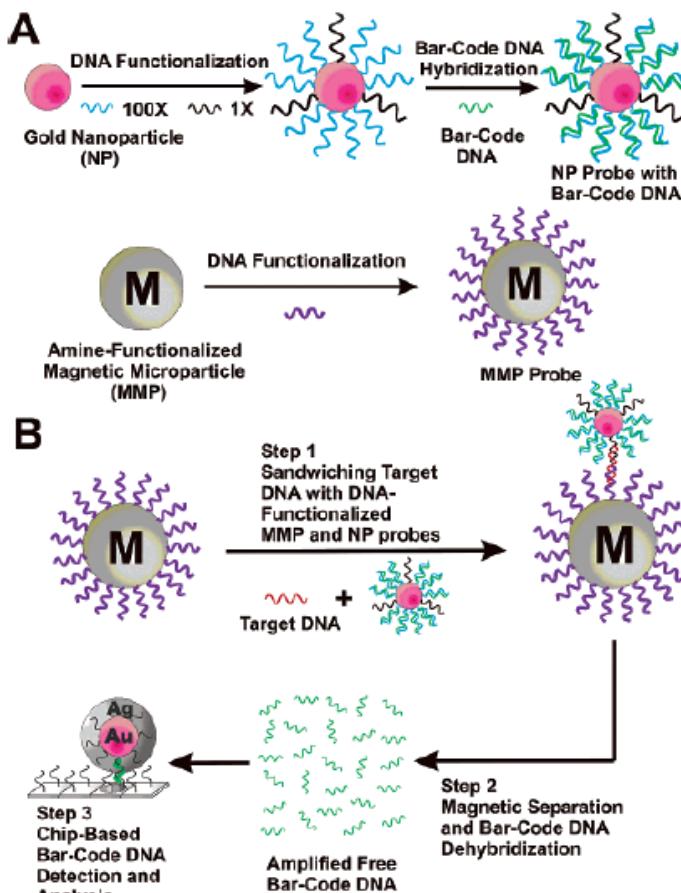
Fig. 3. (A) Flatbed scanner images of Ag-enhanced microarrays and **(B)** corresponding Raman spectra. The colored boxes correlate with the color-coded Raman spectra in Fig. 2. No false-positives or false-negatives were observed.



Bio-Bar-Code-Based DNA Detection with PCR-like Sensitivity

Jwa-Min Nam, Savka I. Stoeva, and Chad A. Mirkin*

J. AM. CHEM. SOC. 2004, 126, 5932–5933



Target DNA (anthrax): 5' GGATTATTGTTAAATATTGATAAGGAT 3'
Bar-Code DNA: 5' AGCTACGAGTTGAGAATCCTGAAATGCGACG 3'

Figure 1. The DNA-BCA assay. (A) Nanoparticle and magnetic microparticle probe preparation. (B) Nanoparticle-based PCR-less DNA amplification scheme.



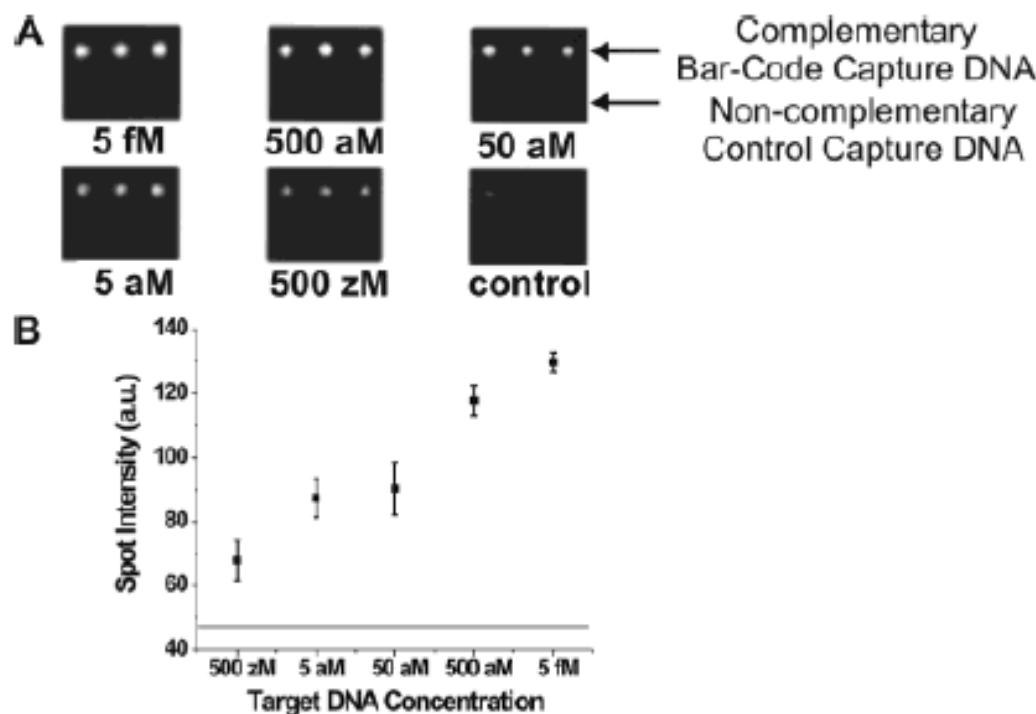


Figure 2. Amplified anthrax bar-code DNA detection with the Verigene ID system. (A) Anthrax bar-code DNA detection with 30 nm NP probes. (B) Quantitative data of spot intensities with 30 nm NP probes (Adobe Photoshop, Adobe Systems, Inc., San Jose, CA). The horizontal line represents control signal intensity (47 ± 2).

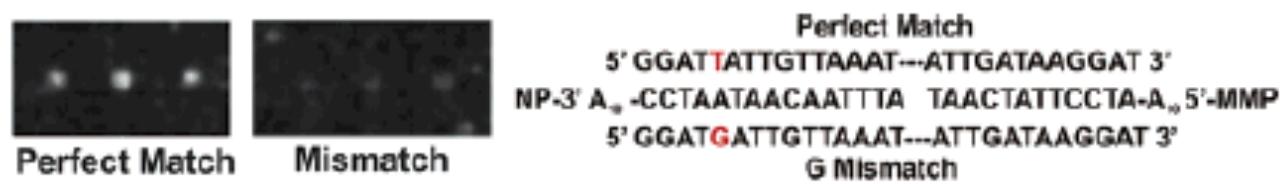


Figure 3. Single base mismatch experiment.



Nanoparticle-Based Bio-Bar Codes for the Ultrasensitive Detection of Proteins

26 SEPTEMBER 2003 VOL 301 SCIENCE

Jwa-Min Nam,* C. Shad Thaxton,* Chad A. Mirkin†

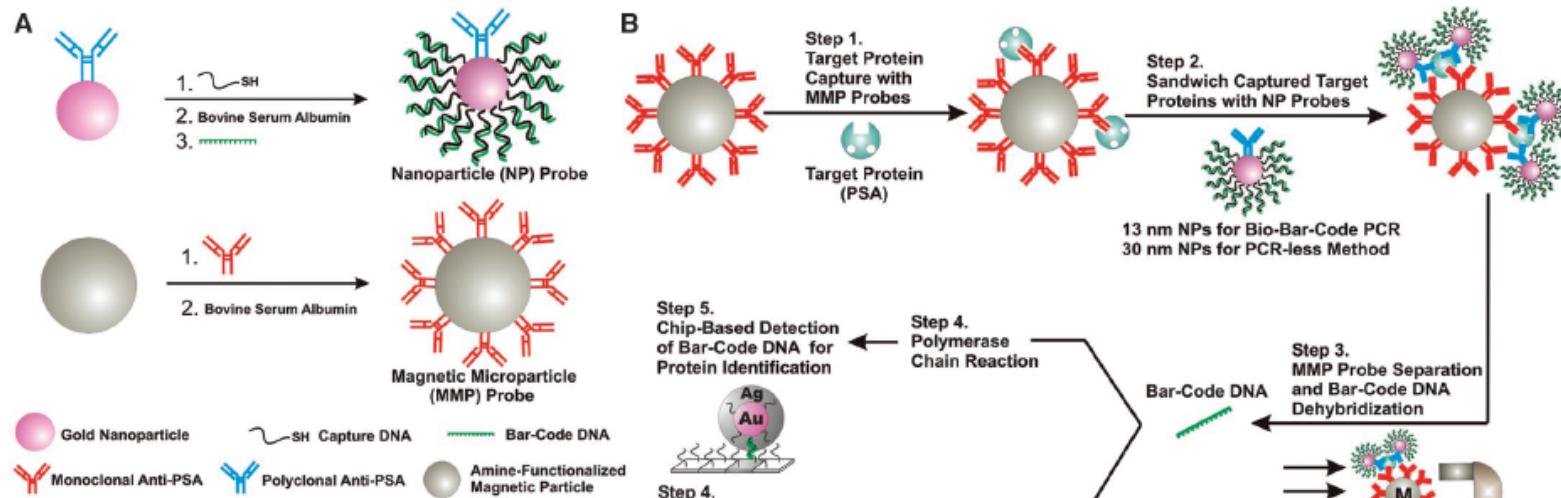


Fig. 1. The bio-bar-code assay method. (A) Probe design and preparation. (B) PSA detection and bar-code DNA amplification and identification. In a typical PSA-detection experiment, an aqueous dispersion of MMP probes functionalized with mAbs to PSA (50 μ l of 3 mg/ml magnetic probe solution) was mixed with an aqueous solution of free PSA (10 μ l of PSA) and stirred at 37°C for 30 min (Step 1). A 1.5-ml tube containing the assay solution was placed in a BioMag microcentrifuge tube separator (Polysciences, Incorporated, Warrington, PA) at room temperature. After 15 s, the MMP-PSA hybrids were concentrated on the wall of the tube. The supernatant (solution of unbound PSA molecules) was removed, and the MMPs were resuspended in 50 μ l of 0.1 M phosphate-buffered saline (PBS) (repeated twice). The NP probes (for 13-nm NP probes, 50 μ l at 1 nM; for 30-nm NP probes, 50 μ l at 200 pM), functionalized with polyclonal Abs to PSA and hybridized bar-code DNA strands, were then added to the assay solution. The NPs reacted with the PSA immobilized on the MMPs and provided DNA strands for signal amplification and protein identification (Step 2). This solution was vigorously stirred at 37°C for 30 min. The MMPs were then washed with 0.1 M PBS with the magnetic separator to isolate the mag-

netic particles. This step was repeated four times, each time for 1 min, to remove everything but the MMPs (along with the PSA-bound NP probes). After the final wash step, the MMP probes were resuspended in NANOpure water (50 μ l) for 2 min to dehybridize bar-code DNA strands from the nanoparticle probe surface. Dehybridized bar-code DNA was then easily separated and collected from the probes with the use of the magnetic separator (Step 3). For bar-code DNA amplification (Step 4), isolated bar-code DNA was added to a PCR reaction mixture (20- μ l final volume) containing the appropriate primers, and the solution was then thermally cycled (20). The bar-code DNA amplicon was stained with ethidium bromide and mixed with gel-loading dye (20). Gel electrophoresis or scanometric DNA detection (24) was then performed to determine whether amplification had taken place. Primer amplification was ruled out with appropriate control experiments (20). Notice that the number of bound NP probes for each PSA is unknown and will depend upon target protein concentration.



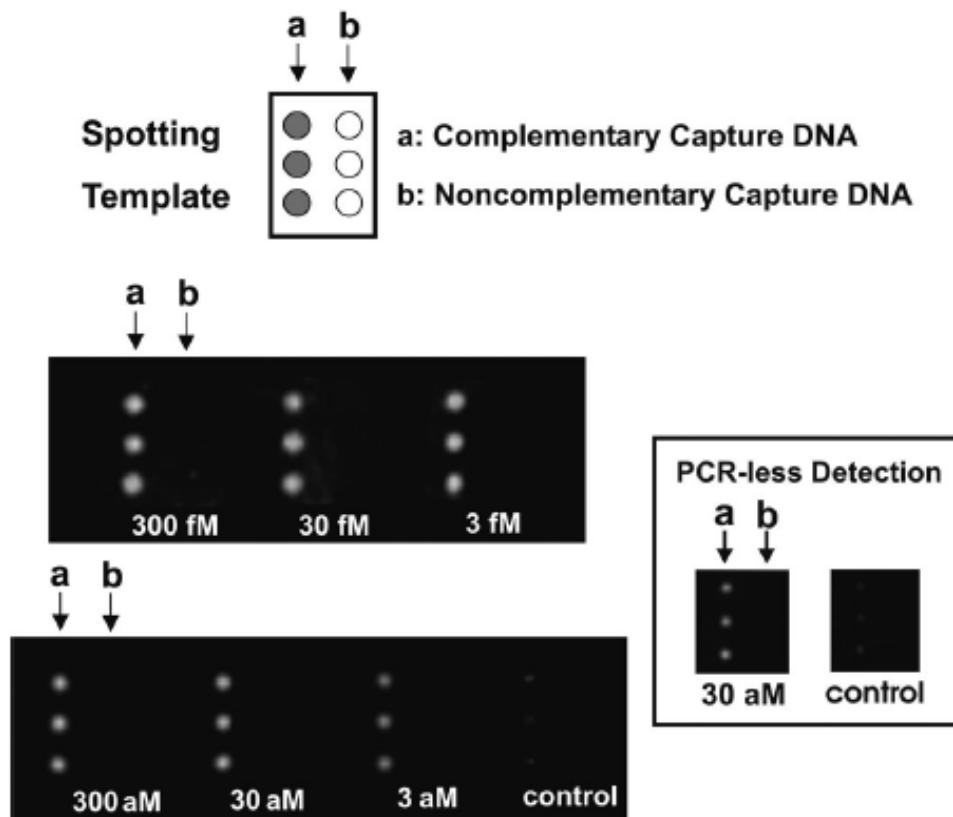


Fig. 2. Scanometric detection of PSA-specific bar-code DNA. PSA concentration (sample volume of 10 μ l) was varied from 300 fM to 3 aM and a negative control sample where no PSA was added (control) is shown. For all seven samples, 2 μ l of anti-dinitrophenyl (10 pM) and 2 μ l of β -galactosidase (10 pM) were added as background proteins. Also shown is PCR-less detection of PSA (30 aM and control) with 30 nm NP probes (inset). Chips were imaged with the Verigene ID system (20).



Table 1. Detection Limits of Nucleic Acid Assays^a

	assay	ss DNA	PCR products	genomic DNA
nanostructure-based methods	colorimetric ²⁹ (cross-linked Au nanoparticles)	~10 nM		
	colorimetric ³⁶ (non-cross-linked Au nanoparticles)	60 nM		
	magnetic relaxation ⁹⁷ (iron oxide nanoparticles)	20 pM		
	electrochemical ⁹⁶ (nanoparticles)	270 pM		
	scanometric ^{35,66,67} (Au nanoparticles with Ag amplification)	50 fM	100 aM ^b	200 fM
	Raman spectroscopy ⁶⁸ (Au nanoparticles with Ag amplification)	~1 fM		
	electrical ⁹³ (Au nanoparticles with Ag amplification)	500 fM		
	electrical ⁹⁹ (Si nanowire)	10 fM		
	electrical ¹⁰³ (carbon nanotube)	54 aM		
	resonant light-scattering ⁶¹⁻⁶⁶ (metal nanoparticles)	170 fM ^b		33 fM
	fluorescence ⁵⁶ (ZnS and CdSe quantum dots)	2 nM		
	surface plasmon resonance ⁴¹ (Au nanoparticles)	10 pM		
	quartz crystal microbalance ⁹⁴ (Au nanoparticles)	~1 fM		
other non-enzymatic based methods	laser diffraction ⁴² (Au nanoparticles)	~50 fM		
	fluorescence ⁴⁵ (fluorescent nanoparticles)	~1 fM		
	bio-bar-code amplification ⁷¹ (Au nanoparticles with Ag amplification)	500 zM		
	fluorescence ³⁵ (molecular fluorophores)		~600 fM ^b	
	fluorescence (dendrimer amplification) ¹³⁴		2.5 μ g	
	electrochemical amplification ¹³⁶ (electroactive reporter molecules)	100 aM		

^a Detection limits can vary based on target length and sequence; therefore, it is difficult to compare assays without testing them using identical targets and conditions. ^b Values taken from ref 34.



Table 2. Detection Limits of Protein Assays

	assay	target	protein in saline	protein in serum
nanostructure-based methods	optical ⁷² (Au nanoshells)	rabbit IgG	0.88 ng/mL (~4.4 pM) ^a	0.88 ng/mL (~4.4 pM) ^a
	optical ⁷⁴ (Au nanoparticles)	IgE and IgG1	~20 nM	
	magnetic relaxation ⁹⁸ (iron oxide nanoparticles)	adenovirus (ADV) and herpes simplex virus (HSV)	100 ADV/ 100 μL	50 HSV/ 100 μL
	scanometric ⁷⁹ (Au nanoparticles with Ag amplification)	mouse IgG	200 pM	
	Raman ⁸² (Au nanoparticles with Raman labels)	prostate-specific antigen		30 fM
	surface plasmon resonance ^{83,84} (triangular Ag particles on surfaces)	streptavidin(S A) and anti-biotin (AB)	~1 pM SA and ~700 pM AB	
	electrical ¹¹⁰ (single-walled carbon nanotubes)	10E3 antibody to U1A RNA splicing factor	~1 nM	
	electrical ²⁰ (Si nanowires)	streptavidin	10 pM	
molecular fluorophore methods	bio-bar-code amplification ⁷⁵ (Au nanoparticles with Ag amplification)	prostate-specific antigen	30 aM (3 aM) ^b	(30 aM) ^b
	enzyme-linked immunosorbent assay	various	pM range	pM range
electrochemical methods	electrochemical amplification ¹³⁷ (oligonucleotide reporter molecules)	IgG	13 fM	
enzyme-based amplification methods	immuno-PCR ⁷⁶	bovine serum albumin	2 fM	
	rolling circle amplification ⁷⁷	prostate-specific antigen	3 fM	

^a Reported in ng/mL; authors converted to molar concentration for ease of comparison. ^b These values are the lower limits when PCR is used to amplify the bar-code DNA prior to scanometric detection of bar codes.



Surface Plasmon

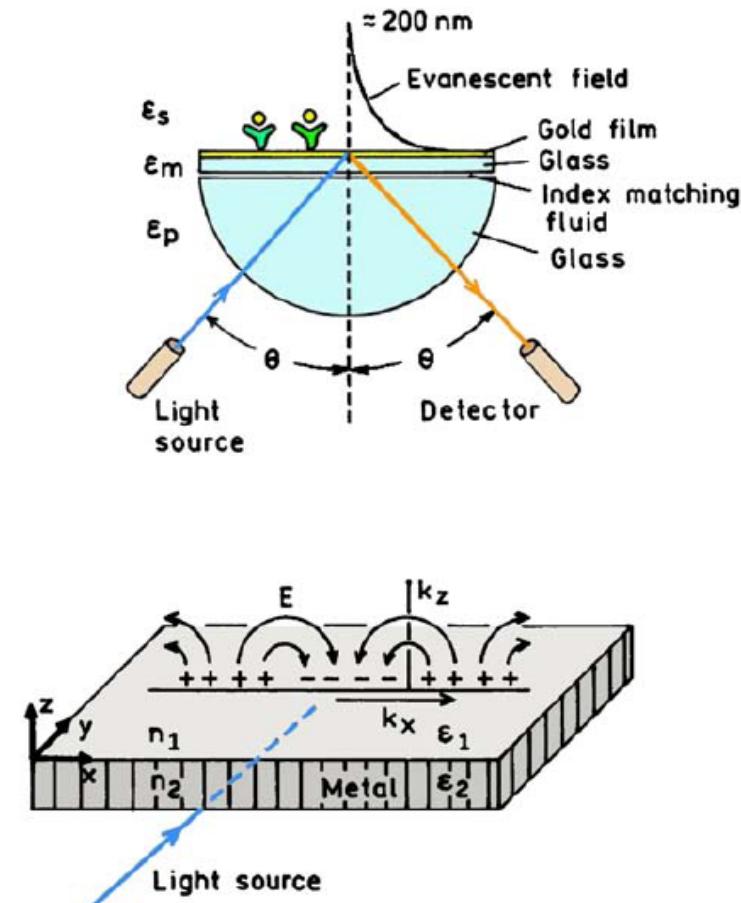
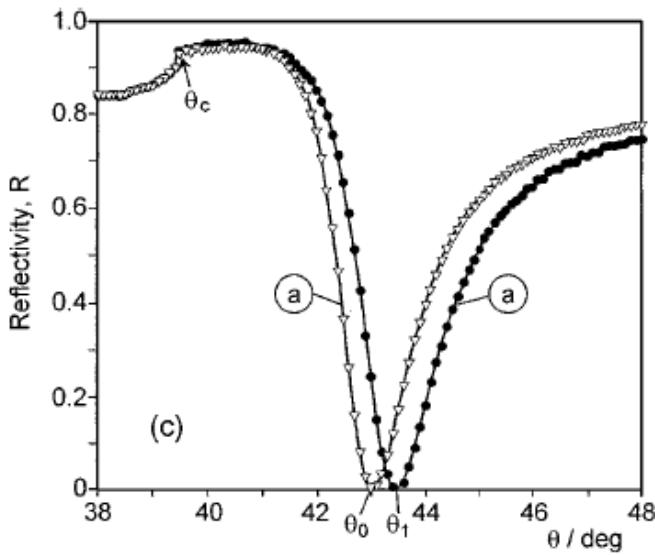
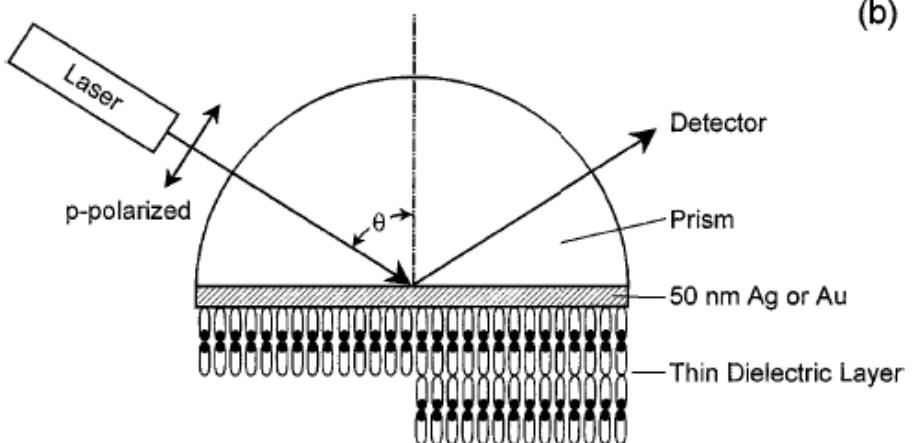
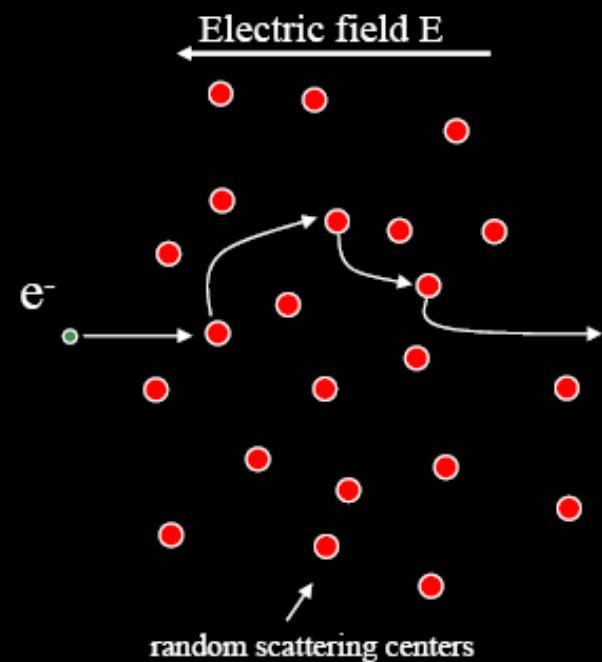


Figure 3. Schematics of an SPR experiment (top) and of the light-induced surface plasmons (bottom).



Drift: Drude model



$$F = ma$$

$$eE = m \frac{\partial v}{\partial t}$$

$$v_{avg} = \underbrace{\frac{e\tau}{m}}_{\mu} E$$

$$j = ne v_{avg} = \underbrace{\frac{ne^2\tau}{m}}_{\sigma} E$$

$$m \frac{\partial}{\partial t} \langle \vec{v} \rangle = q \vec{E} - \gamma \langle \vec{v} \rangle$$

$$\sigma(\omega) = \frac{\sigma_0}{1 + i\omega\tau}$$



AC Dielectric Response

$$\epsilon_m = 1 - \frac{\omega_p^2}{\omega^2} \quad \text{Plasma frequency}$$

polarizability of a small metal sphere with dielectric function $\epsilon(\lambda)$

$$\alpha = R^3 \frac{\epsilon - 1}{\epsilon + 2}.$$

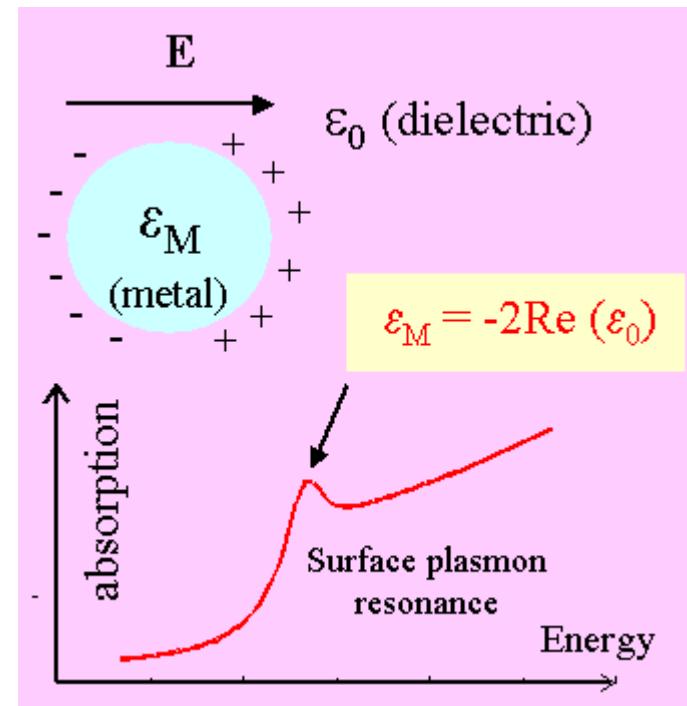
$$\epsilon = \epsilon_b + 1 - \frac{\omega_p^2}{\omega^2 + i\omega\gamma},$$

$$\alpha = \frac{R^3(\epsilon_b\omega^2 - \omega_p^2) + i\omega\gamma\epsilon_b}{[(\epsilon_b + 3)\omega^2 - \omega_p^2] + i\omega\gamma(\epsilon_b + 3)}.$$

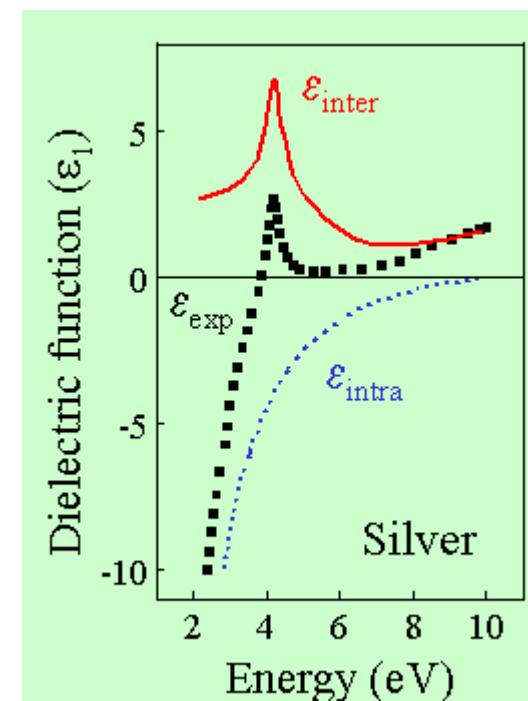
$$\omega_R = \frac{\omega_p}{\sqrt{\epsilon_b + 3}}$$

$$\gamma(\epsilon_b + 3)$$



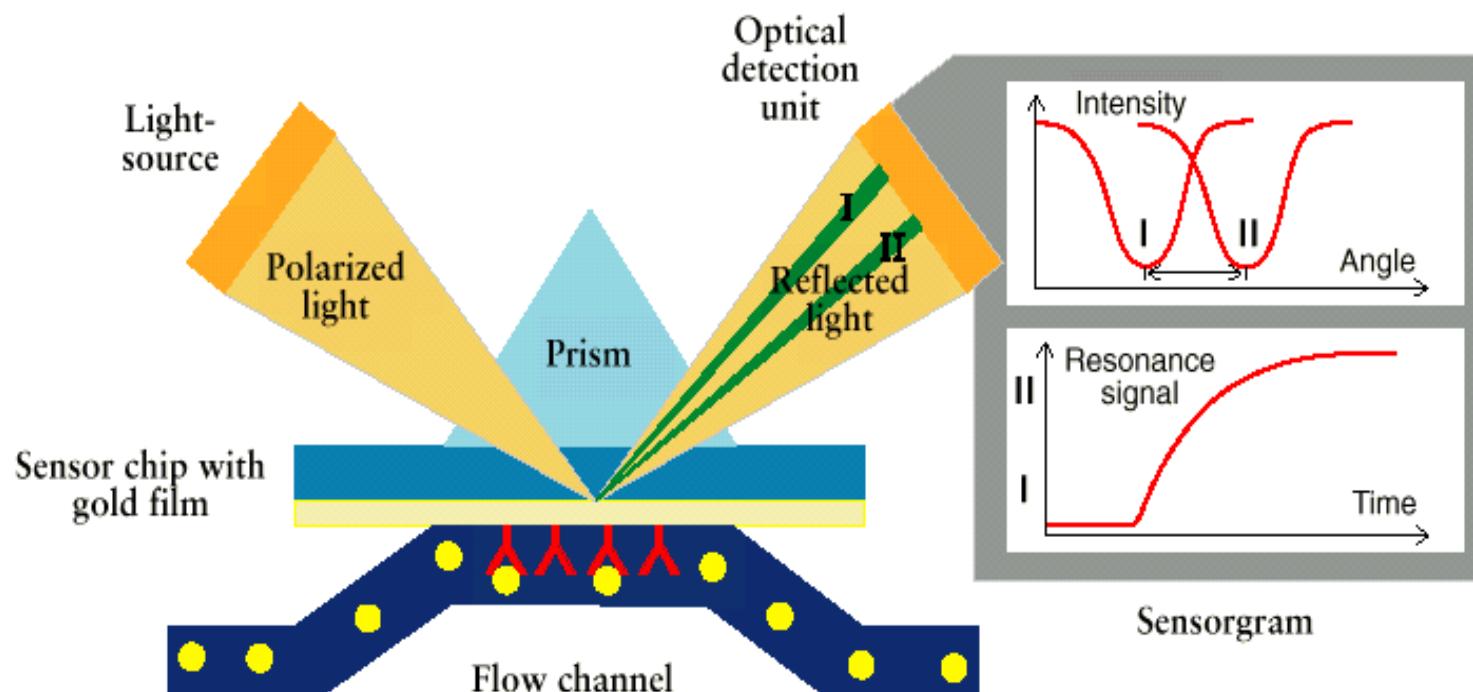


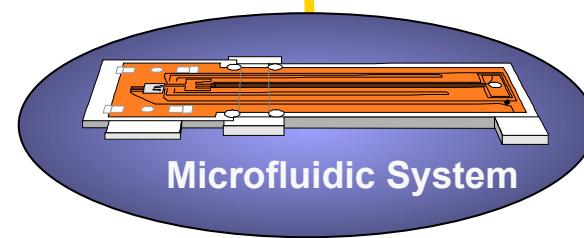
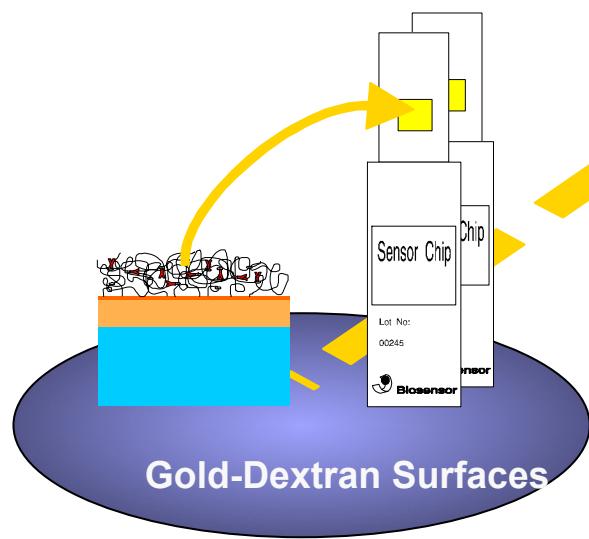
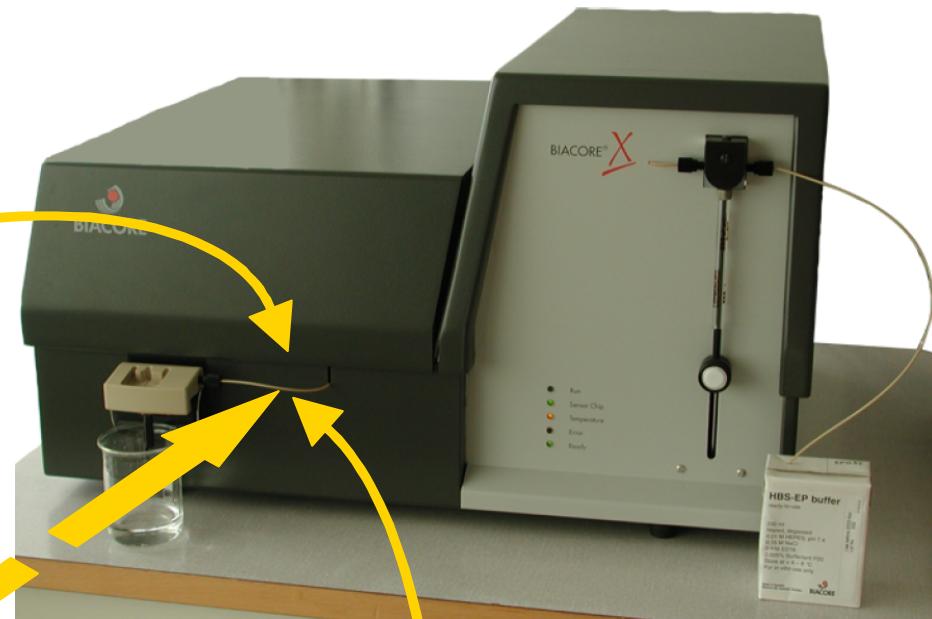
$$\epsilon_{eff} = \epsilon_0 + 3N\epsilon_0 \frac{\epsilon_M - \epsilon_0}{\epsilon_M + 2\epsilon_0}$$



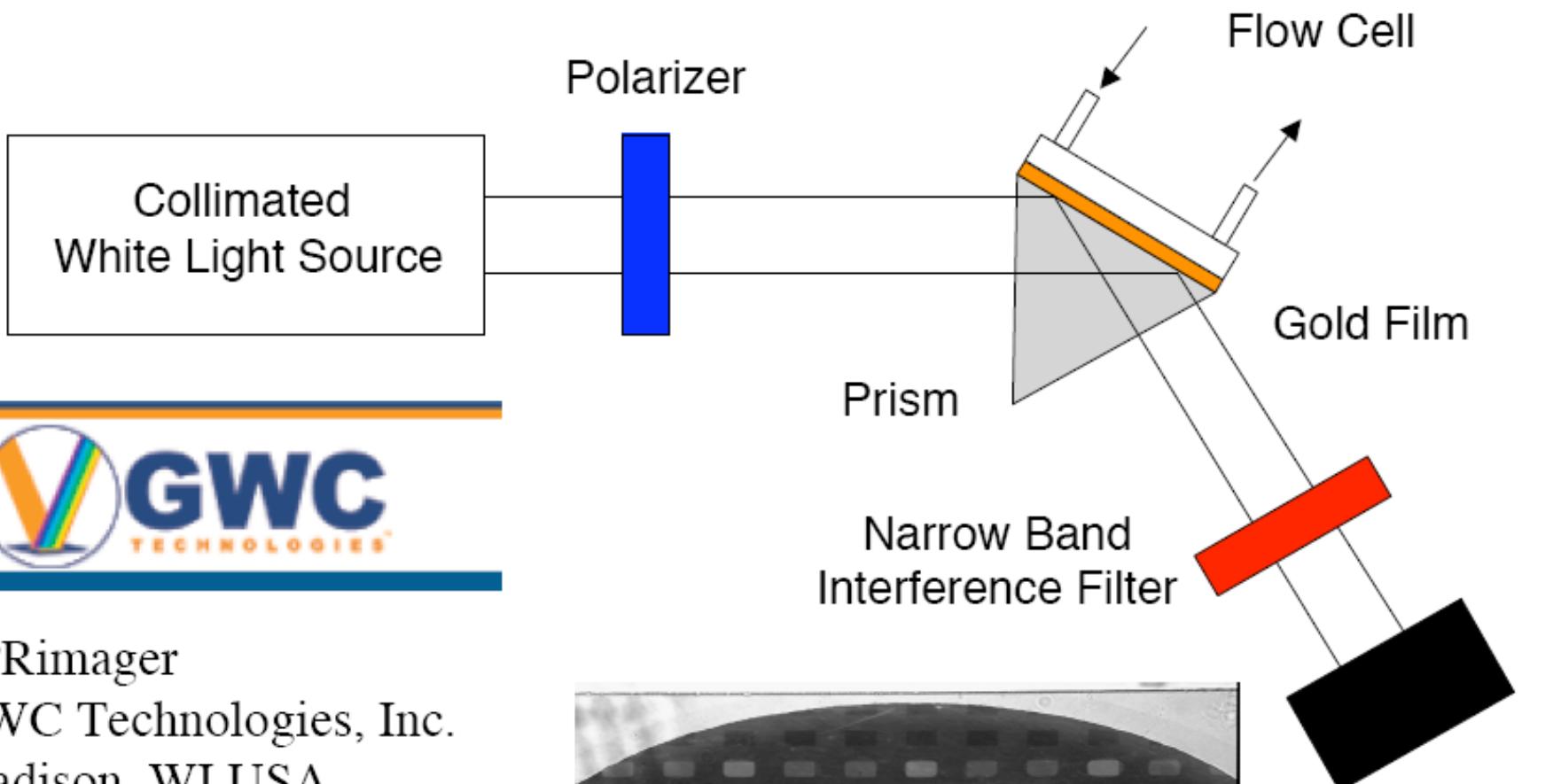
Biomolecular Binding in Real Time

Principle of Detection - SPR (Surface Plasmon Resonance)



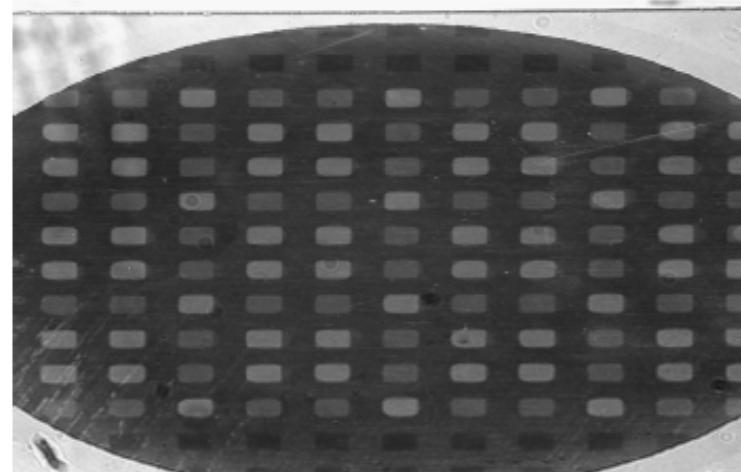


SPR Imaging Apparatus



SPRimager
GWC Technologies, Inc.
Madison, WI USA
www.gwctechnologies.com

Raw Image



UCIrvine
University of California, Irvine

Localized Plasmon

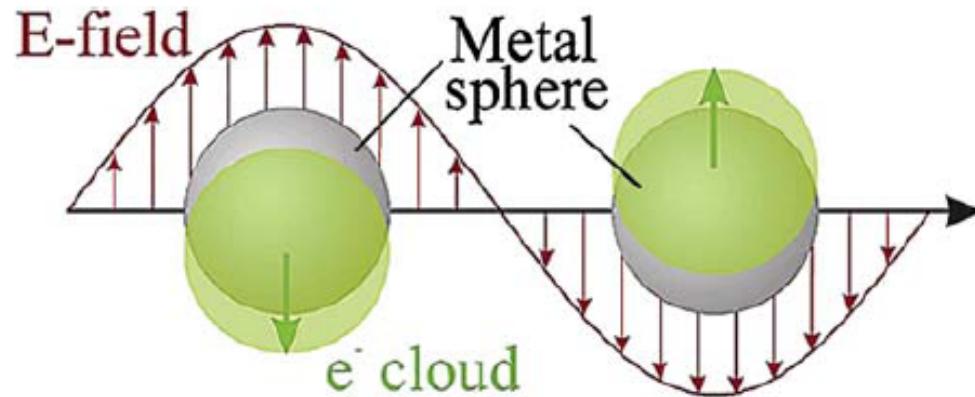
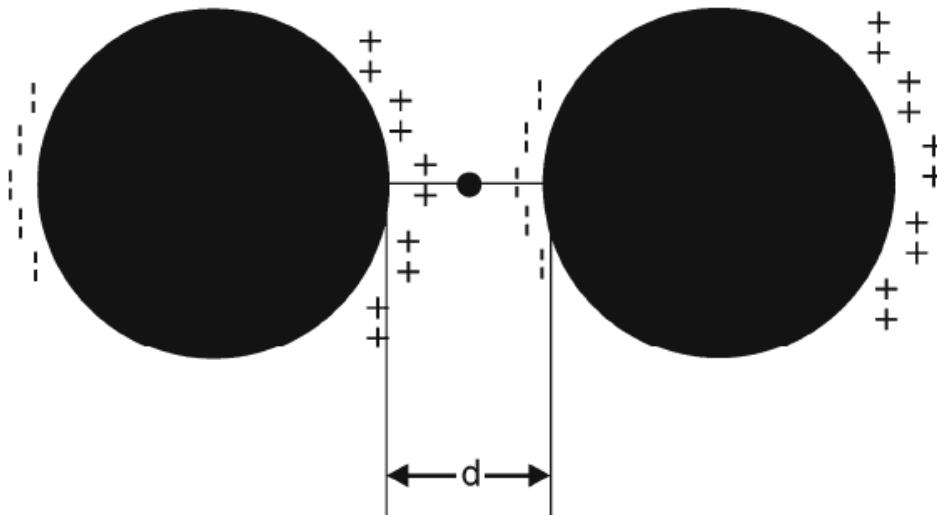
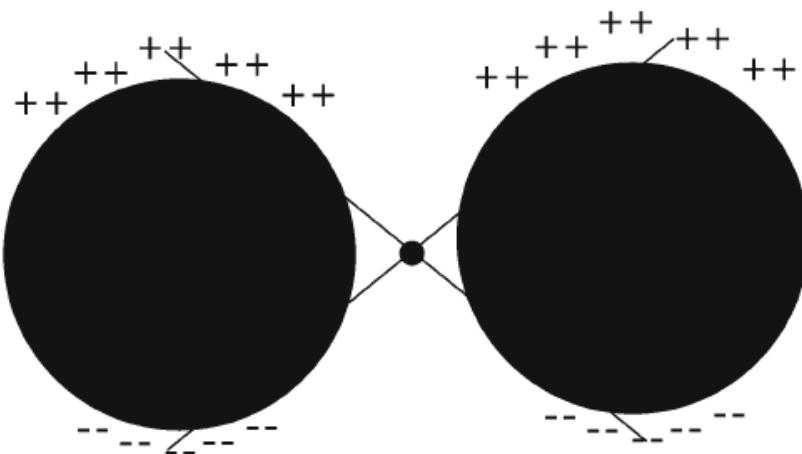


Figure 6. Schematic of plasmon oscillation for a sphere. From [39].





field enhancement

$E_s = gE_0$, where E_0 is the magnitude of the incident field

$$E_R \propto \alpha_R E_s \propto \alpha_R g E_0$$

$$E_{SERS} \propto \alpha_R g g' E_0$$

$$I_{SERS} \propto |\alpha_R|^2 |g g'|^2 I_0$$

$$g \cong g'$$

$$|E_L|^4 = |g|^4$$



Probing Single Molecules and Single Nanoparticles by Surface-Enhanced Raman Scattering

SCIENCE • VOL. 275 • 21 FEBRUARY 1997

Shuming Nie* and Steven R. Emory

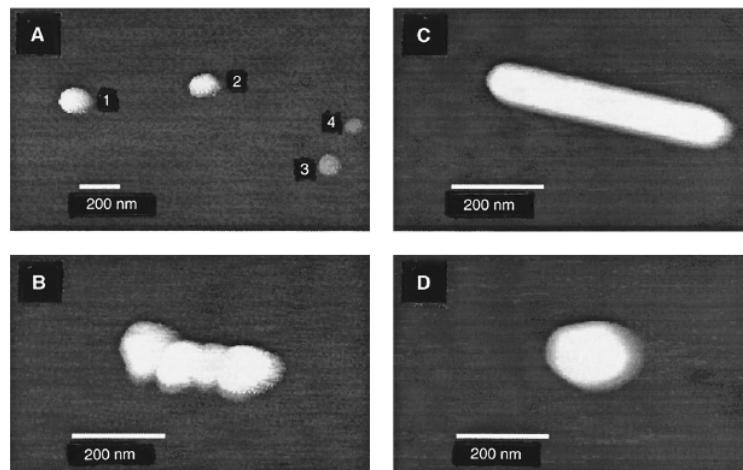
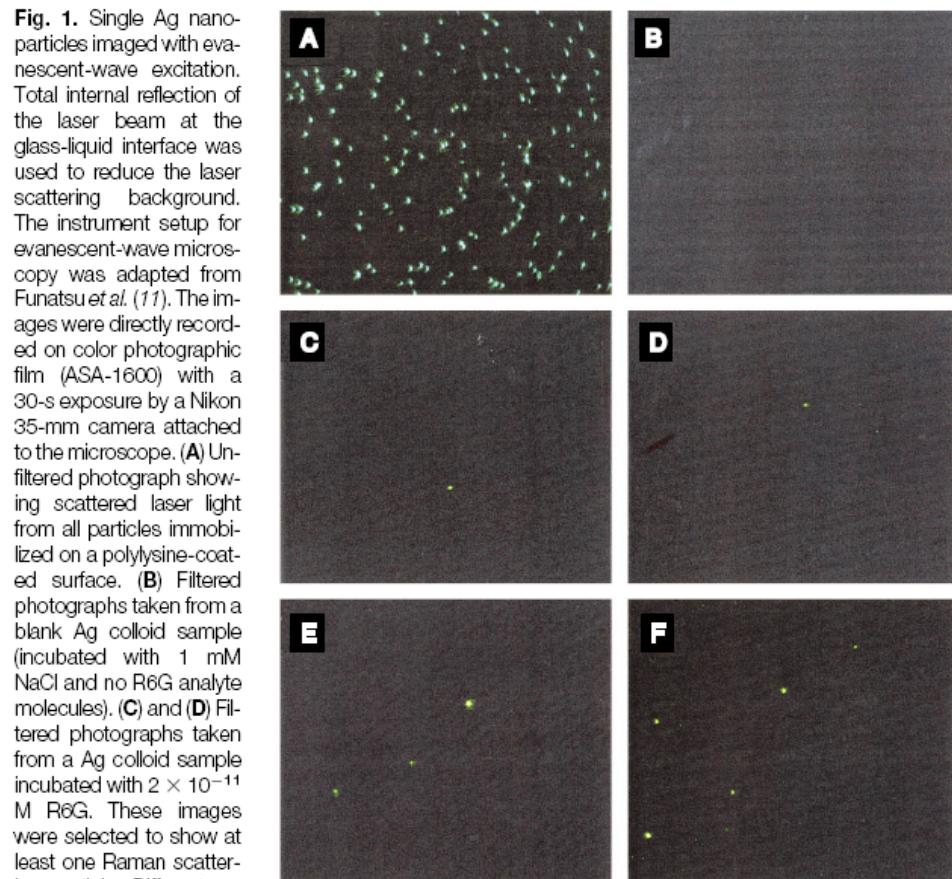
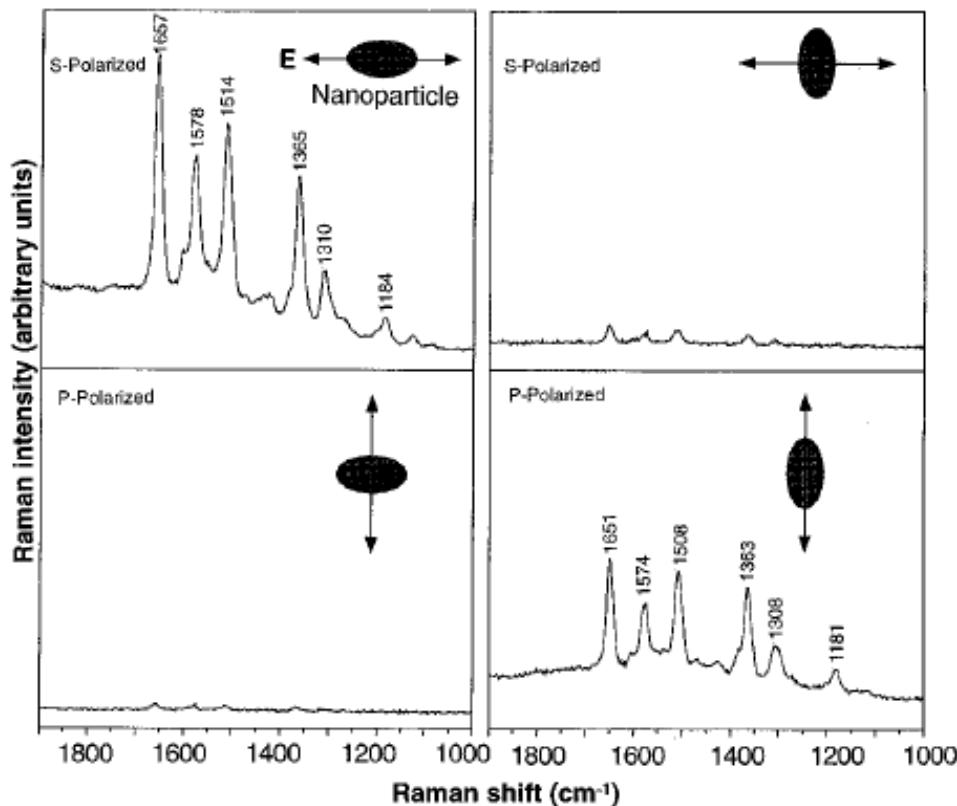


Fig. 3. Surface-enhanced Raman spectra of R6G obtained with a linearly polarized confocal laser beam from two Ag nanoparticles. The R6G concentration was 2×10^{-11} M, corresponding to an average of 0.1 analyte molecule per particle. The direction of laser polarization and the expected particle orientation are shown schematically for each spectrum. Laser wavelength, 514.5 nm; laser power, 250 nW; laser focal radius, ~ 250 nm; integration time, 30 s. All spectra were plotted on the same intensity scale in arbitrary units of the CCD detector readout signal.



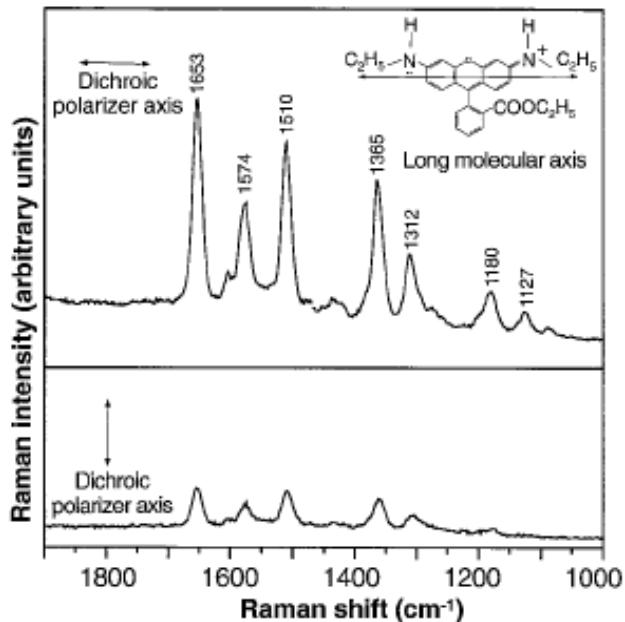


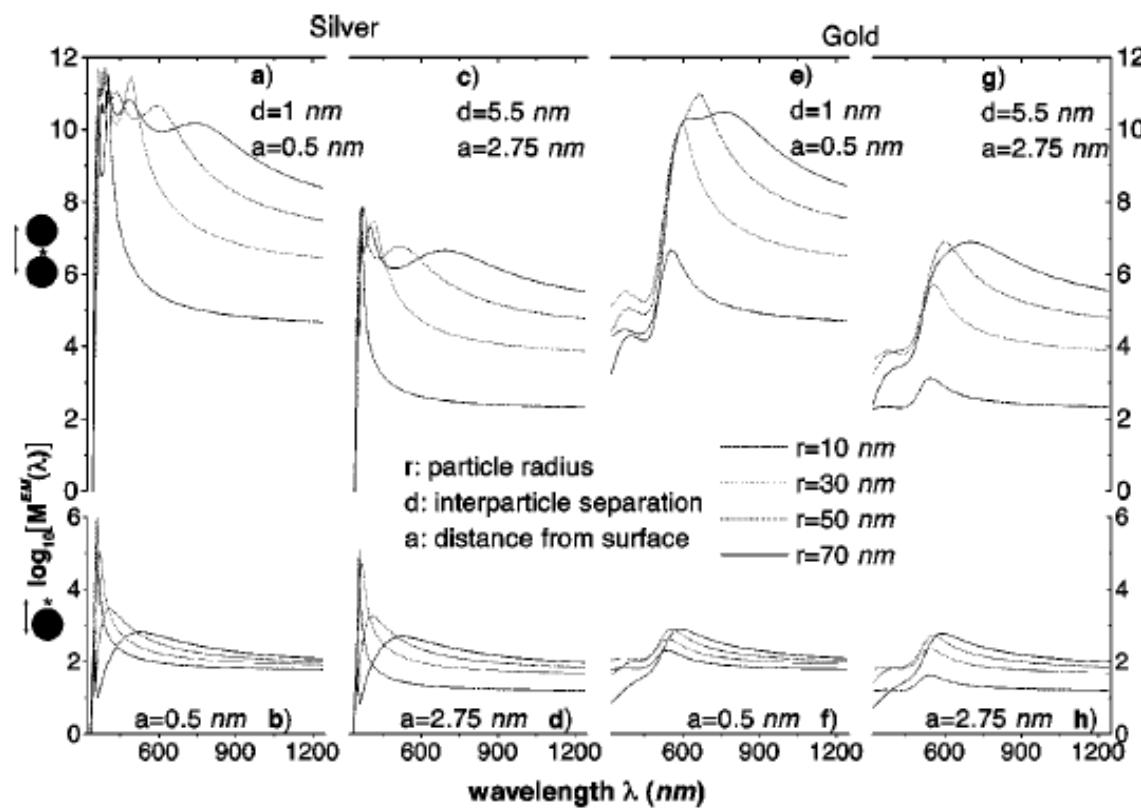
Fig. 4. Emission-polarized surface-enhanced Raman signals of R6G observed from a single Ag nanoparticle with a polarization-scrambled confocal laser beam. A dichroic sheet polarizer was rotated 90° to select Raman scattering signals polarized parallel (upper spectrum) or perpendicular (lower spectrum) to the long molecular axis of R6G. (**Inset**) Structure of R6G, the electronic transition dipole (along the long axis when excited at 514.5 nm), and the dichroic polarizer orientations. Other conditions as in Fig. 3.

troscopic signatures of adsorbed molecules. For single rhodamine 6G molecules adsorbed on the selected nanoparticles, the intrinsic Raman enhancement factors were on the order of 10^{14} to 10^{15} , much larger than the ensemble-averaged values derived from conventional measurements. This enormous enhancement leads to vibrational Raman signals that are more intense and more stable than single-molecule fluorescence.



Electromagnetic contributions to single-molecule sensitivity in surface-enhanced Raman scattering

PRE 62 4318



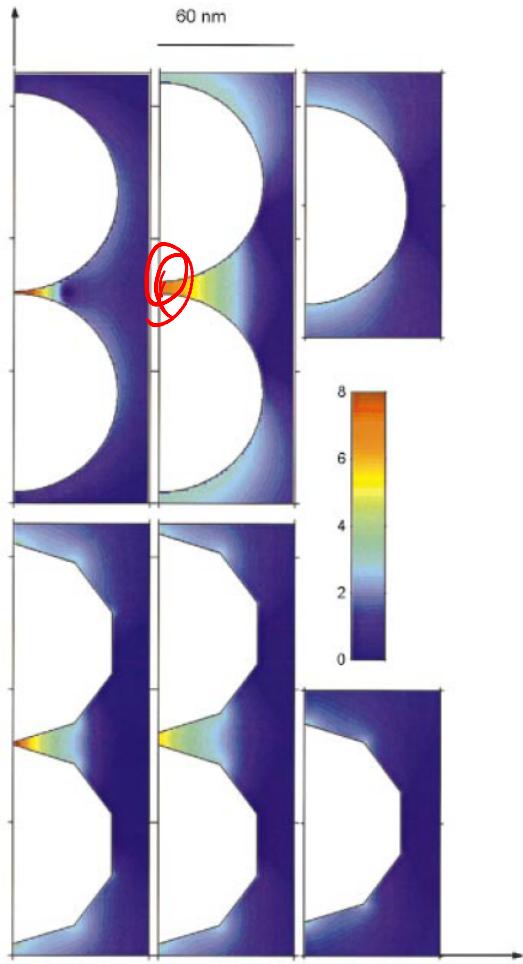


FIG. 3. (Color) EM-enhancement factor M^{EM} at a cross section through six different silver particle configurations. The wavelength of the incident field is $\lambda = 514.5$ nm with vertical polarization. The left-hand column illustrates the EM enhancement for dimer configurations of two spheres (top) and two polygons (bottom) with a separation of 1 nm. The middle column shows the same situation, but with a separation distance of 5.5 nm. The right-hand column shows the case of an isolated single particle. All particles share a common largest dimension of 90 nm. Note that the color scale from dark blue to dark red is logarithmic, covering the interval $10^0 < M^{EM} < 10^8$. Regions with enhancement outside this interval are shown in dark blue and dark red, respectively.

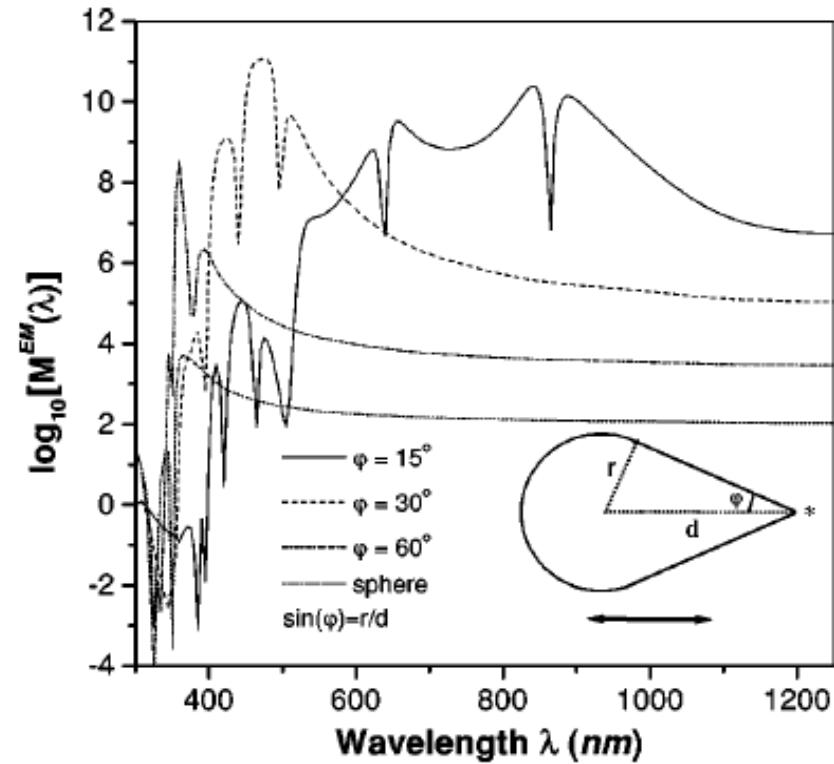


FIG. 5. EM-enhancement factor for a rotationally symmetric silver droplet as a function of the angle defining the opening edge ϕ . The field is polarized parallel to the axis of the droplet and the evaluation position (star) is located 0.5 nm outside the tip. As the droplet becomes sharper the enhancement increases several orders of magnitude.



Nanosphere Arrays with Controlled Sub-10-nm Gaps as Surface-Enhanced Raman Spectroscopy Substrates

J. AM. CHEM. SOC. 2005, 127, 14992–14993

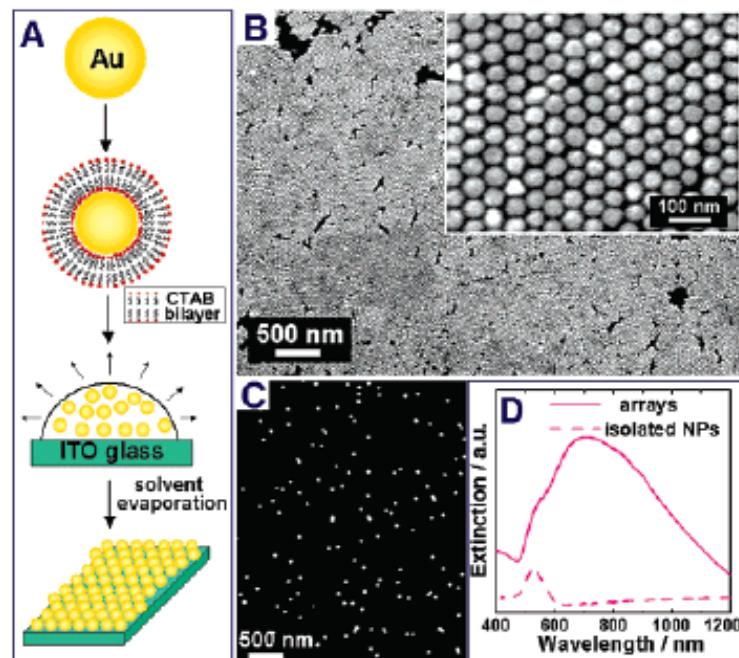


Figure 1. (A) Schematic illustration of the fabrication of sub-10-nm gap Au NP arrays. (B) SEM image of the arrays. (C) SEM image of monolayer of isolated Au NPs on ITO glass. (D) Vis–NIR extinction spectrum of the monolayer of isolated Au NPs and arrays.

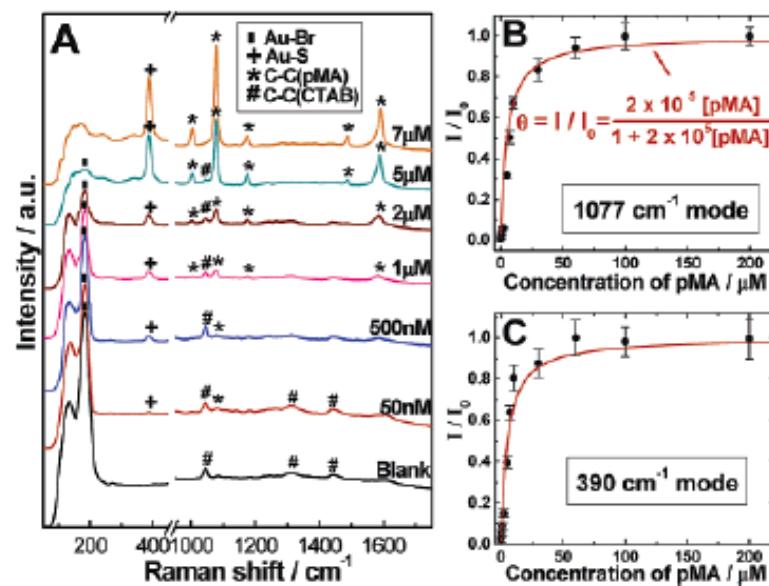
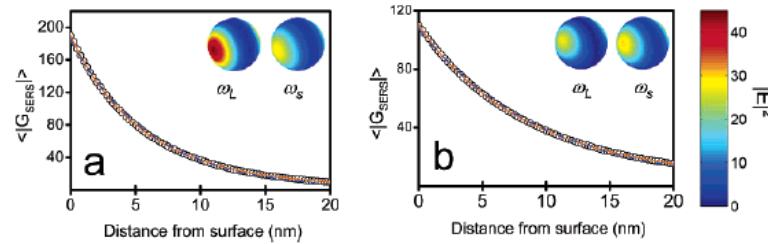
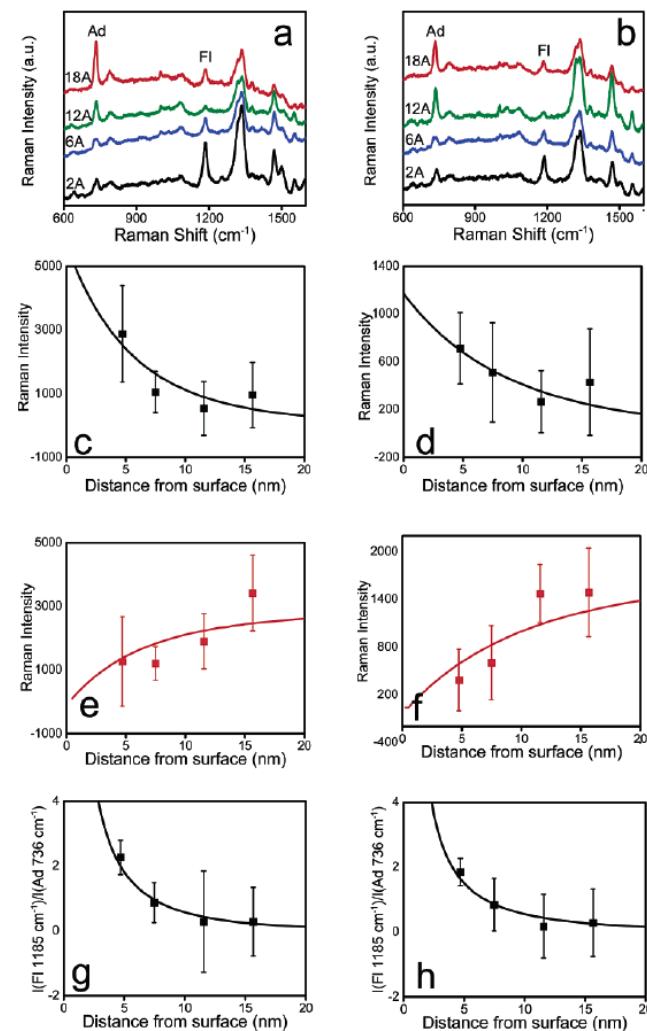
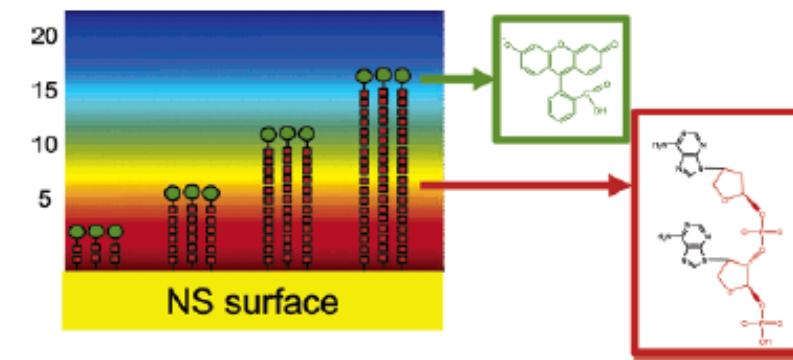


Figure 2. (A) SERS spectra of 5 μL of pMA with different concentrations deposited on the NP arrays. The excitation laser wavelength is 785 nm. Adsorption isotherm of pMA on the NP arrays obtained according to (B) 1077 and (C) 390 cm⁻¹ modes in the SERS spectra. I_0 is the peak intensity of a saturated pMA monolayer.

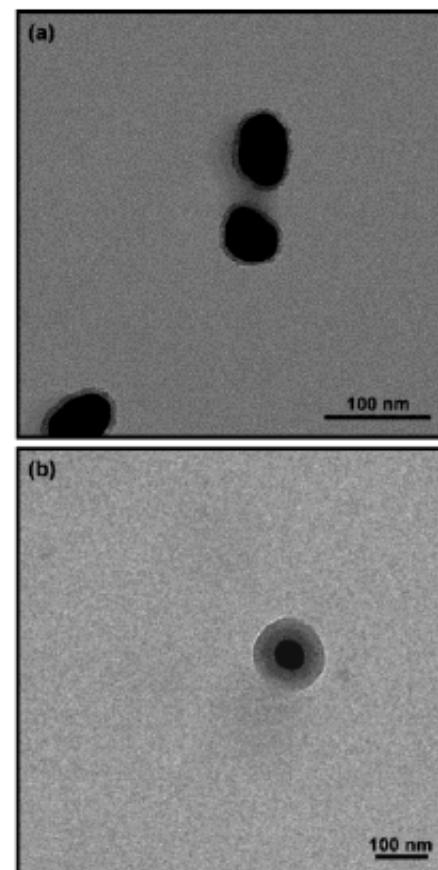
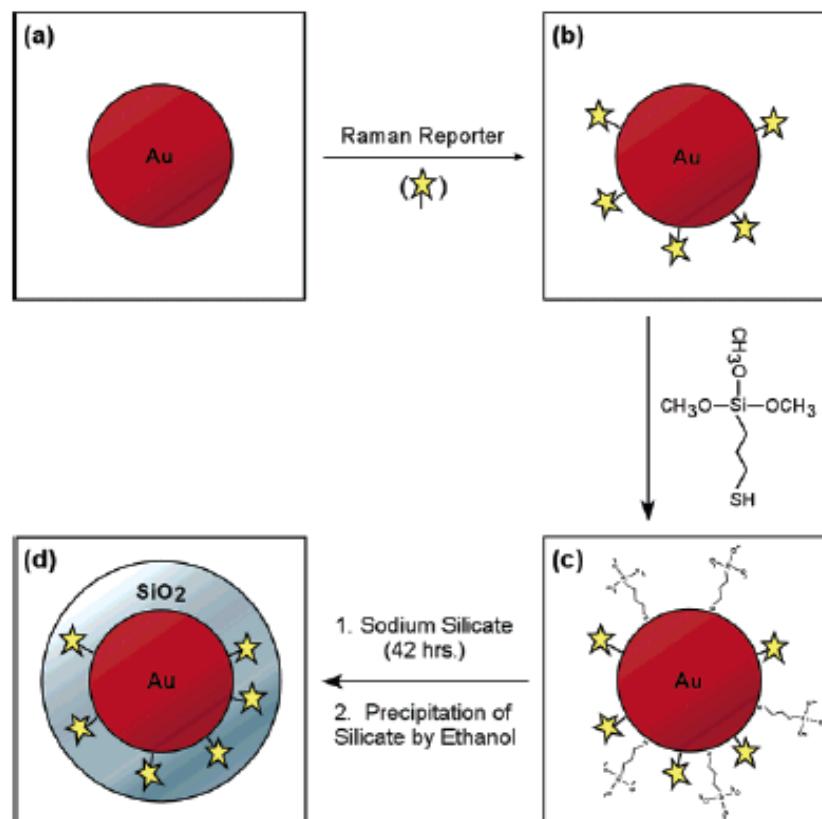


Profiling the Near Field of a Plasmonic Nanoparticle with Raman-Based Molecular Rulers

NANO
LETTERS
2006
Vol. 6, No. 10
2338–2343



Spectroscopic Tags Using Dye-Embedded Nanoparticles and Surface-Enhanced Raman Scattering



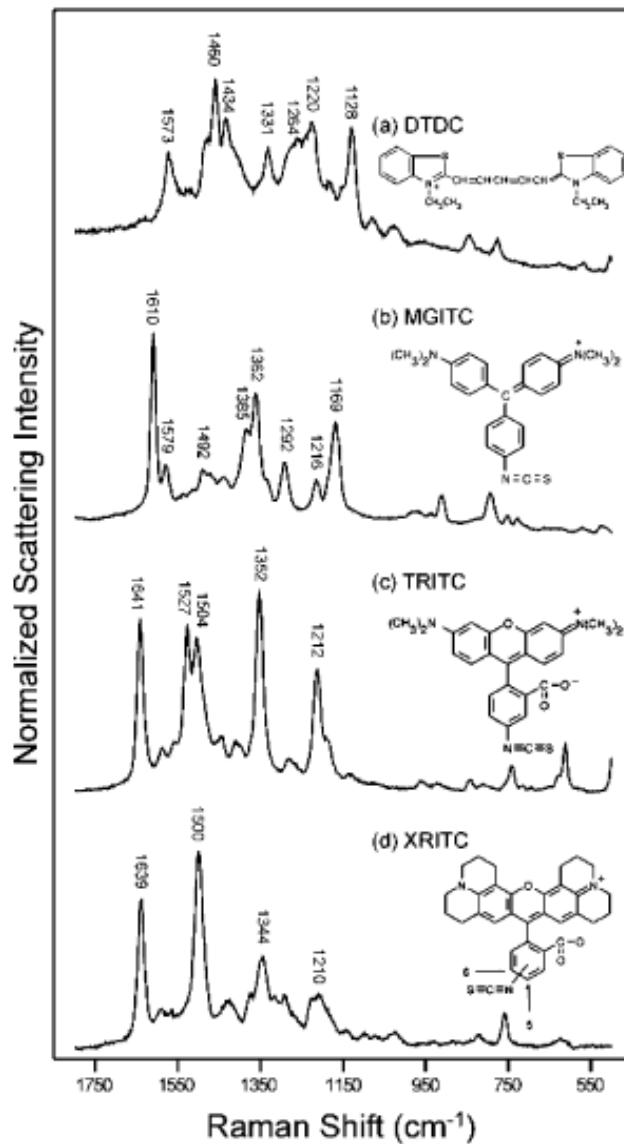
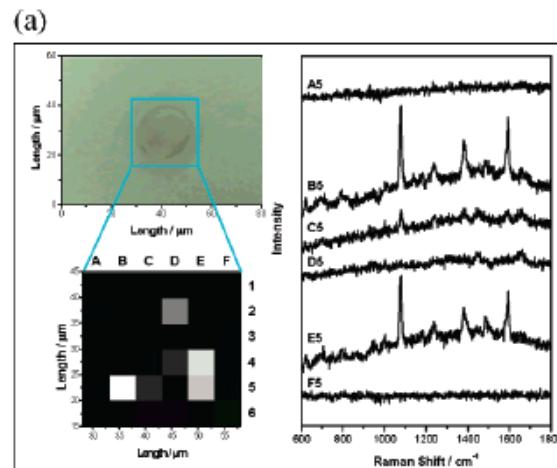
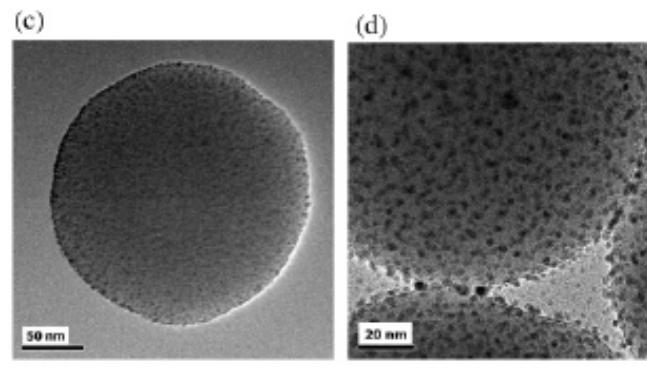
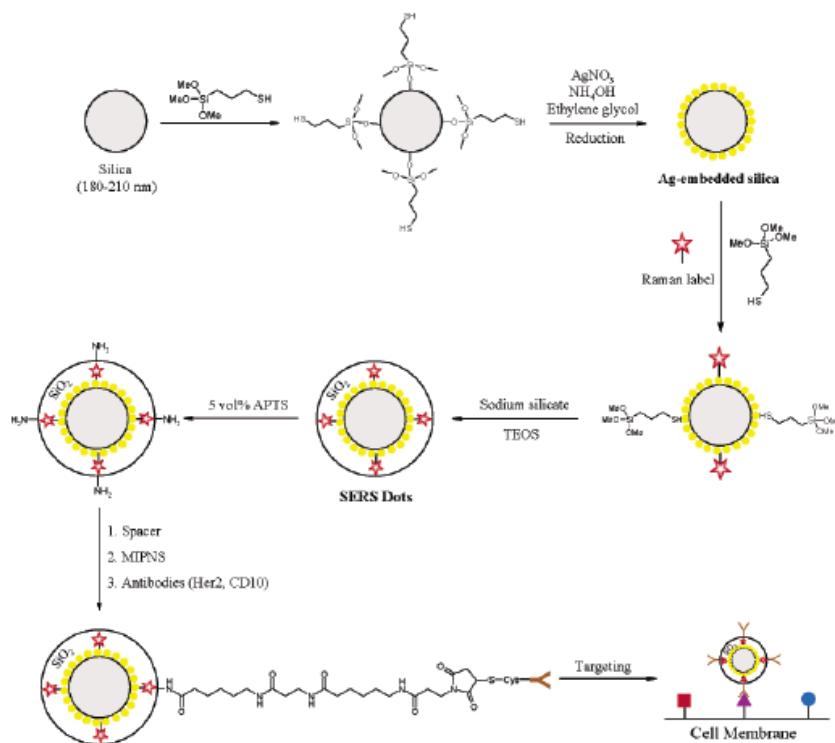


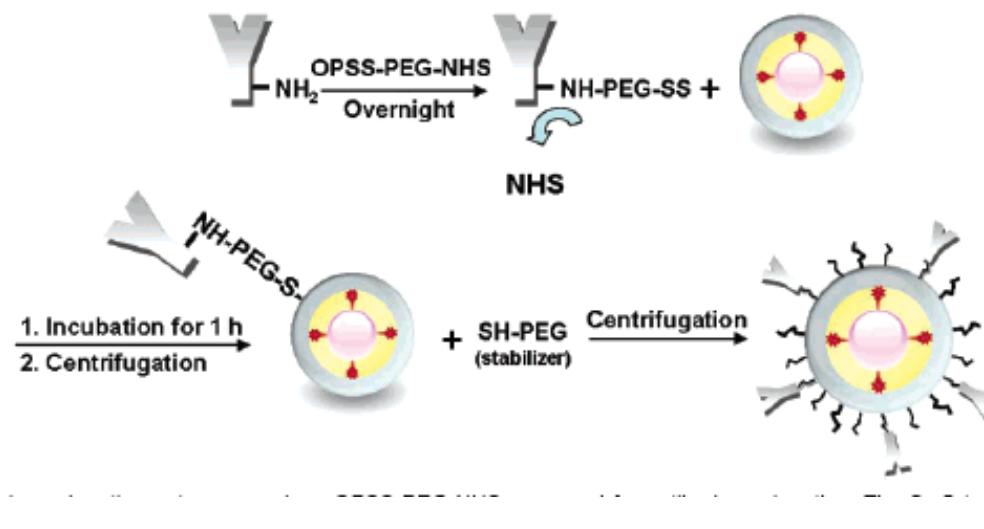
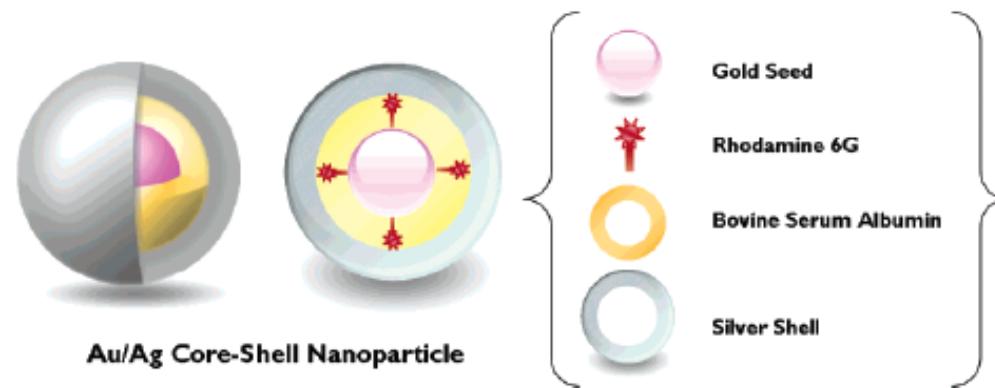
Figure 4. Chemical structures of four Raman reporters and their surface-enhanced resonance Raman spectra: (a) 3,3'-Diethylthiadicarbocyanine iodide (DTDC); (b) malachite green isothiocyanate (MGITC); (c) tetramethylrhodamine-5-isothiocyanate (TRITC); and (e) rhodamine-5-(and-6)-isothiocyanate (XRITC).



Nanoparticle Probes with Surface Enhanced Raman Spectroscopic Tags for Cellular Cancer Targeting



Biological Imaging of HEK293 Cells Expressing PLC γ 1 Using Surface-Enhanced Raman Microscopy



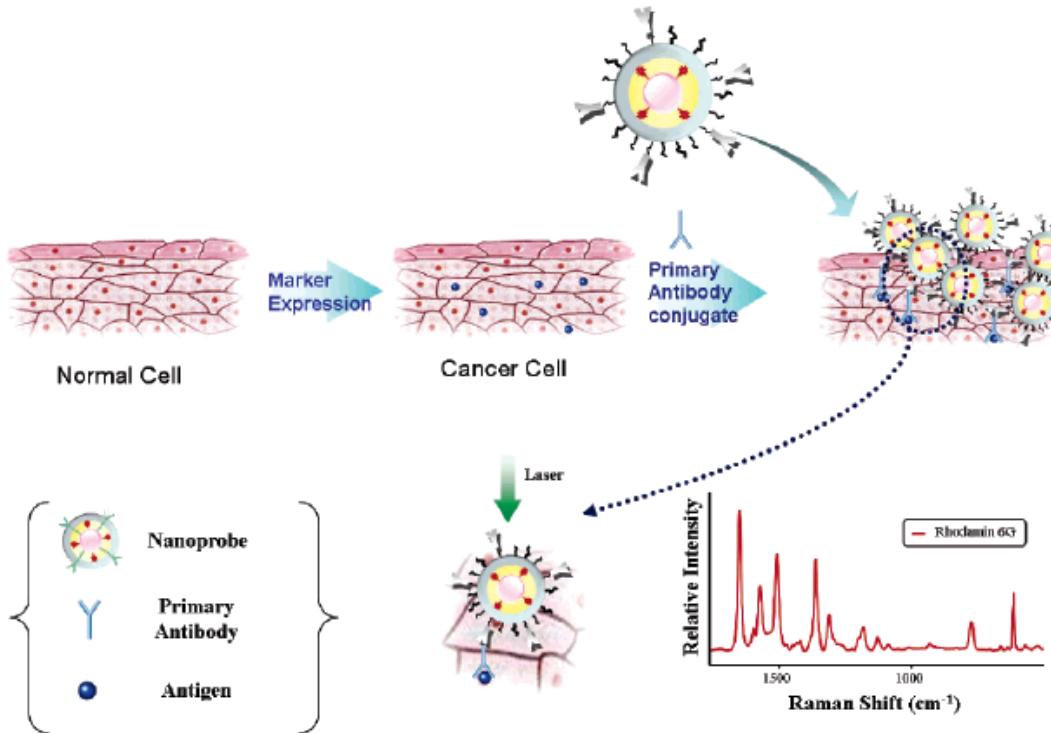


Figure 4. Schematic diagram depicting immobilization of Au/Ag core–shell nanoprobes on PLC γ 1-expressing HEK293 cells and their SERS detection.

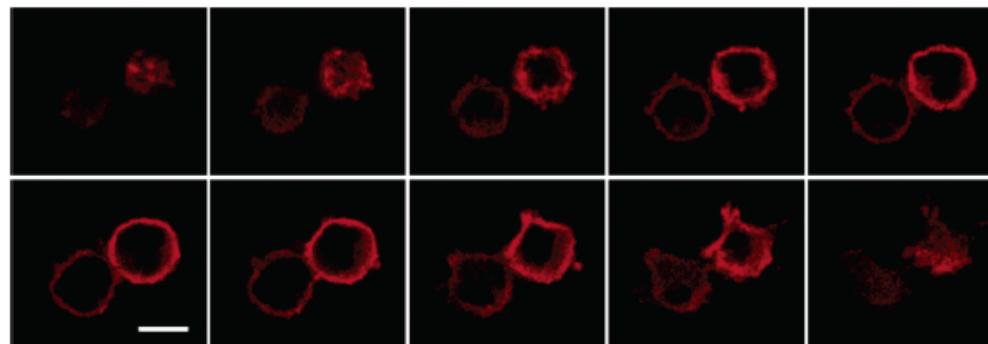


Figure 5. Serial fluorescence optical sections of PLC γ 1-expressing HEK293 cells using red QDs. The z-axis interval of optical slices is 1.3 μm . Cells were incubated for 30 min in red QDs, after which the free QDs were washed away. These fluorescence images indicate that PLC γ 1 markers are only expressed on the surface membranes. Scale bar, 10 μm .



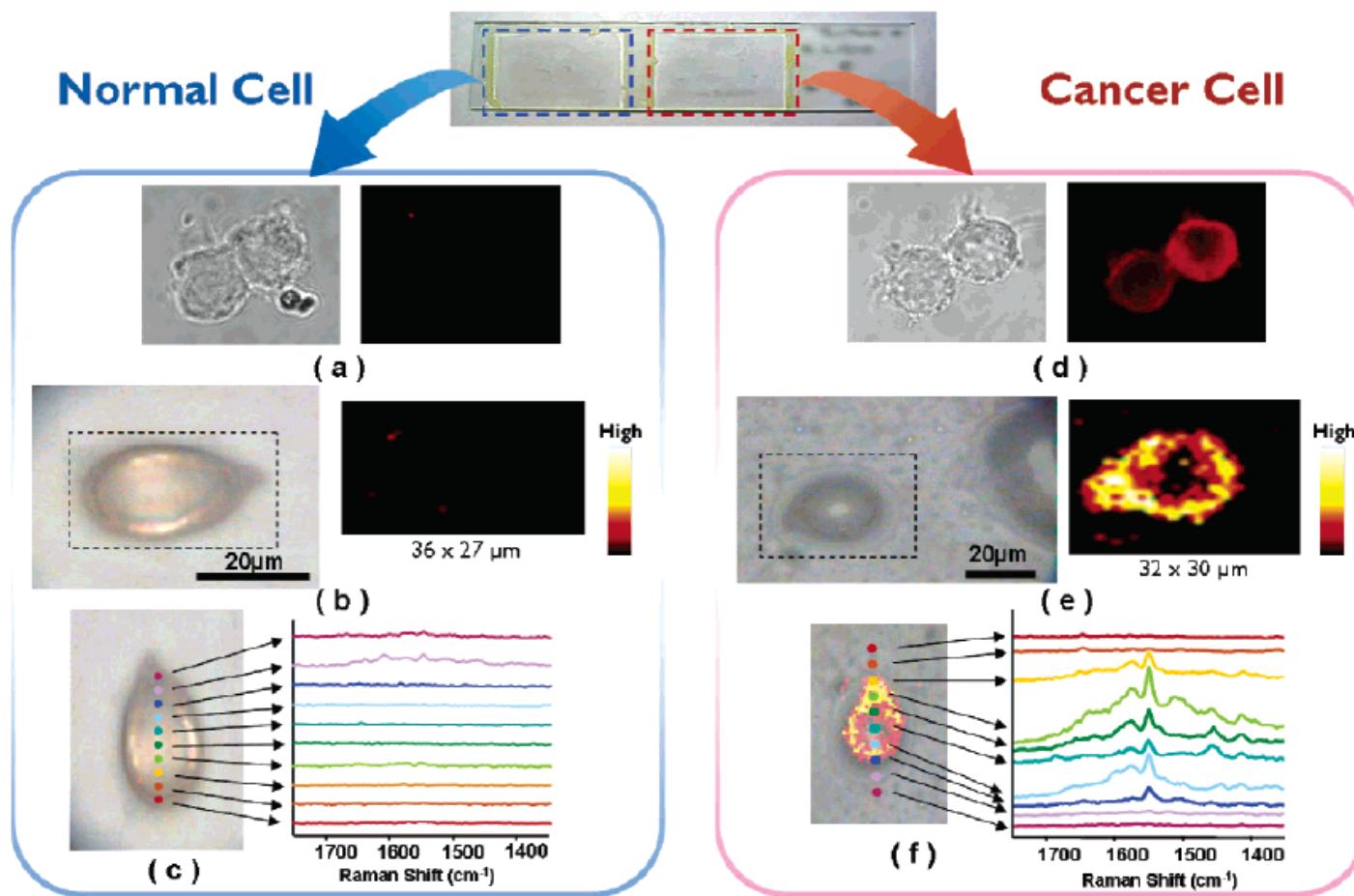


Figure 6. Fluorescence and SERS images of normal HEK293 cells and PLC γ 1-expressing HEK293 cells. (a) QD-labeled fluorescence images of normal cells: (left) brightfield image, (right) fluorescence image. (b) SERS images of single normal cell: (left) brightfield image, (right) Raman mapping image of single normal cell based on the 1650- cm^{-1} R6G peak. The cell area was scanned with an interval of 1 μm . Intensities are scaled to the highest value in each area. (c) Overlay image of brightfield and Raman mapping for single normal cell. Colorful spots indicate the laser spots across the middle of the cell along the y axis. (d) QD-labeled fluorescence images of cancer cells: (left) brightfield image, (right) fluorescence image. (e) SERS images of single cancer cell: (left) brightfield image, (right) Raman mapping image of single cancer cell based on the 1650- cm^{-1} R6G peak. The cell area was scanned with an interval of 1 μm . Intensities are scaled to the highest value in each area. (f) Overlay image of brightfield and Raman mapping for single cancer cell. Colorful spots indicate the laser spots across the middle of the cell along the y axis.



Mammalian Cell Surface Imaging with Nitrile-Functionalized Nanoprobes:
Biophysical Characterization of Aggregation and Polarization Anisotropy in
SERS Imaging

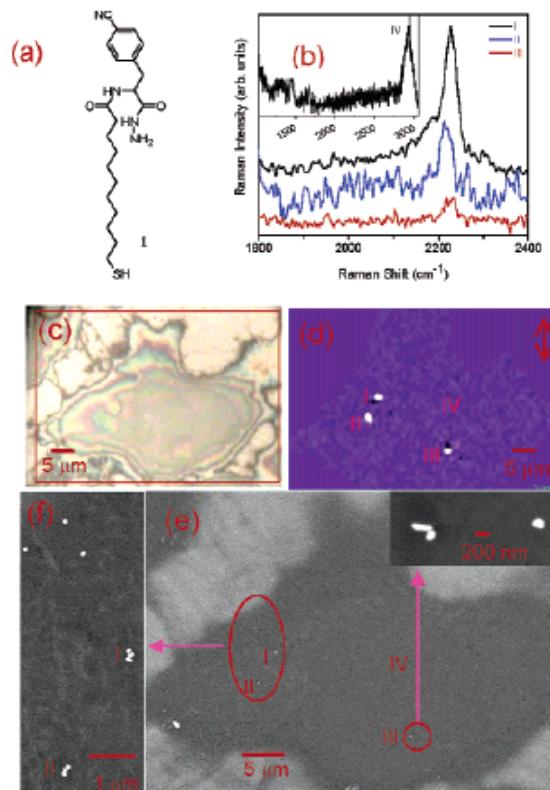


Figure 1. (a) The chemical structure of Raman reporter 1; (b) Raman spectra of the CN vibration mode extracted from positions I, II, and III of the cell shown in the optical image (c). Inset of (b) is a cellular Raman spectrum taken from spot IV of the same cell. (d) Raman intensity map of the C≡N band of the same cell, and (e) the corresponding SEM image. Inset in (e) showed the NPs in the lower right circle. (f) The group of NPs as shown in the large oval of (e).

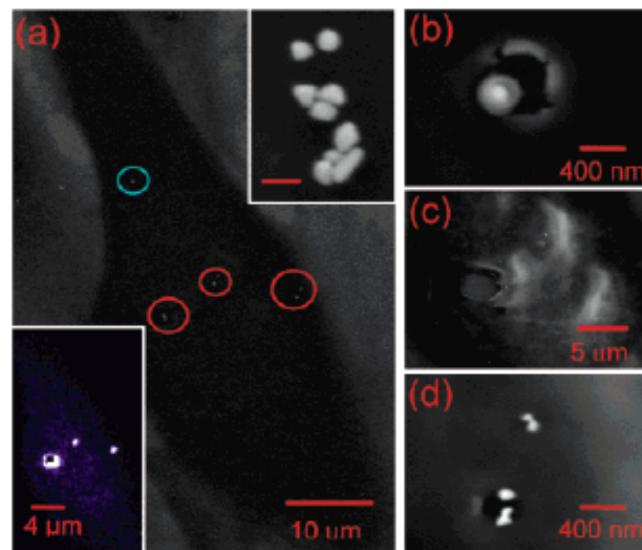


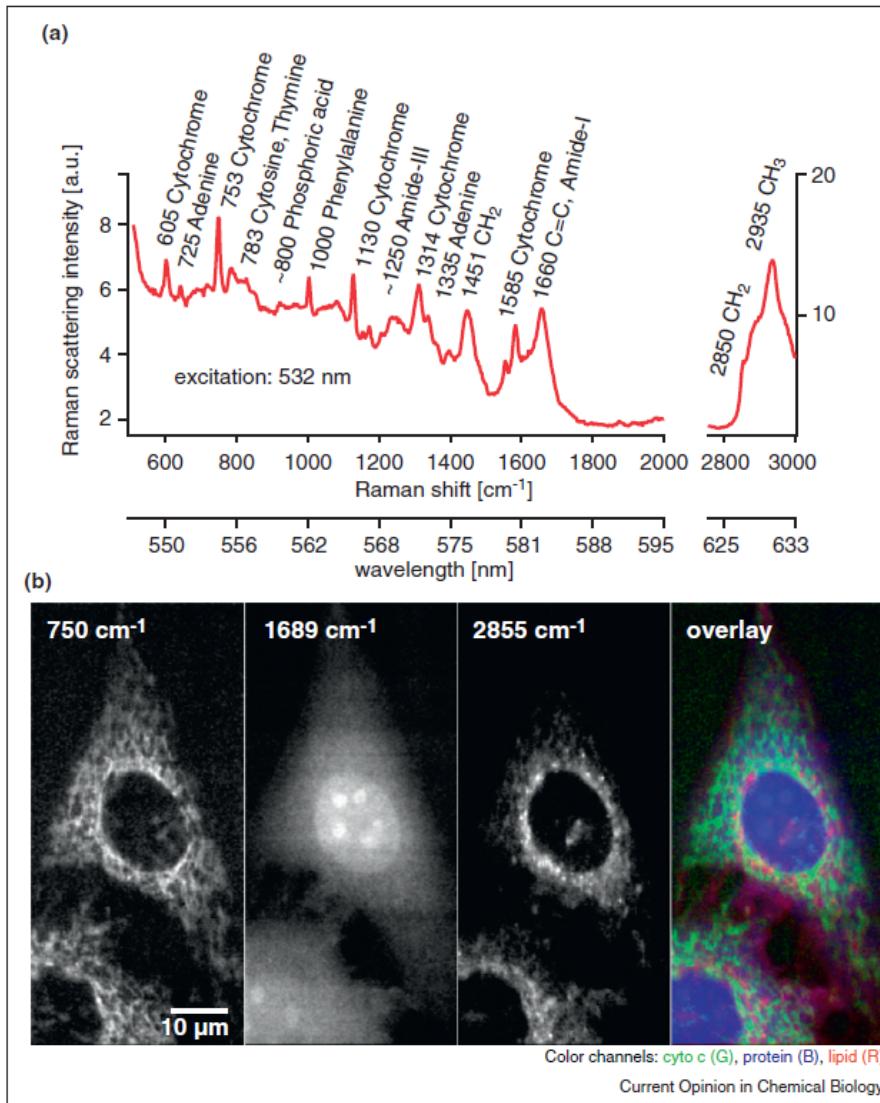
Figure 2. (a) SEM image of a cell. Upper right inset: magnification of a group of aggregated NPs. The scale bar is 200 nm. Lower left inset: the corresponding Raman intensity image of the same cell obtained with a power density of 10^5 W/cm 2 . Laser-induced damage to the cell is shown in (b) the monomer (blue circle in a), (c) the aggregates, and (d) a pair of dimers.

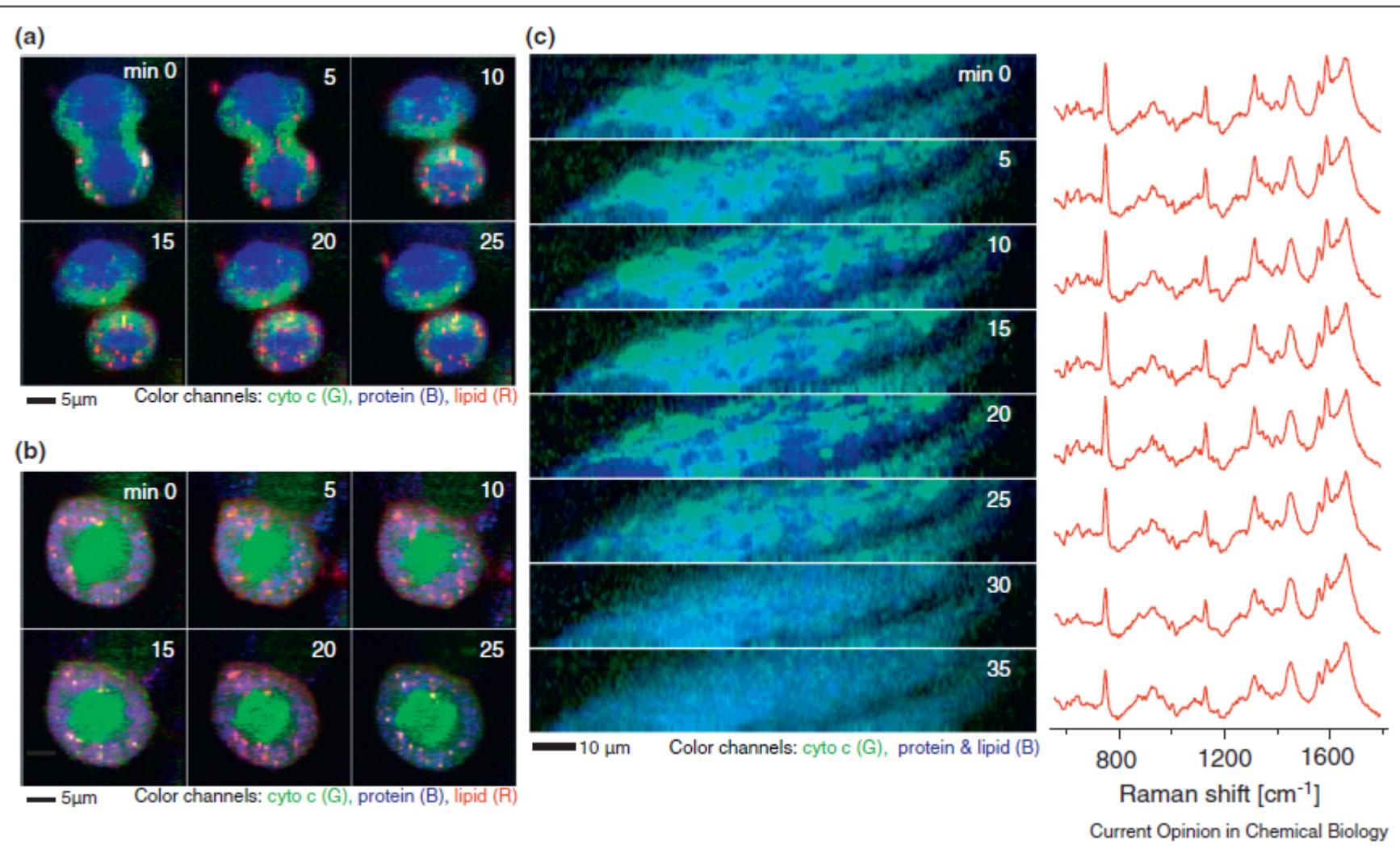


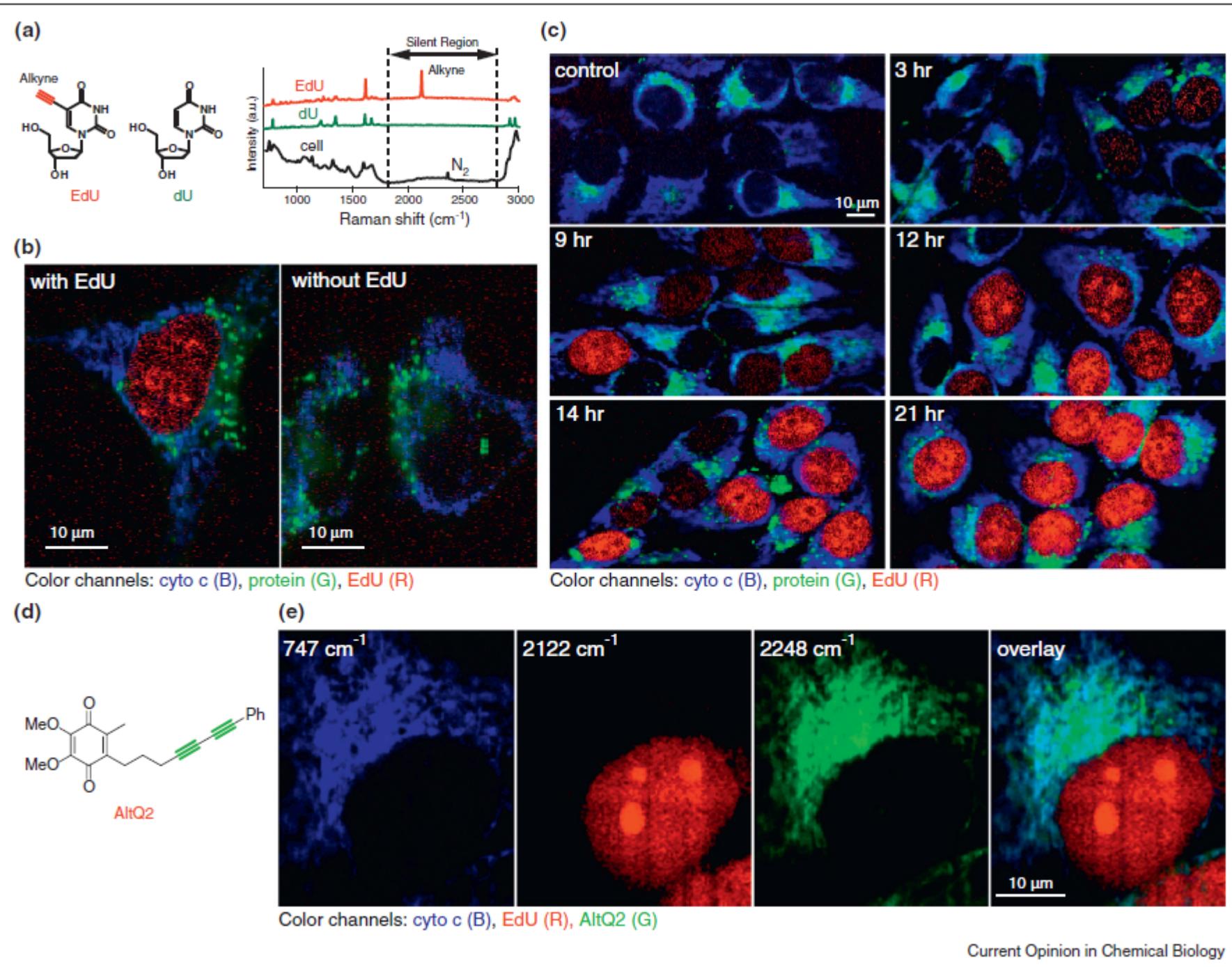
Molecular imaging of live cells by Raman microscopy

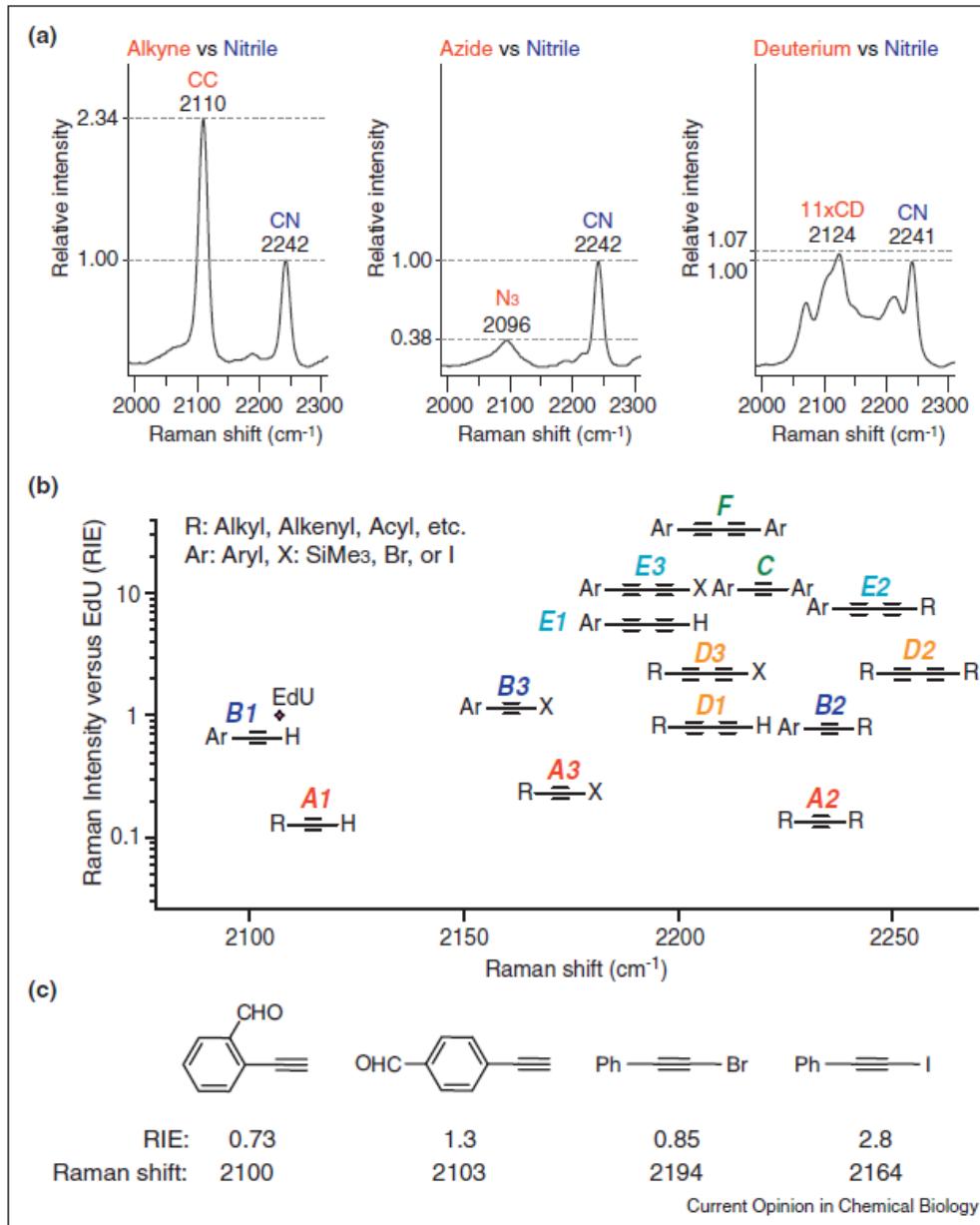
Almar F Palonpon^{1,2}, Mikiko Sodeoka^{2,3} and Katsumasa Fujita^{1,2}

Current Opinion in Chemical Biology 2013, 17:708–715

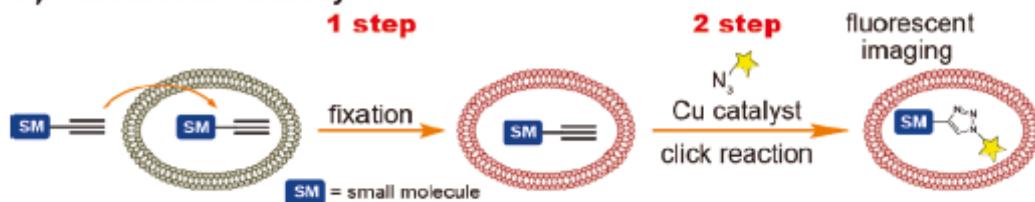




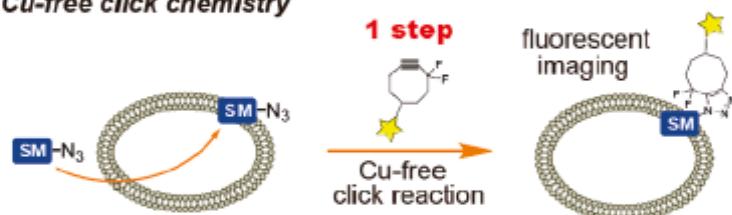




A) General click chemistry



B) Cu-free click chemistry



C) This time (Click-free)



Figure 1. Concept of click-free imaging.

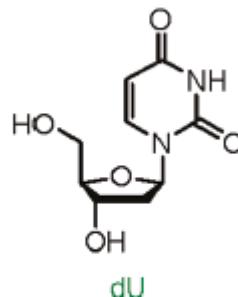
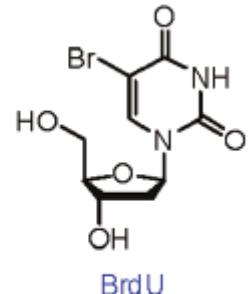
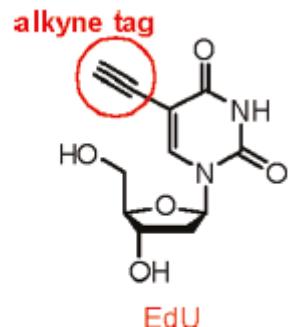
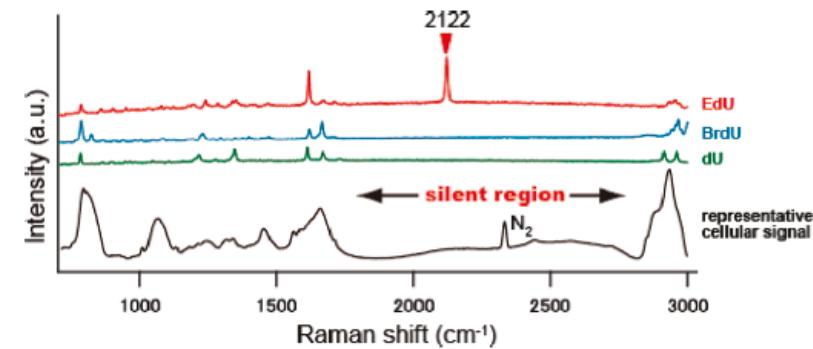
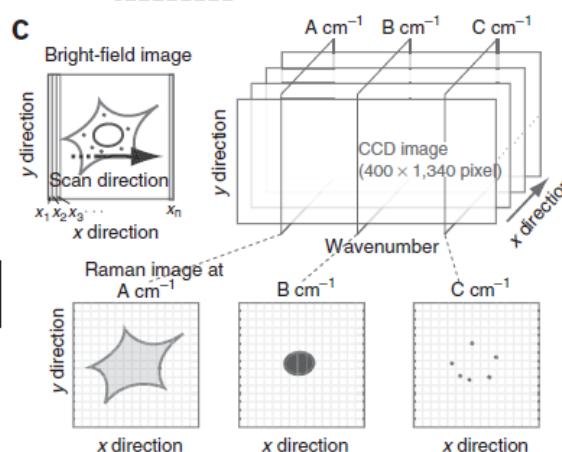
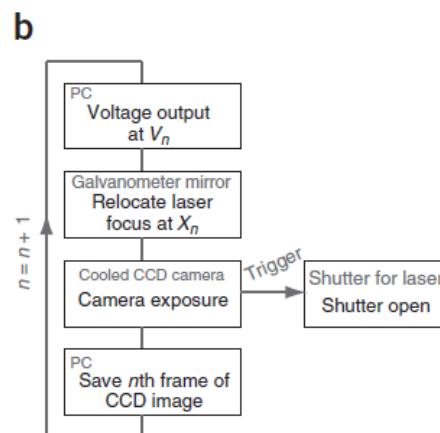
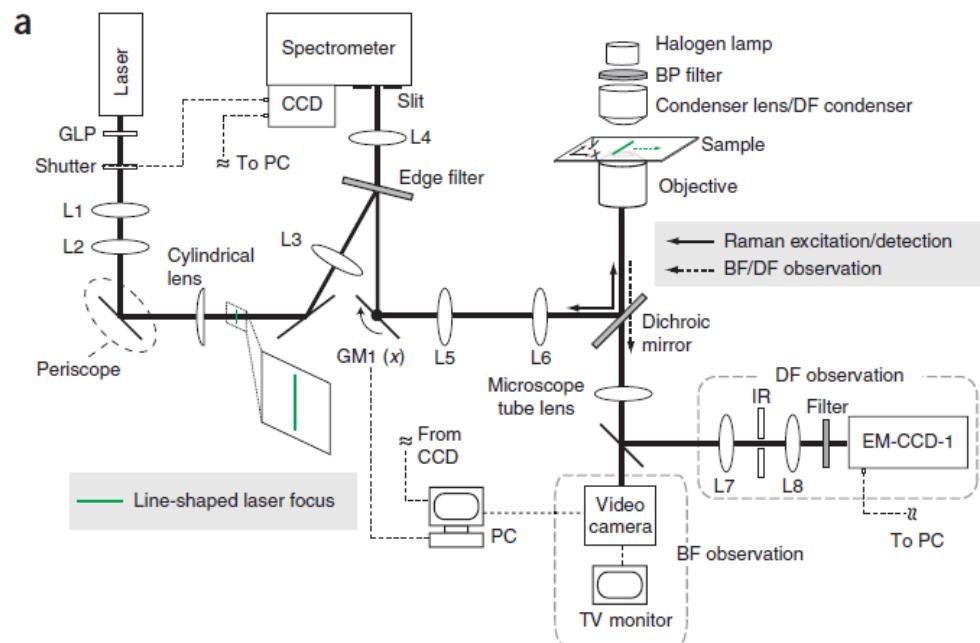


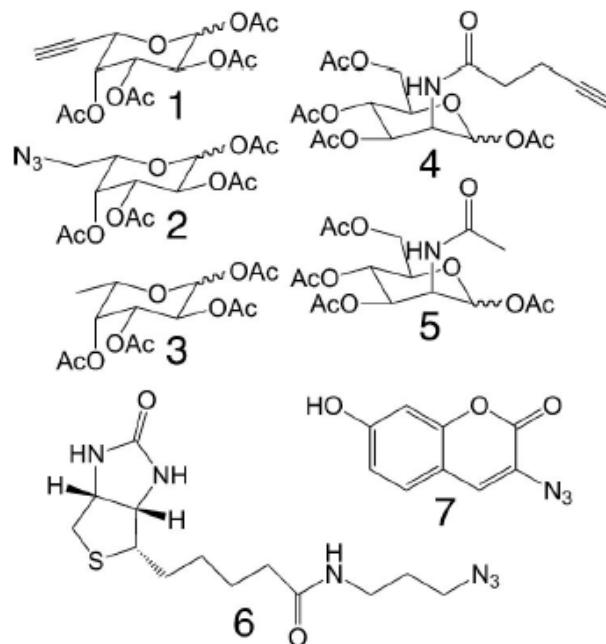
Figure 2. Structures of thymidine analogues.



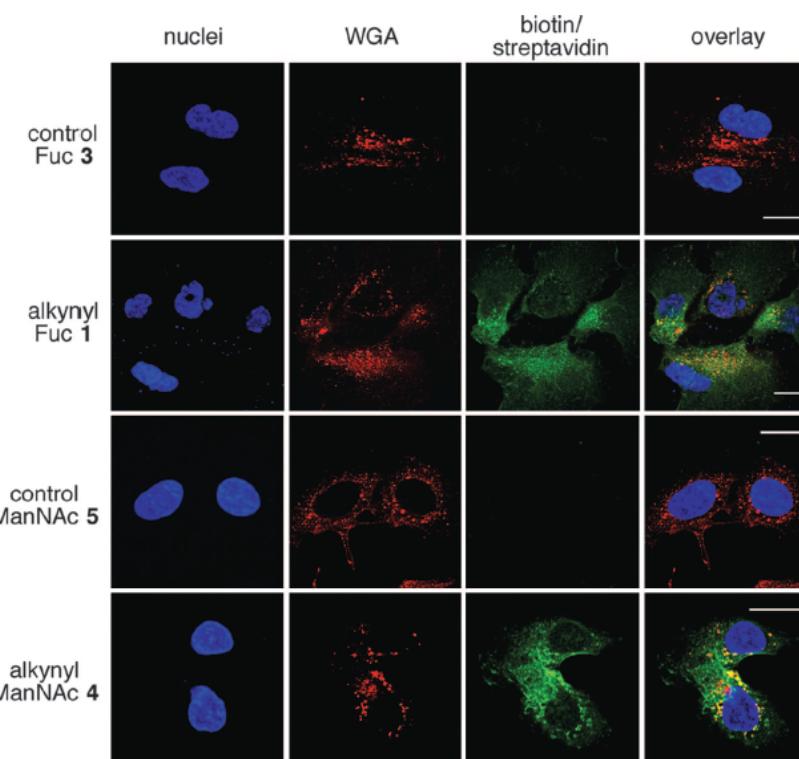
PROTOCOL |



Alkynyl sugar analogs for the labeling and visualization of glycoconjugates in cells



Scheme 1. Modified sugar analogs and probes used in this study.



Cell-permeable probe for identification and imaging of sialidases

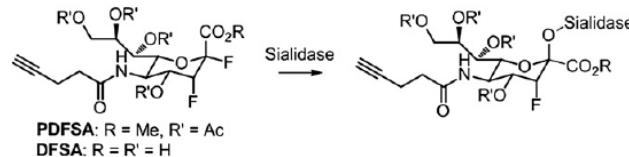
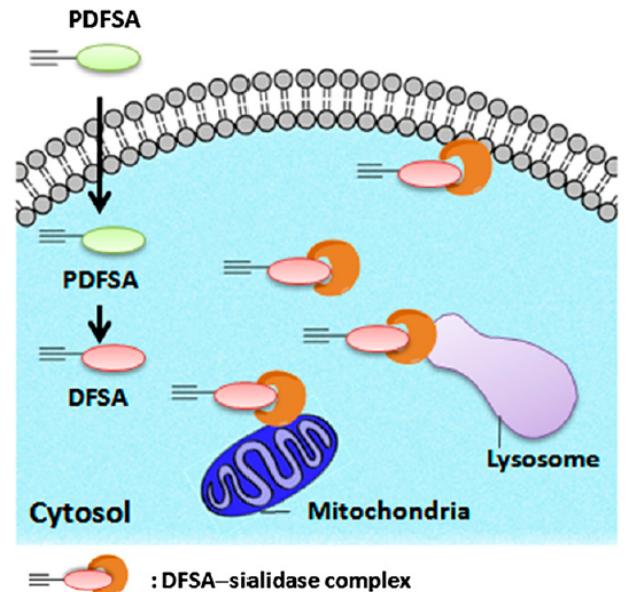


Fig. 1. Identification and imaging of sialidase with activity changes using these activity-based sialidase probes.

