

**Cell Surface-based Sensing with Metallic Nanoparticles**

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Complete List of Authors:	Jiang, Ziwen; University of Massachusetts--Amherst, Chemistry Le, Ngoc; University of Massachusetts--Amherst, Chemistry Gupta, Akash; University of Massachusetts--Amherst, Chemistry Rotello, Vincent M; University of Massachusetts at Amherst, Dept. of Chemistry (710A LGRT)

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1 **Cell Surface-based Sensing with Metallic Nanoparticles**

2 **Ziwen Jiang, Ngoc D. B. Le,[†] Akash Gupta,[†] Vincent M. Rotello***

3 Department of Chemistry, University of Massachusetts Amherst, 710 North Pleasant St, Amherst,
4 Massachusetts, MA 01003, USA

5 [†]These authors equally contributed to this work.

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7 * Correspondence should be addressed to Vincent M. Rotello (V. M. R.).

8 **Address:**

9 Department of Chemistry

10 710 North Pleasant Street

11 University of Massachusetts

12 Amherst, MA 01003, USA.

13 **Tel.:** +1 413 545 2058; **fax:** +1 413 545 4490.

14 **E-mail:** rotello@chem.umass.edu.

15

16 **Abstract**

17 Metallic nanoparticles provide versatile scaffolds for biosensing applications. In this
18 review, we focus on the use of metallic nanoparticles for cell surface sensings. Examples of the
19 use of both specific recognition and array-based “chemical nose” approaches to cell surface
20 sensing will be discussed.

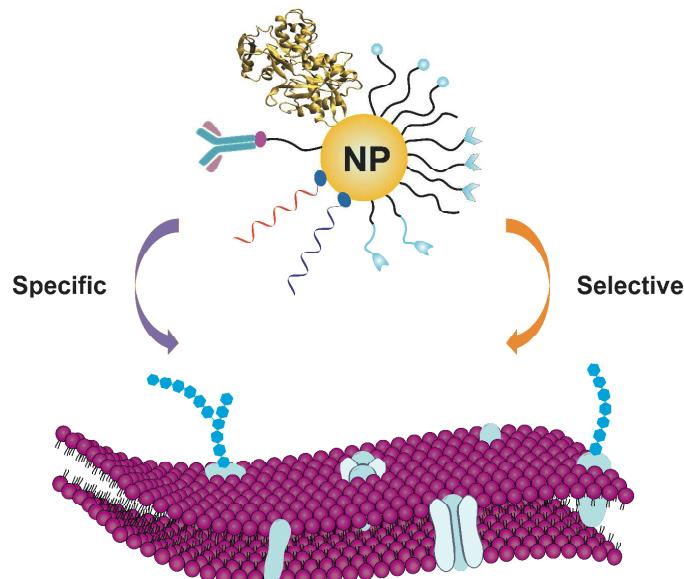
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22 **Key learning points**

- 23 1. Both cell surface biomarkers (such as carbohydrates and proteins) and the overall cell surface
24 signatures provide crucial information for identifying cell types.
- 25 2. Metallic nanoparticles provide multiple modes of signal transduction for biosensing
26 applications.
- 27 3. Surface functionalization determines how nanoparticles interact with cell surfaces.
- 28 4. The specific recognition capabilities of biomacromolecules such as antibodies, lectins,
29 aptamers, and DNAzymes can be coupled with nanoparticle transduction processes to design cell
30 sensing strategies.
- 31 5. Nanoparticle surface can be functionalized with a variety of small molecule ligands to provide
32 the selective recognition required for array-based sensing.

33

34 **TOC**



35

36

37 **1. Introduction**

38 Cell surface sensors for disease and detection of infection have direct access to the
39 sensing target, in contrast to approached that detect intracellular proteins, nucleic acids, or other
40 markers buried inside the cells. This ready access has the potential to provide rapid sensing with
41 minimal processing. The rich environment presented by the cell exterior also gives cell surface
42 sensors the capability to read out the phenotypes of cells, a property that is the final outcome of
43 multiple factors including both genetic and epigenetic variations.¹ For example, in the case of
44 cancer, abnormal cells have been found to overexpress specific glycosylated proteins at their
45 plasma membrane such as epithelial cell adhesion molecule (EpCAM) or carcinoembryonic
46 antigen (CEA).²⁻⁴ Therefore, targeting cell surface phenotype provides a strategy for simple,
47 rapid, and robust diagnostic pathways in diverse areas such as cancer and pathogenic bacteria.

48

49 Three integrated components are necessary to fabricate an effective sensor: (1) a
50 recognition element to interact with a target analyte, (2) a signal transduction element to generate
51 a measurable signal from an analyte-receptor binding event, and (3) a device that outputs a result.
52 Metallic nanoparticles (NPs) can be easily engineered to provide scaffolds for recognition
53 processes, with their physical properties facilitating the transduction process, making them
54 excellent platforms for cell surface sensing.^{5,6}

55

56 In this review, we will focus on the use of metallic NPs for the detection and
57 quantification of cell properties, based on cell surface components. We will discuss examples of
58 different engineered metallic NP systems^{7,8} that provide cell sensing through specific and
59 selective interactions with the cell surfaces.

60

61 **2. Cell surface and nanoparticle interactions**

62 Enormous cell surface diversity exists among cells from plants, bacteria, and animals.
63 The surface of a mammalian cell is composed of a complex structure featuring the lipid bilayer,
64 proteins, nucleic acids as well as a range of polysaccharide structures that comprise the
65 glycocalyx.⁹ This glycocalyx is composed of glycoproteins, proteoglycans and glycolipids.¹⁰
66 Phenotypically altered expression of each of these components provides diagnostic information
67 for diseases such as cancer, Gaucher's, and Tay-Sachs diseases.^{11,12} Taken together, the complex

68 array of biomolecules that comprise cell surfaces make them excellent targets for both specific
69 biomarker sensing and selective “chemical nose” based methods.

70

71 The interaction between nanomaterials and cells is an important issue for designing
72 systems not only for sensing, but also for imaging and delivery. In general, the following factors
73 need to be taken into account: (1) specific receptors (biomarkers) on the cell membrane, (2) the
74 size, shape, surface charge, roughness and hydrophobicity of nanoparticles and their role in
75 selective interactions. While these topics are all central to the sensing described here, the in-
76 depth discussion required for understanding this interaction is beyond the scope of this tutorial
77 discussion. Nel and coworkers have provided a comprehensive review to help understand the
78 biophysicochemical interactions at the nano-bio interface, which discussed cell-nanoparticle
79 interactions in detail.¹³

80

81 **3. Specific sensing**

82 We will focus on spherical metallic nanoparticles in this review, as these systems have
83 been widely employed for cell surface sensing. Metallic nanoparticles can be functionalized with
84 small molecules¹⁴ and biomacromolecules¹⁵ to achieve the specific interactions with the
85 biological targets. However, the vast majority of specific-based sensors have been using
86 biomacromolecules to functionalize metallic nanoparticles, so we will focus on these
87 bioconjugate systems. These platforms provide highly adaptable tools for rapid and/or point-of-
88 care tools that provide alternatives to more complex and instrument-intensive techniques such as
89 flow cytometry.¹⁶

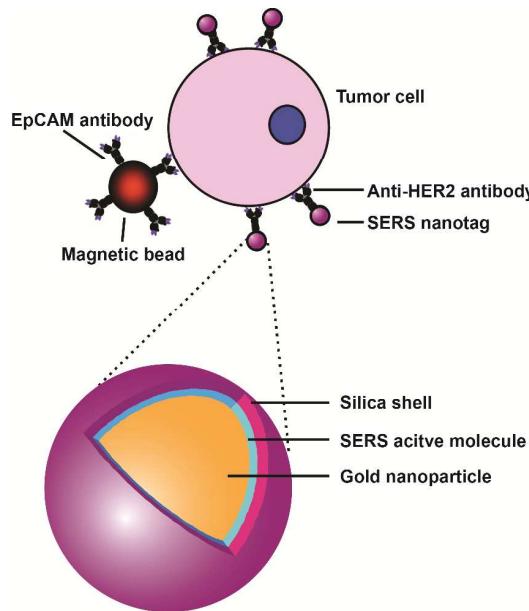
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91 **3.1. Antibody-based sensing**

92 Antibodies are widely used as recognition elements in diagnostic and therapeutic
93 applications.¹⁷ There are two key components of antibodies: the Fab (fragment, antigen-binding)
94 region of an antibody that recognizes the antigen and the Fc (fragment, constant) that can be used
95 for conjugation without disrupting the recognition process. Conjugation of either complete
96 antibodies or Fab fragments to NPs provides an effective means of recognizing cell surface
97 functionality. As described below, these binding events can be detected *via* various tools such as
98 surface-enhanced Raman scattering (SERS) and electrochemistry.

99

SERS is a technique in which the Raman signal can be dramatically amplified through surface plasmon resonance of metallic NPs.⁶ SERS-based techniques utilizing antibodies have been successfully applied to immunoassay-based methodologies.¹⁸ In one example, Sha and coworkers detected cancer circulating cells (CTCs) by combining capturing capability of a magnetic bead and specific labeling of SERS nanotags.¹⁹ This bead was conjugated with anti-EpCAM antibody to capture SKBR3 cancer cells. These cells were then labeled for SERS detection by AuNPs functionalized with anti-HER2 antibody (human epidermal growth factor receptor-2). A silica shell was subsequently coated on this complex to enable the functionalization of antibody on the particle without interfering with the Raman response (Figure 1). In a similar study, SERS-based systems were further employed for *in vivo* tumor targeting. Poly(ethylene glycol)-capped AuNPs were used to stabilize Raman-active reporter molecules. The SERS-NPs were conjugated with antibodies specifically targeting the overexpressed epidermal growth factor receptor on tumor cells, resulting in highly specific *in vivo* tumor detection.²⁰



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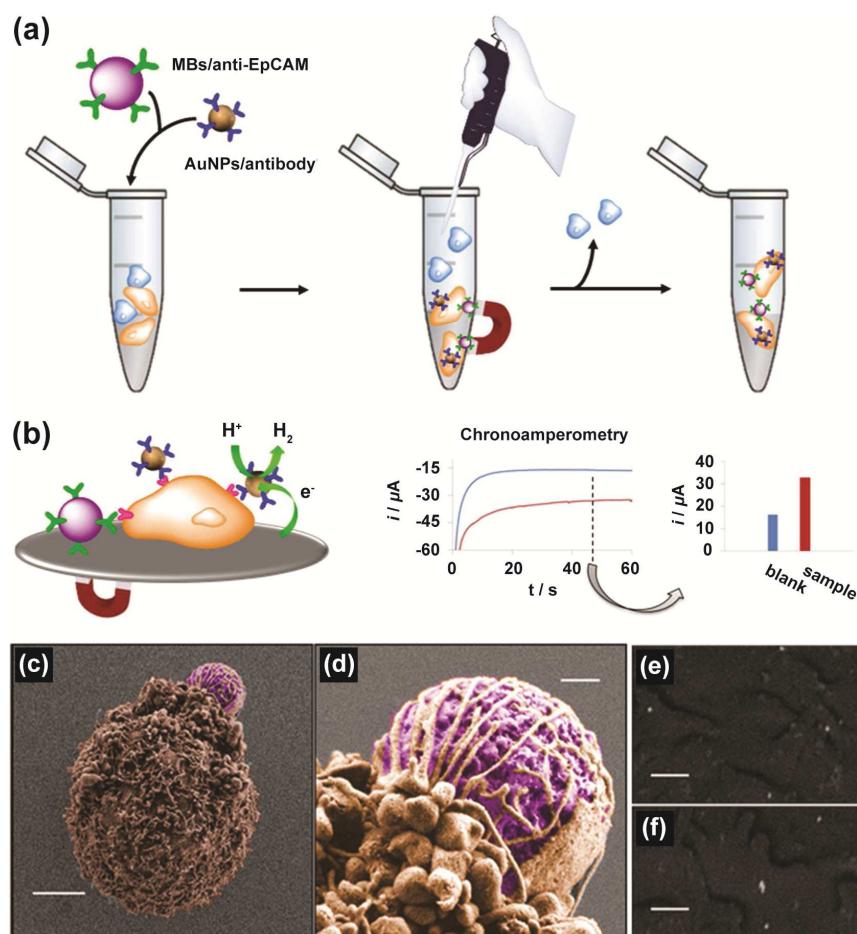
Figure 1 Tumor cell detection using anti-HER2 antibody-conjugated magnetic beads with SERS nanotags. Reprinted with permission from ref. 19. Copyright 2008 American Chemical Society.

117

118 Electrochemical detection technique can utilize the electrocatalytic properties of AuNPs
119 to provide rapid cell surface sensing.²¹ This strategy has the advantages of simpler

120 instrumentation coupled with the direct connection of sensor output with devices/computers. For
121 example, Merkoçi and coworkers employed antibody-conjugated magnetic beads and AuNPs for
122 the detection of CTC (Figure 2).²² AuNPs fabricated with anti-EpCAM antibody were used for
123 targeting EpCAM, an overexpressed transmembrane glycoprotein on human colon
124 adenocarcinoma cells (Caco2 cells). The AuNP-antibody conjugates were used to generate an
125 electrochemical signal through electrocatalytic hydrogen evolution. The signals generated from
126 AuNP-antibody conjugates could detect 2.2×10^2 Caco2 cells in the presence of other interfering
127 cells such as monocytes (THP-1).

128



129

130 **Figure 2** (a) Capture of Caco2 cells by magnetic beads conjugated to anti-EpCAM.
131 Simultaneously, cells were labeled with AuNP-specific antibodies in the presence of interfering
132 cells (THP-1). (b) Chronoamperometry of the hydrogen evolution reaction (HER)
133 electrocatalyzed by AuNPs. (c), (d) False colors scanning electron microscopy (SEM) images of
134 a Caco2 cell captured by magnetic beads (MBs)/anti-EpCAM. (e), (f) Backscattered images

135 showing AuNPs distributed along the cell plasma membrane of Caco2 cells. Scale bars, 3 μm (c),
136 400 nm (d), and 200 nm (e, f). Reprinted with permission from ref. 22. Copyright 2012
137 American Chemical Society.

138

139 **3.2. Lectin-based sensing**

140 Lectins are proteins that exhibit strong and specific or selective binding towards
141 carbohydrate moieties. Targeting carbohydrates has been the useful strategy for diagnosis
142 because the alterations of carbohydrates found on plasma membrane have been correlated with
143 disease, such as liver fibrosis, pancreatic cancer, and cervical cancers.²³ Thus, NPs
144 functionalized with lectins can be a powerful tool for cell surface sensing.²⁴

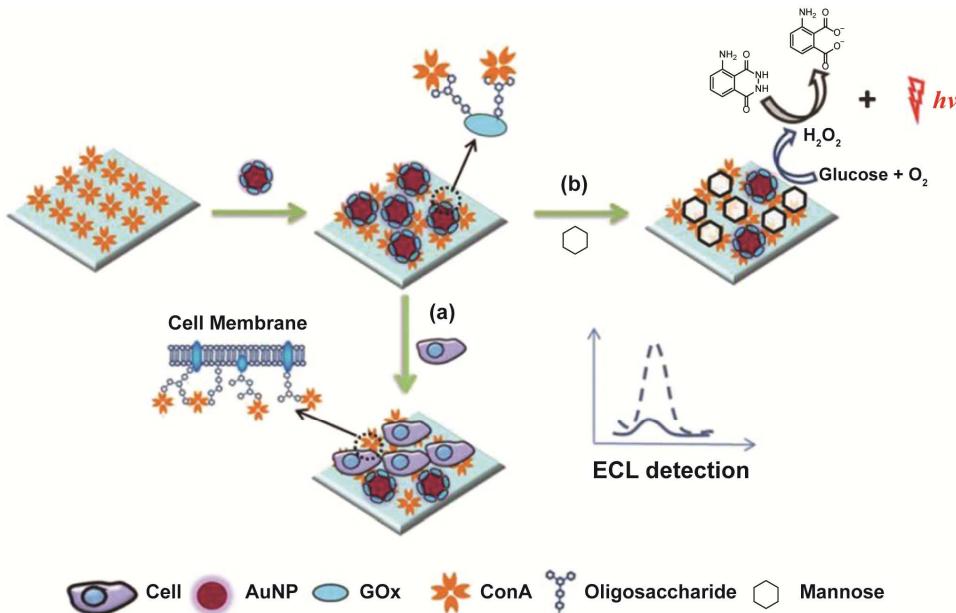
145

146 In a recent study, Liu and coworkers designed a sensitive electrochemiluminescence²⁵
147 (ECL)-based biosensor using a displacement assay that relies on the interaction between NP-
148 bound lectins and carbohydrates on the cell surface (Figure 3).²⁶ In this system, a gold electrode
149 immersed in luminol solution was coated with Concanavalin A (ConA), a lectin that recognizes
150 mannose (a carbohydrate type found on the cell surface). These mannose moieties can also be
151 found in glucose oxidase molecules (GOx), an enzyme that can catalyze the luminol ECL
152 reaction. By coupling GOx with AuNPs (GOx-Au), they were able to fabricate a multifunctional
153 probe. This GOx-Au probe can both compete with mannose moieties on the analyte cells for
154 ConA-coated gold electrode and improve the ECL signal of luminol. In the presence of the target
155 cells, the competition between GOx-Au and mannose-containing cells generates the alterations in
156 ECL signal intensity, providing the ability to profile carbohydrate-lectin interaction and *in situ*
157 cell surface carbohydrate expression.

158

159

160



161

162 **Figure 3** Schematic illustration of the lectin-based sensing strategy for (a) carbohydrate-ConA
 163 interaction analysis and (b) cell surface carbohydrate expression. Reprinted with permission from
 164 ref. 26. Copyright 2013 Royal Society of Chemistry.

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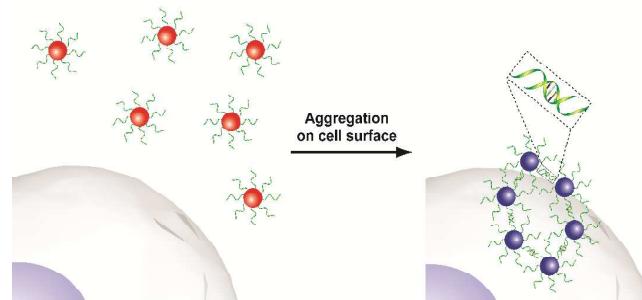
166 3.3. Aptamer-based sensing

167 Short, single-stranded oligonucleotides (ssDNA or ssRNA), known as aptamers, provide
 168 an emerging strategy for biorecognition. Aptamers are produced from an *in vitro* method known
 169 as SELEX (systematic evolution of ligands by exponential enrichment). In this process, SELEX
 170 uses polymerase chain reaction (PCR) to specifically amplify the sequence that has high affinity
 171 and selectivity to the target analyte. The iterative process generates aptamers that often fold into
 172 unique three-dimensional conformations. Aptamers can bind to target molecules ranging from
 173 small organic molecules to biomacromolecules,²⁷ making them promising candidates to serve as
 174 recognition elements in biosensors.

175

176 The recognition capabilities of aptamers can be combined with the spectroscopic
 177 advantages of AuNPs for cell detection applications. AuNPs possess strong distance dependent
 178 optical properties due to surface plasmon resonance.²⁸ The aggregation of aptamer-conjugated
 179 AuNPs causes a shift in their absorption spectra, resulting in a change in their scattering profile
 180 and color from red to blue/purple.²⁹ This colorimetric sensing method has been applied using
 181 aptamers for the detection of cancer cells. For instance, Tan and coworkers successfully applied

182 the aggregation-based colorimetric sensing platform to detect cancer cells using AuNPs
183 functionalized with the aptamers of interest.³⁰ The specific interaction between AuNP-aptamer
184 conjugates and the target cells (CCRF-CEM acute leukemia cell) induced a distinct color change
185 (Figure 4).

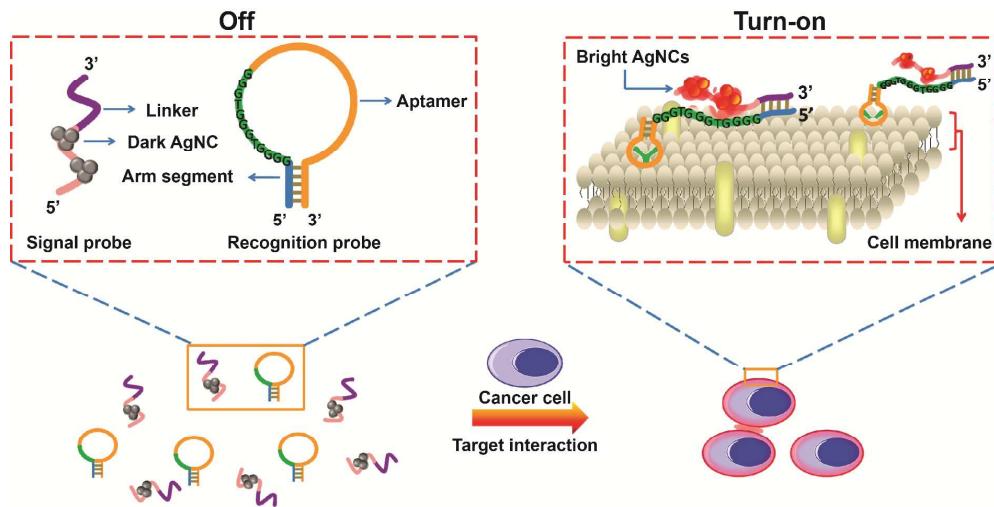


186

187 **Figure 4** Aptamer-conjugated gold nanoparticles used in colorimetric sensing of cancer cells.

188

189 Recently, DNA-templated silver nanoclusters (DNA-AgNCs) were used as a signal
190 transduction element for use with aptamers. The fluorescence of DNA-AgNCs can be
191 significantly amplified in proximity of guanine-rich DNA sequences,³¹ a phenomena Wang and
192 coworkers have applied in cell surface-based sensing.³² They designed a “turn-on” system for
193 cancer cell detection, utilizing the fluorescence enhancement of DNA-AgNCs and the
194 recognition capability of aptamers. Two separate DNA-based probes were involved in this
195 system, denoted as the recognition probe and signal probe. The recognition probe was designed
196 as a hairpin-shaped structure that contains a CCRF-CEM cancer cell specific aptamer sequence,
197 a guanine-rich DNA sequence and an arm segment. The signal probe contains a sequence for
198 AgNC-templated synthesis and a link sequence that is complementary to the arm segment of the
199 recognition probe. Once the aptamer sequence from the recognition probe recognizes and binds
200 to CCRF-CEM cells, the recognition probe undergoes a conformational alteration. This
201 conformational alteration then initiates the hybridization of the two probes and consequently
202 brings DNA-AgNCs close to the guanine-rich DNA sequence, resulting in an enhanced
203 fluorescence readout (Figure 5).

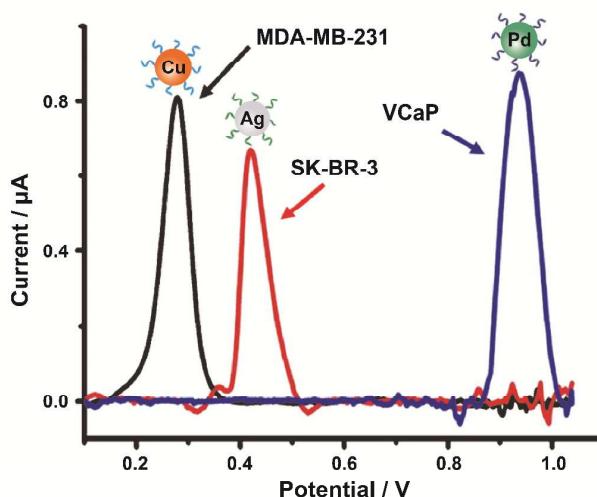


204

205 **Figure 5** Schematic representation of cancer cell detection based on DNA-templated silver
206 nanoclusters (AgNCs). Reprinted with permission from ref. 32. Copyright 2013 American
207 Chemical Society.

208

209 Aptamer-based specific sensing can also be used to detect different cancer cells using an
210 electrochemical approach. By combining multiple metallic NPs with electrochemical analysis,
211 Kelley and coworkers designed a chip-based strategy for the analysis of cancer cells associated
212 with different tumor phenotypes.³³ Pd, Ag, and Cu NPs were chosen as signal reporters since
213 they have well-separated potentials as redox-active probes. Biomarker-specific aptamers were
214 conjugated with these three types of metallic NPs to form Pd-anti-PSMA, Ag-anti-HER2, and
215 Cu-anti-MUC1 NPs. A mixture of these three NPs were successfully used for the specific
216 detection of different prostate and breast cancer cell lines such as VCaP, SK-BR-3, and MDA-
217 MB-231 (Figure 6).



218

219 **Figure 6** Linear-sweep voltammetry of specific cancer cell detection with a mixture of Pd-anti-
220 PSMA, Cu-anti-MUC1, and Ag-anti-HER2 nanoparticles: VCaP (blue), MDA-MB-231 (black)
221 and SK-BR-3 (red) cells. Reprinted with permission from ref. 33. Copyright 2014 WILEY-VCH
222 KGaA, Weinheim.

223

224 **3.4. DNAzyme-based sensing**

225 Deoxyribozymes, known as DNAzymes or catalytic DNAs, provide an alternative
226 approach to biosensing. DNAzymes are selected from random DNA sequences through
227 combinatorial screening techniques for catalytic and ligand-binding activities.³⁴ DNAzyme-
228 based sensing relies on the optical property of AuNPs for target recognition role for analytes
229 such as metal ions and small organic molecules. Using such DNAzyme-functionalized AuNPs,
230 the DNAzyme-catalyzed cleavage or ligation of the nucleic acid substrates affects the assembly
231 of AuNPs, resulting in a colorimetric readout for the cofactors.³⁴

232

233 DNAzymes can also be used for signal amplification by behaving as peroxidase mimics.
234 For example, it has been found that when one certain DNA sequence binds with hemin,
235 DNAzyme can be formed with G-quadruplex motifs. This type of DNAzyme can catalyze the
236 generation and enhancement of chemiluminescence (CL) signals in the presence of luminol and
237 H₂O₂. In this process, AuNPs are employed as carriers for these horseradish peroxidase (HRP)-
238 mimicking DNAzymes.³⁵ In a recent study, Zhang and coworkers applied HRP-mimicking
239 DNAzyme-functionalized NPs to cancer cell detection through the amplified CL signals.³⁶

240

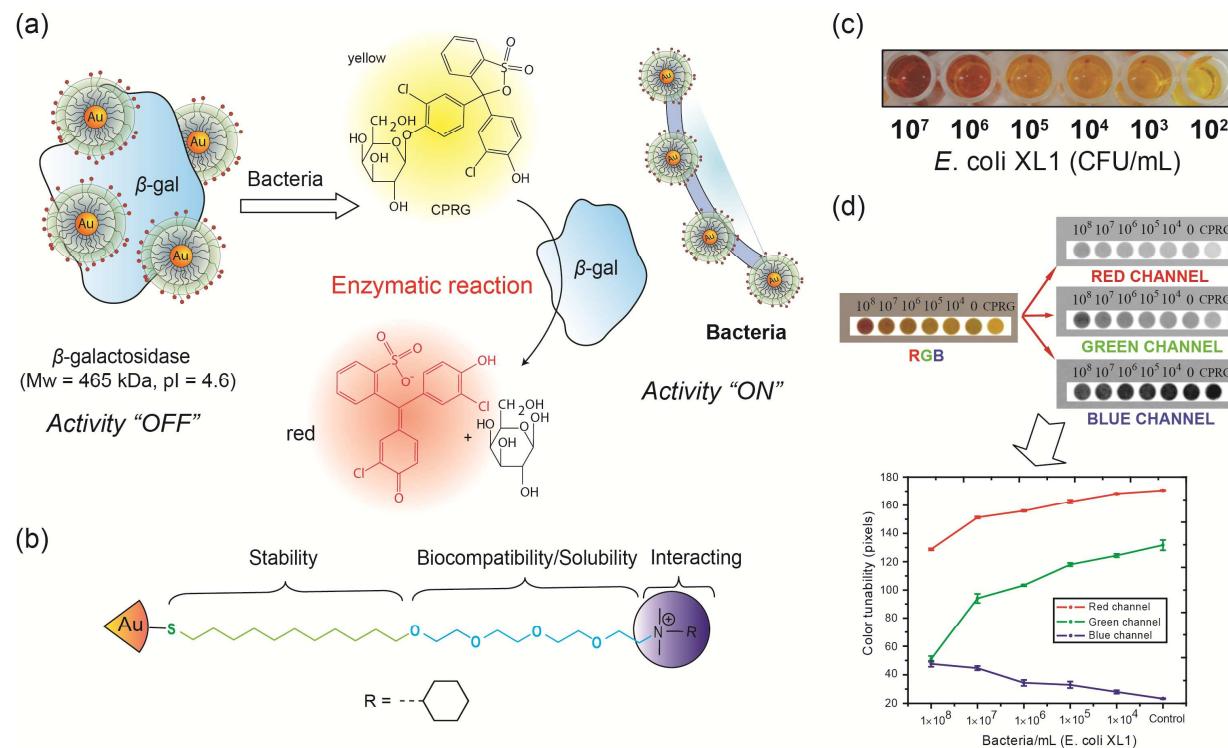
241 **4. Selective sensing**

242 Specific recognition-based sensors require pre-identification of the biomarkers, and face
243 certain limitations when used in systems containing multiple analytes. For example, cancer cells
244 present multiple biomarkers on the cell surface. The level of biomarkers may vary among cell
245 populations. In addition, subtle changes in the biomarker levels may be indicative of dramatic
246 phenotypic differences. As an alternative, sensors using *selectivity*-based modality do not require
247 the knowledge of a specific biomarker. On the contrary, selectivity-based approaches capture the
248 responses from complex analytes to generate a signature for each sample. Such selective sensing
249 approach can be utilized to detect non-specific analytes with either a single recognition element
250 or more commonly an array of recognition elements. In a typical array-based sensor, a set of
251 recognition elements interacts with a number of different analytes or classes of analytes,
252 providing a process reminiscent of mammalian olfaction.³⁷ This mechanistic similarity is why
253 array-based sensors are often denoted as chemical “noses” or “tongues”.

254

255 **4.1. Single recognition element system**

256 Rotello and coworkers have developed an enzyme amplification sensor using cationic
257 AuNP to inhibit the activity of β -galactosidase (β -gal) based on electrostatic interaction.³⁸ Such
258 enzyme catalysis can amplify the weak signals generated by the system. Bacterial cell surfaces
259 are negatively charged which can disrupt the AuNP- β -gal conjugates. During the sensing process,
260 bacteria cells replace β -gal from the NP- β -gal conjugates, restoring the activity of β -gal towards
261 the chromogenic substrate. Finally, the enzymatic reaction on the substrate gives the
262 corresponding readout to quantify the analytes (Figure 7).³⁹ The enzyme-amplified colorimetric
263 readout was able to detect 10^2 CFU/mL of *Escherichia coli* (*E. coli*) in solution. Furthermore, the
264 performance of this methodology was tested on a paper strip format against concentrations of
265 bacteria ranging from 10^4 ~ 10^8 CFU/mL. The designed bacteria test strips demonstrate the
266 potential for field applications such as a test of drinking water safety. However, this strategy
267 displays limitations in sensitivity and multiple analyte detection capability due to insufficient
268 interactions between the single recognition element and the analytes.



269

270 **Figure 7** (a) Schematic demonstration of enzyme amplified sensing of bacteria using gold
 271 nanoparticles. (b) The structure of quaternary amine functionalized gold nanoparticles. (c) The
 272 colorimetric sensing of *Escherichia coli* (*E. coli*) in solution. (d) Schematic illustration of the
 273 RGB analysis for monitoring color changes on test strips for different concentrations of *E. coli*.
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275

276 4.2. Array based sensing systems

277 Multiple recognition elements can maximize the variation in interactions between sensors
 278 and analytes. This array-based strategy combines responses from many individual sensors and
 279 analytes to generate a distinct pattern (fingerprint) for each analyte, either based on specific or
 280 selective interactions. Since multiple responses can be obtained from array-based sensors, these
 281 data matrices are generally analyzed using a variety of multivariate analyses such as principal
 282 component analysis (PCA) or linear discriminant analysis (LDA).⁴⁰

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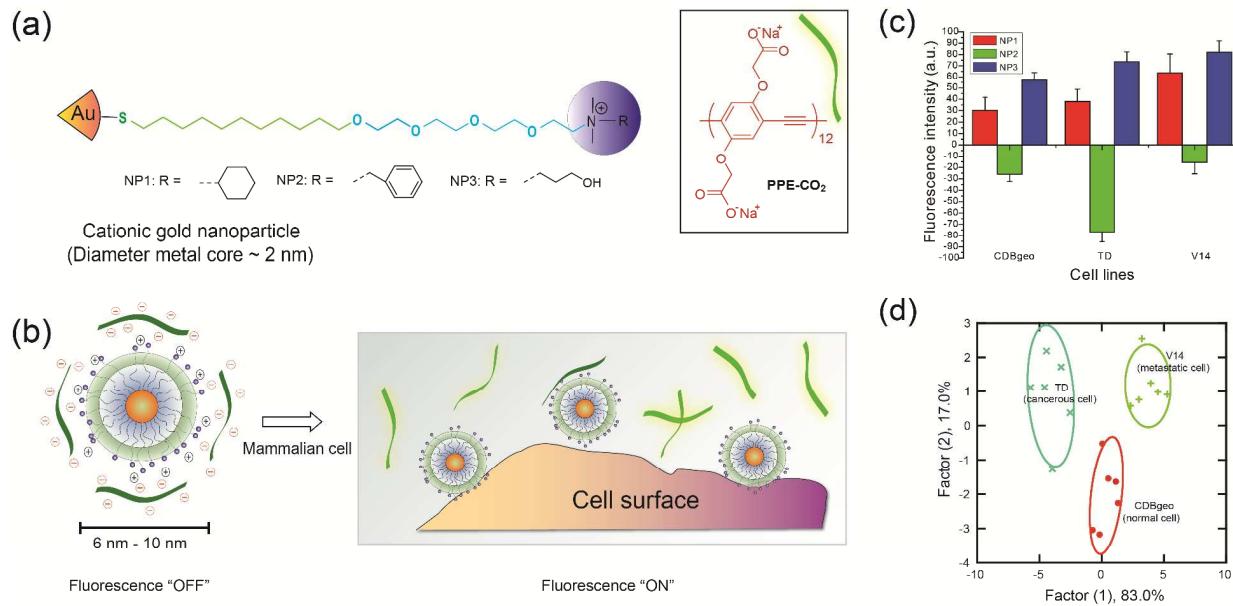
284 NPs can be readily functionalized with ligands to generate diverse sensor elements. These
 285 head groups exhibit differential affinity towards various analytes, leading to variations in the
 286 sensor that can be correlated to cellular signatures. Rotello and coworkers have designed several

287 small molecule based sensor arrays that used the phenomenon of fluorescence quenching.⁴¹
288 Fluorescence quenching is reduction in the fluorescence quantum yield due to energy transfer
289 from the photo-excited fluorophore to the AuNPs. In one approach, gold nanoparticles (AuNPs)
290 with a fluorescent polymer [carboxylate poly(para-phenyleneethynylene) (PPE-CO₂)] were used
291 to discriminate a series of cell lines (Figure 8).⁴² The sensor was comprised of three NP types
292 with different quaternary amine functional head groups. An array of AuNPs was used to quench
293 the intensity of the fluorescent polymers *via* electrostatic interaction. The subsequent binding of
294 cells disrupted the AuNP-polymer complex, thereby generating different fluorescence response
295 patterns for each cell lines. This AuNP-polymer complex was able to identify human cancerous
296 (MCF-7), metastatic (MDA-MB231) and normal (MCF10A) breast cell lines. Since these cell
297 lines came from different individuals, their differentiation might be originating from genetic
298 variation. To avoid this possibility, isogenic cell lines [CDBgeo (normal), TD (cancerous) and
299 V14 (metastatic)] derived from BALB/c mice were used to validate the sensor. Fluorescent
300 proteins can also be used as a transducer in an array based sensor. Cell differentiation using
301 AuNP-green fluorescent protein (GFP) conjugates resulted in four-fold enhancement in the
302 sensitivity of “chemical nose”-based sensor.⁴³ Moreover, in a subsequent study, AuNP-GFP
303 constructs were used to discriminate site specific metastases and healthy state using cell lysates
304 as well as tissue lysates, providing a promising strategy for medical diagnosis.⁴⁴

305

306

307



308

Figure 8 (a) Cationic gold nanoparticles (NP1-NP3) and the fluorescent polymer, carboxylate poly(para-phenyleneethynylene) (PPE-CO₂). (b) Fluorescence quenching of the polymers and the restoration of fluorescence after AuNP-polymer complex disrupted by the incubation with cells (dark green strips, fluorescence off; light green strips, fluorescence on). (c) Detection of three isogenic mammalian cell lines (CDBgeo, TD cell and V14) determined by fluorescence change using nanoparticle-polymer supramolecular complexes. (d) Canonical score plot using linear discrimination analysis (LDA) for the first two factors of simplified fluorescence response patterns obtained with NP-polymer assembly arrays against isogenic cell types. Reprinted with permission from ref. 42. Copyright 2009 National Academy of Sciences, USA.

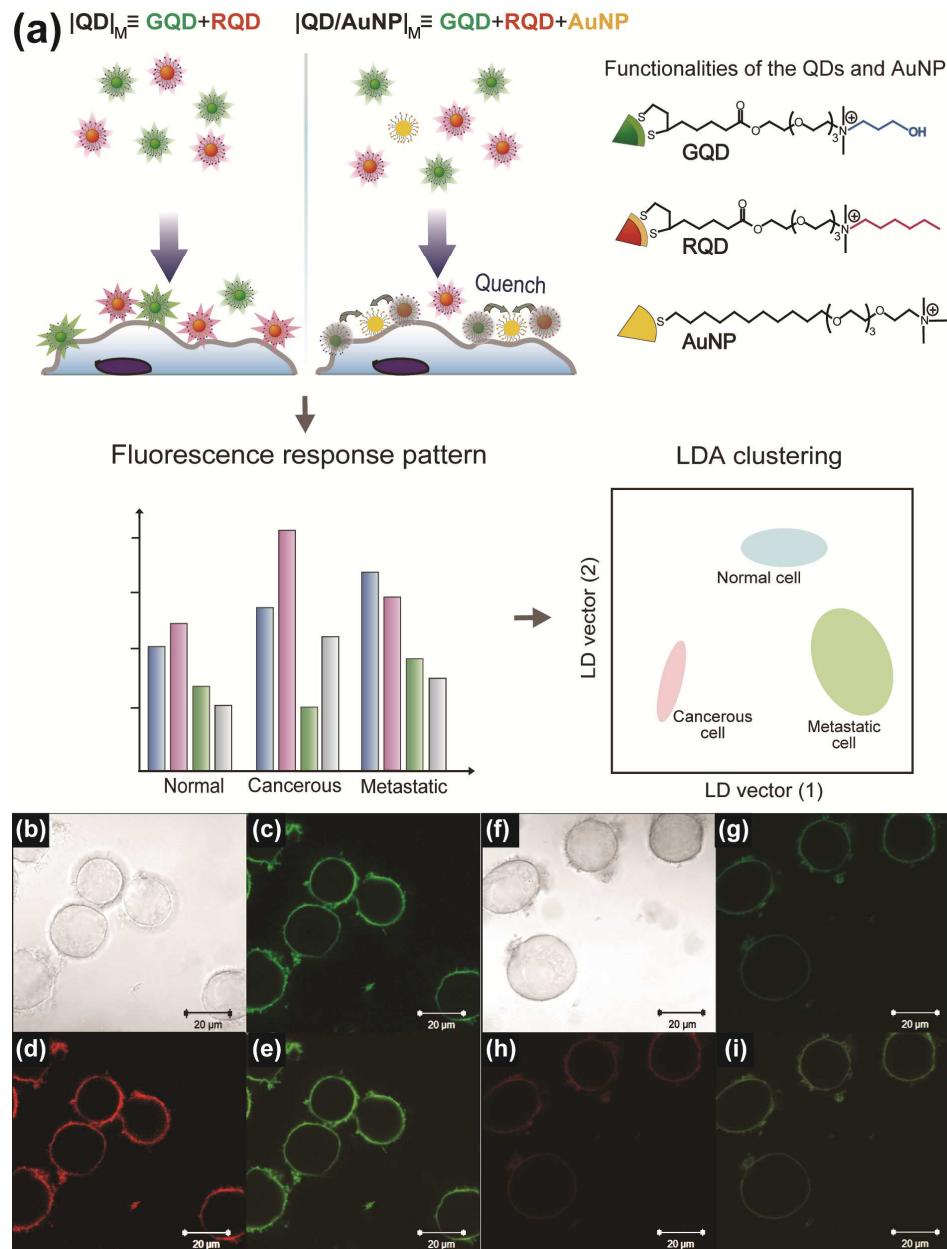
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In an analogous study, a swallowtail substituted carboxylate PPE (Sw-CO₂) was used as a transducer for AuNP-polymer sensor.⁴⁵ This array-based system comprised of amine functionalized hydrophobic NPs that served as recognition units for microorganisms such as bacteria. Bacterial cell walls are negatively charged and furnish a polyvalent environment to interact selectively with AuNP-polymer complex. For example, the Gram-positive microorganisms are highly negatively charged due to the presence of teichoic acid residue, whereas *E. coli* bacteria possess pili (rich in lectins) emanating from the surface. This array-based sensor enables the detection of bacteria cells within minutes. The AuNP-polymer complex was disrupted *via* competitive binding of different bacteria strains with the AuNP. Twelve

328 different bacteria strains including Gram-positive strains such as *Bacillus subtilis*, *Amycolatopsis*
329 *azurea* and Gram-negative bacteria such as *E. coli*, and *Pseudomonas putida* were identified.

330

331 Recently, quantum dots (QDs) were used in an array-based sensor as a recognition
332 element alongside AuNPs. When added together to the cells, co-localization of AuNPs and QDs
333 resulted in quenching, generating different patterns based on cell type/state. This sensor was used
334 to differentiate four different types of cancer cells as well as isogenic normal, cancer and
335 metastatic cells (Figure 9).⁴⁶ The dual channel fluorescence response obtained from the QD-
336 AuNP sensor array could identify 30 unknown samples with 100% accuracy. Besides pairing
337 with QDs, AuNPs can also be combined with upconversion nanoparticles (UCNPs) as a
338 fluorescence resonance energy transfer (FRET) couple to design biosensors.⁴⁷



339

340 **Figure 9** (a) Schematic illustration of the interaction between the nanoparticles and cell surface.
 341 The sensing system generated differential quenching and provided distinct patterns to discern
 342 different types/states of cells. Two arrays ($|QD|_M$ and $|QD/AuNP|_M$) were used in the system and
 343 placed in separated wells, with each array providing two fluorescence responses. $|QD|_M$, the
 344 mixture of GQD and RQD; $|QD/AuNP|_M$, the mixture of GQD, RQD, and AuNP. (b)-(i)
 345 Confocal microscopy images of (b)-(e) $|QD|_M$ and (f)-(i) $|QD/AuNP|_M$ after the incubation with
 346 HeLa cells for 15 min: (b), (f) bright field; (c), (g) green channel; (d), (h) red channel; (e), (i)
 347 merged images. Reprinted with permission from ref. 46. Copyright 2013 Elsevier.

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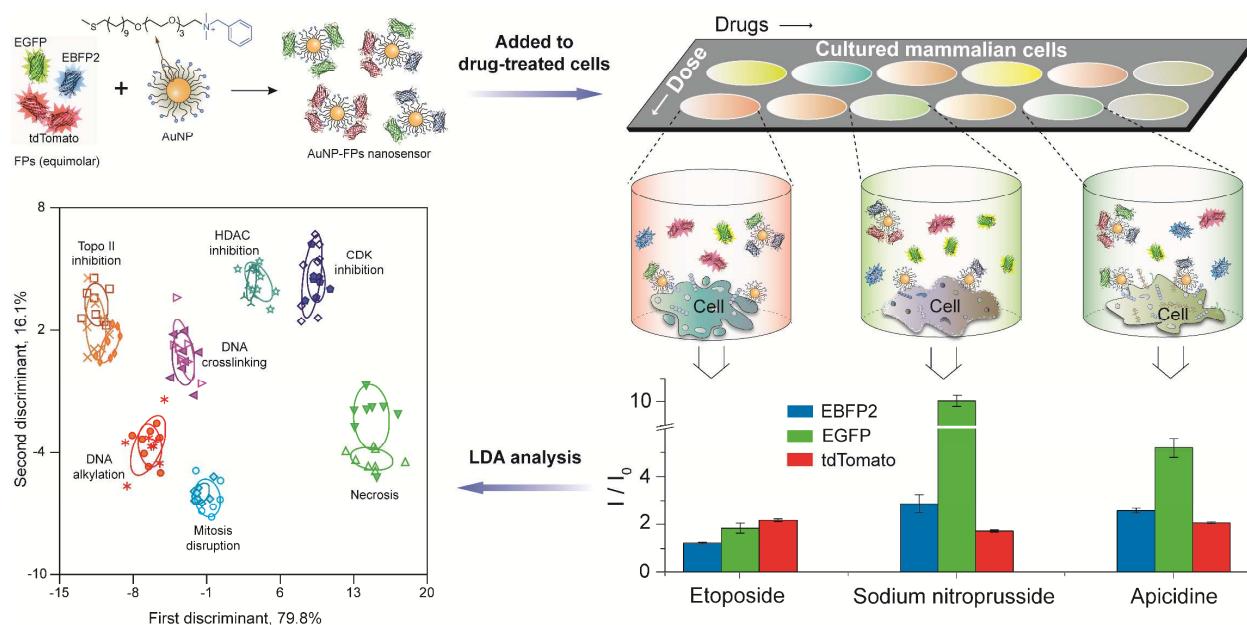
349 Functionalization of biomolecules such as aptamers on NPs can also be used for selective
350 identification of target analytes. Aptamers with selective binding properties towards were coated
351 on citrate capped AuNPs.⁴⁸ Upon addition of the target cells, aptamer-protected AuNPs
352 displayed different aggregation, generating different color patterns. Human cancer cells (Jurkat,
353 Reh, Raji) and normal human cells (WIL2-S) were distinguished with the array-based approach
354 by using one human immunoglobulin E aptamer (HIgE-1) and two thrombin aptamers (Tro-1
355 and Tro-2).

356

357 **4.3. Multiplexed output sensing system**

358 The previous array-based sensing examples use separate recognition elements to generate
359 the multiple sensor outputs required for identification of analytes. An alternative strategy to
360 generating information-rich data would be to use a single recognition element with multiple
361 outputs. Very recently, Rotello and coworkers have developed a high-throughput multi-channel
362 sensor that classifies the mechanism of chemotherapeutic drugs in minutes.⁴⁹ This sensor
363 consists of a single AuNP complexed with three different fluorescent proteins (FPs) that is used
364 to sense drug-induced physicochemical changes on cell surfaces. In the presence of cells,
365 differential displacement of the fluorophores with concomitant fluorogenesis provide a
366 ratiometric output that is measurable from a single readout (Figure 10). This result demonstrates
367 the ability of cell surface sensing to be used for high throughput screening of therapeutics, and
368 suggests the utility of these sensors for applications in toxicology and related fields.

369



370

Figure 10 Multi-channel sensor fabricated by incubating AuNP to an equimolar mixture of three fluorescent proteins (FPs): tdTomato (red), EBFP2 (blue) and EGFP (green). Different drug-treated cells result in distinct cell surface phenotypes, leading to different FP displacement patterns as schematically shown for the three wells. The bar plot shows differential fluorescence responses for three representative drugs. These fluorescence responses were further analyzed by linear discriminant analysis (LDA) to generate different clusters corresponding to different categories of drug mechanisms. Each ellipse represents each drug in that mechanism category.

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372

380 5. Conclusions and prospects

381 Metallic NPs present a versatile platform for the creation of recognition elements for
 382 analyzing the biological targets. NPs can be fabricated with different recognition elements to
 383 provide specific or selective interactions with the target analytes. Moreover, physicochemical
 384 properties of the NPs such as fluorescence quenching or enhancement, surface enhanced Raman
 385 scattering and electrochemical activity can be harnessed to signal the transduction of the binding
 386 events. Hence, inclusion of NPs can simplify the system design as well as increase the sensitivity
 387 of the biosensors.

388

389 Incorporation of metallic nanoparticles in diagnostic techniques has opened promising
 390 avenues for a wide range of sensing strategies that feature combinations of simplicity, rapid

391 output, low cost platforms and multiplexing. As we develop better strategies for particle
392 functionalization and signal transduction, a wide range of platforms ranging from microfluidic
393 sensors through inexpensive paper test strips will be enabled, provide solutions to address health
394 issues worldwide.

395

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401

402 References

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- 1 H. Shen and P. W. Laird, *Cell*, 2013, **153**, 38-55.
 - 2 M. Ferrari, *Nat. Rev. Cancer*, 2005, **5**, 161-171.
 - 3 L. Harris, H. Fritzsche, R. Mennel, L. Norton, P. Ravdin, S. Taube, M. R. Somerfield, D. F. Hayes and R. C. Bast, Jr., *J. Clin. Oncol.*, 2007, **25**, 5287-5312.
 - 4 M. Trzpis, P. M. J. McLaughlin, L. M. F. H. de Leij and M. C. Harmsen, *Am. J. Pathol.*, 2007, **171**, 386-395.
 - 5 P. K. Jain, X. H. Huang, I. H. El-Sayed and M. A. El-Sayed, *Acc. Chem. Res.*, 2008, **41**, 1578-1586.
 - 6 R. A. Sperling, P. Rivera Gil, F. Zhang, M. Zanella and W. J. Parak, *Chem. Soc. Rev.*, 2008, **37**, 1896-1908.
 - 7 E. Katz and I. Willner, *Angew. Chem. Int. Ed.*, 2004, **43**, 6042-6108.
 - 8 R. Mout, D. F. Moyano, S. Rana and V. M. Rotello, *Chem. Soc. Rev.*, 2012, **41**, 2539-2544.
 - 9 M. Edidin, *Nat. Rev. Mol. Cell Biol.*, 2003, **4**, 414-418.
 - 10 M. D. Mager, V. LaPointe and M. M. Stevens, *Nat. Chem.*, 2011, **3**, 582-589.
 - 11 D. H. Dube and C. R. Bertozzi, *Nat. Rev. Drug Discov.*, 2005, **4**, 477-488.
 - 12 K. Ohtsubo and J. D. Marth, *Cell*, 2006, **126**, 855-867.
 - 13 A. E. Nel, L. Maedler, D. Velegol, T. Xia, E. M. V. Hoek, P. Somasundaran, F. Klaessig, V. Castranova and M. Thompson, *Nat. Mater.*, 2009, **8**, 543-557.
 - 14 J. Sudimack and R. J. Lee, *Adv. Drug Del. Rev.*, 2000, **41**, 147-162.
 - 15 N. L. Rosi and C. A. Mirkin, *Chem. Rev.*, 2005, **105**, 1547-1562.
 - 16 N. Rifai, M. A. Gillette and S. A. Carr, *Nat. Biotechnol.*, 2006, **24**, 971-983.
 - 17 P. Holliger and P. J. Hudson, *Nat. Biotechnol.*, 2005, **23**, 1126-1136.
 - 18 M. D. Porter, R. J. Lipert, L. M. Siperko, G. Wang and R. Narayanan, *Chem. Soc. Rev.*, 2008, **37**, 1001-1011.
 - 19 M. Y. Sha, H. Xu, M. J. Natan and R. Cromer, *J. Am. Chem. Soc.*, 2008, **130**, 17214-17215.

-
- 20 X. Qian, X.-H. Peng, D. O. Ansari, Q. Yin-Goen, G. Z. Chen, D. M. Shin, L. Yang, A. N. Young, M. D. Wang and S. Nie, *Nat. Biotechnol.*, 2008, **26**, 83-90.
- 21 E. Katz, I. Willner and J. Wang, *Electroanalysis*, 2004, **16**, 19-44.
- 22 M. Maltez-da Costa, A. de la Escosura-Muniz, C. Nogues, L. Barrios, E. Ibanez and A. Merkoci, *Nano Lett.*, 2012, **12**, 4164-4171.
- 23 R. Jelinek and S. Kolusheva, *Chem. Rev.*, 2004, **104**, 5987-6016.
- 24 L. Ding, W. Cheng, X. Wang, S. Ding and H. Ju, *J. Am. Chem. Soc.*, 2008, **130**, 7224-7225.
- 25 L. Hu and G. Xu, *Chem. Soc. Rev.*, 2010, **39**, 3275-3304.
- 26 Y. Wang, Z. Chen, Y. Liu and J. Li, *Nanoscale*, 2013, **5**, 7349-7355.
- 27 X. Fang and W. Tan, *Acc. Chem. Res.*, 2010, **43**, 48-57.
- 28 S. Srivastava, B. L. Frankamp and V. M. Rotello, *Chem. Mater.*, 2005, **17**, 487-490.
- 29 J. Liu and Y. Lu, *Nat. Protoc.*, 2006, **1**, 246-252.
- 30 C. D. Medley, J. E. Smith, Z. Tang, Y. Wu, S. Bamrungsap and W. Tan, *Anal. Chem.*, 2008, **80**, 1067-1072.
- 31 H.-C. Yeh, J. Sharma, J. J. Han, J. S. Martinez and J. H. Werner, *Nano Lett.*, 2010, **10**, 3106-3110.
- 32 J. Yin, X. He, K. Wang, F. Xu, J. Shangguan, D. He and H. Shi, *Anal. Chem.*, 2013, **85**, 12011-12019.
- 33 Y. Wan, Y.-G. Zhou, M. Poudineh, T. S. Safaei, R. M. Mohamadi, E. H. Sargent and S. O. Kelley, *Angew. Chem. Int. Ed.*, 2014, **53**, 13145-13149.
- 34 J. Liu, Z. Cao and Y. Lu, *Chem. Rev.*, 2009, **109**, 1948-1998.
- 35 T. Niazov, V. Pavlov, Y. Xiao, R. Gill and I. Willner, *Nano Lett.*, 2004, **4**, 1683-1687.
- 36 S. Bi, J. Zhang and S. Zhang, *Chem. Commun.*, 2010, **46**, 5509-5511.
- 37 N. D. B. Le, M. Yazdani and V. M. Rotello, *Nanomedicine*, 2014, **9**, 1487-1498.
- 38 O. R. Miranda, H.-T. Chen, C.-C. You, D. E. Mortenson, X.-C. Yang, U. H. F. Bunz and V. M. Rotello, *J. Am. Chem. Soc.*, 2010, **132**, 5285-5289.
- 39 O. R. Miranda, X. Li, L. Garcia-Gonzalez, Z.-J. Zhu, B. Yan, U. H. F. Bunz and V. M. Rotello, *J. Am. Chem. Soc.*, 2011, **133**, 9650-9653.
- 40 A. M. Martinez and A. C. Kak, *IEEE Trans. Pattern Anal.*, 2001, **23**, 228-233.
- 41 U. H. F. Bunz and V. M. Rotello, *Angew. Chem. Int. Ed.*, 2010, **49**, 3268-3279.
- 42 A. Bajaj, O. R. Miranda, I.-B. Kim, R. L. Phillips, D. J. Jerry, U. H. F. Bunz and V. M. Rotello, *Proc. Natl. Acad. Sci. U. S. A.*, 2009, **106**, 10912-10916.
- 43 A. Bajaj, S. Rana, O. R. Miranda, J. C. Yawe, D. J. Jerry, U. H. F. Bunz and V. M. Rotello, *Chem. Sci.*, 2010, **1**, 134-138.
- 44 S. Rana, A. K. Singla, A. Bajaj, S. G. Elci, O. R. Miranda, R. Mout, B. Yan, F. R. Jirik and V. M. Rotello, *ACS Nano*, 2012, **6**, 8233-8240.
- 45 R. L. Phillips, O. R. Miranda, C.-C. You, V. M. Rotello and U. H. F. Bunz, *Angew. Chem. Int. Ed.*, 2008, **47**, 2590-2594.
- 46 Q. Liu, Y.-C. Yeh, S. Rana, Y. Jiang, L. Guo and V. M. Rotello, *Cancer Lett.*, 2013, **334**, 196-201.

-
- 47 L. Y. Wang, R. X. Yan, Z. Y. Hao, L. Wang, J. H. Zeng, J. Bao, X. Wang, Q. Peng and Y. D. Li, *Angew. Chem. Int. Ed.*, 2005, **44**, 6054-6057.
- 48 Y. Lu, Y. Liu, S. Zhang, S. Wang, S. Zhang and X. Zhang, *Anal. Chem.*, 2013, **85**, 6571-6574.
- 49 S. Rana, N. D. B. Le, R. Mout, K. Saha, G. Y. Tonga, R. E. S. Bain, O. R. Miranda, C. M. Rotello and V. M. Rotello, *Nat. Nanotechnol.*, 2015, **10**, 65-69.