

Rapid ratiometric biomarker detection with topically applied SERS nanoparticles

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Multiplexed surface-enhanced Raman scattering (SERS) nanoparticles (NPs) offer the potential for rapid molecular phenotyping of tissues, thereby enabling accurate disease detection as well as patient stratification to guide personalized therapies or to monitor treatment outcomes. The clinical success of molecular diagnostics based on SERS NPs would be facilitated by the ability to accurately identify tissue biomarkers under time-constrained staining and detection conditions with a portable device. *In vitro*, *ex vivo* and *in vivo* experiments were performed to optimize the technology and protocols for the rapid detection (0.1-s integration time) of multiple cell-surface biomarkers with a miniature fiber-optic spectral-detection probe following a brief (5 min) topical application of SERS NPs on tissues. Furthermore, we demonstrate that the simultaneous detection and ratiometric quantification of targeted and nontargeted NPs allows for an unambiguous assessment of molecular expression that is insensitive to nonspecific variations in NP concentrations.

INNOVATION

Multidisciplinary advances have been made to develop a technology for the multiplexed molecular phenotyping of fresh *ex vivo* and *in vivo* tissues under time-constrained conditions that are relevant for point-of-care clinical applications. By developing high-affinity targeted SERS NPs, a sensitive portable spectral-detection device, and an optimized topical-delivery protocol, we demonstrate for the first time a ratiometric method to rapidly quantify the specific binding of a panel of biomarker-targeted NPs on fresh tissues, thereby eliminating the ambiguities that often arise due to nonspecific sources of contrast.

INTRODUCTION

A major focus of biomedical optics has been to develop technologies for the detection of some of the most prevalent diseases worldwide such as epithelial cancers of the colon, esophagus, oral cavity, cervix and skin, as well as to image surgical margins to guide tumor-resection procedures^{1–6}. The general approach of optical diagnostics is to deduce tissue status through the measurement of optical signals generated either intrinsically by cell and tissue constituents^{7,8} or extrinsically by targeted contrast agents with known signatures^{9–11}. While the simplicity and regulatory ease of imaging intrinsic signatures is compelling, the use of exogenous contrast agents allows for the assessment of highly informative biomarkers such as cell-surface receptors. A challenging issue is that molecular biomarkers of disease vary greatly among subjects, between disease subtypes, and even within a single subject over time¹². Thus, exogenous probes should ideally be capable of being multiplexed to simultaneously detect multiple biomarkers. A technology for the rapid molecular phenotyping of fresh tissues at the point of care could enable accurate disease diagnosis, the

monitoring of treatment response, and patient stratification to guide personalized therapies.

Although numerous molecular probes are being developed to label disease biomarkers^{13–15}, their utility for cancer detection is often limited by various factors. For example, fluorescent dyes are easily photobleached, have a wide emission spectrum, and must often be excited at disparate wavelengths when combined, thus limiting their multiplexing capability. Although quantum dots (QD) offer a narrower emission bandwidth, higher sensitivity and higher photostability than fluorescent dyes¹⁶, their potential toxicity has thus far precluded their *in vivo* use in humans¹⁷. Surface-enhanced Raman-scattering (SERS) nanoparticles (NPs), hereafter referred to as “SERS NPs” or “NPs”, have attracted interest due to their brightness, low toxicity, and potential for sensitive and multiplexed biomarker detection¹⁸. The SERS NPs utilized in this study are available in multiple “flavors,” each of which emits a characteristic Raman fingerprint spectrum that allows for the identification and quantification of large multiplexed mixtures of different NP flavors when illuminated at a single wavelength^{19–21}. It is important to emphasize that these SERS NPs are engineered to emit a stable and unique Raman spectra that is insensitive to the environment¹⁹. This is accomplished by encapsulating the SERS NPs within a protective silica shell, such that their gold core and Raman-active layer are shielded from other NP cores as well as from their surroundings. The gold cores at the center of these SERS NPs provide an electromagnetic enhancement that dramatically increases the Raman fingerprint signal (which uniquely identifies each NP flavor) compared with non-enhanced Raman signals^{19,22}. Since the Raman signals emitted by these SERS NPs are much brighter than background Raman signals from tissue components or buffers, the background Raman signals are negligible at the measurement conditions (laser power, detector

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integration times, and optical set up) utilized in this study. Here we design each flavor of NP to target a unique protein biomarker by conjugating the NPs to monoclonal antibodies. The SERS NP spectra can be easily identified by their distinct and narrow spectral peaks whereby the quantity of each NP flavor in a mixture may be determined through a spectral-demultiplexing software algorithm^{21,23–25}.

In recent years, a number of groups have begun to investigate the advantages of various types of SERS NPs for the detection of cancer biomarkers^{18,26}. For example, a few studies have explored the feasibility of using SERS NPs to specifically label cell-surface protein biomarkers under *in vitro* conditions^{9,27–31}. One study has demonstrated the basic feasibility of targeting a single biomarker using SERS-NPs on *ex vivo* tissues topically stained for 1 hour and detected with a large Raman microscope system⁹. In addition, a few groups have demonstrated the feasibility of the multiplexed detection of large panels of nontargeted SERS NPs^{21,22,32–36}. Finally, studies have utilized large Raman microscope systems to explore the behavior of targeted and nontargeted SERS NPs after intravenous injection in tumor-bearing mice^{26,37}.

Although the basic sensitivity, targeting ability and multiplexing capabilities of SERS NPs have been investigated, there is a need to demonstrate that multiplexed molecular detection is possible under time-limited tissue-staining and detection conditions, as encountered in point-of-care clinical settings, using a miniaturized spectral-detection probe. In particular, in this study we focus on establishing the feasibility of topical application as a means of rapidly delivering targeted SERS NPs to exposed tissue surfaces (e.g. epithelial surfaces and surgical margins), resulting in sufficient contrast to enable *in vivo* molecular detection at short integration times (0.1 s). Topical application potentially offers an expedited route for the regulatory approval of SERS NPs, especially since recent studies have shown that SERS NPs applied topically in the gastrointestinal tract result in negligible systemic uptake^{38,39}. An example application for this technology would be to screen the oral cavity, esophagus, or colon for dysplasia and/or cancer by irrigating the tissue with, or having the patient drink, a cocktail of targeted SERS NPs prior to performing multiplexed molecular imaging with a spectral-imaging endoscope. In the case of oral cancer detection, a major clinical challenge is the fact that many lesions appear suspicious upon visual examination but are actually noncancerous. Physicians (and dentists) are often hesitant to obtain physical biopsy specimens due to the discomfort and potential complications from such biopsies that are often of benign lesions⁴⁰. A noninvasive imaging method to screen for molecular alterations indicative of cancer or precancer could play a valuable role in guiding biopsies (*i.e.* image-guided biopsy). Another example would be the intraoperative molecular phenotyping/imaging of *in vivo* or freshly excised surgical specimens to identify residual tumor at the surgical margins and to improve the completeness of resection¹⁵. In the specific case of breast-conserving surgery (lumpectomy), numerous reports reveal that 20% to 60% of lumpectomy patients must undergo re-excision surgeries after post-operative pathology reveals residual tumor at the surgical margins^{41,42}. With our technology, tissues could potentially be topically stained with a panel of SERS NPs (*in vivo* or *ex vivo*), and imaged after one or more rinse steps to rapidly identify/quantify the presence of tumor-related biomarkers during surgery. In this study, we demonstrate the general feasibility to address both *ex vivo* and *in vivo* imaging applications.

In this study, we also demonstrate the advantages that SERS NPs confer for the ratiometric quantification of molecularly specific vs. nonspecific binding. The importance of distinguishing between specific and nonspecific probe accumulation is critical. For example, in this study we observe that nontargeted NPs that are topically applied on tissues show higher retention in normal tissues versus xenograft tumors, *i.e.* inverse contrast. This same phenomenon has also been seen with intravenously injected contrast agents (e.g. fluorescent dyes) in implanted tumors^{43,44}. By contrast, in many animal models and clinical samples, the opposite effect of enhanced penetration and retention (EPR) in tumors versus

normal tissues is often observed^{10,45,46}. Regardless of the mechanisms for the nonspecific behavior of contrast agents, there is increasing consensus that utilizing a dual-reporter ratiometric detection strategy, in which one contrast agent serves as a negative control for a simultaneously delivered targeted probe, is valuable to accurately identify and quantify molecularly specific binding^{43–45}. Here we show that multiplexed SERS NPs are well suited for this purpose, owing to the fact that one nontargeted NP may serve as a negative control to quantify the specific vs. nonspecific binding of several other targeted NPs. In particular, our SERS NPs are excited under identical illumination conditions from a single laser (785 nm), and emit Raman spectra within the same narrow band of wavelengths (approximately 850 – 900 nm). Therefore, their ratios are insensitive to variables such as tissue optical properties [see **Supplementary Information (SI)** for additional information]. In short, we stain tissues with an equimolar mixture of targeted NPs and one untargeted NP that controls for nonspecific effects. By first obtaining a calibration measurement of an equimolar mixture of NP flavors, it is possible to accurately quantify the relative concentration ratio of the NP flavors when the mixture is used to stain tissues (see **SI**), and consequently to determine the specific vs. nonspecific binding ratio of the NPs. This assumes that the nontargeted NP flavor accurately mimics the nonspecific behavior of the targeted NP flavors in the mixture (see **Supplementary Fig. 3** for information on the optimal design of a nontargeted control NP). We demonstrate the robustness of this ratiometric measurement approach *via in vitro* and *ex vivo*, as well as *in vivo* experiments.

RESULTS

Specific-binding ability of mAb-conjugated NPs

Flow cytometry and confocal microscopy were first performed to investigate the specific-binding abilities of targeted vs. nontargeted SERS NPs. As described in the Methods section, SERS NPs were conjugated with fluorophores (for flow cytometry) as well as with one of the following three monoclonal antibodies (mAbs): an anti-EGFR mAb (to NP flavor S440), an anti-HER2 mAb (to NP flavor S420), or an isotype-control (nontargeted) mAb (to NP flavor S421). Conjugation parameters (e.g. dosage of antibody) and staining conditions (e.g. NP concentration and staining duration) were all optimized to maximize the ratio of specific vs. nonspecific binding of the targeted NPs (**Supplementary Table 1**). In addition, the design of a nontargeted NP to control for nonspecific binding was also optimized (**Supplementary Fig. 3**). For cell labelling, a 200-pM staining concentration and a 15-min staining duration resulted in high levels of specific vs. nonspecific binding. This study utilized three cell lines: (1) A431, which highly overexpresses the epidermal growth factor receptor (EGFR) with $\sim 2 \times 10^6$ receptors/cell and moderately expresses the human epidermal growth factor receptor 2 (HER2) with $\sim 2 \times 10^4$ receptors/cell^{47,48}; (2) SkBr3, which highly overexpresses HER2 with $\sim 1 \times 10^6$ receptors/cell and moderately expresses EGFR with 9×10^4 receptors/cell^{47,48}; and (3) 3T3, a normal mouse fibroblast cell line that expresses negligible amounts of EGFR and HER2⁴⁹. The density of overexpressed cell-surface receptors in the A431 and SkBr3 cell lines is representative of the receptor densities measured in many naturally occurring tumors⁵⁰.

Figure 1 shows the flow cytometry data in the form of fluorescence histograms of a 10,000-cell analysis. The geometric mean of the fluorescent intensity (MFI) of the cell samples is used to compare binding levels. For the negative-control cell line 3T3, the MFI of cells stained with all three NP conjugates is similar. For the A431 cell line, which highly overexpresses EGFR, the MFI of cells stained with EGFR-NPs is 65 times higher than cells stained with isotype-NPs. Since A431 cells express moderate levels of the HER2 receptor^{47,51}, cells stained with HER2-NPs also show a slightly elevated MFI compared to cells stained with isotype-NPs (~ 5 times). SkBr3 cells have been reported to highly express HER2 as well as EGFR at a slightly lower level^{47,51}. Our data in

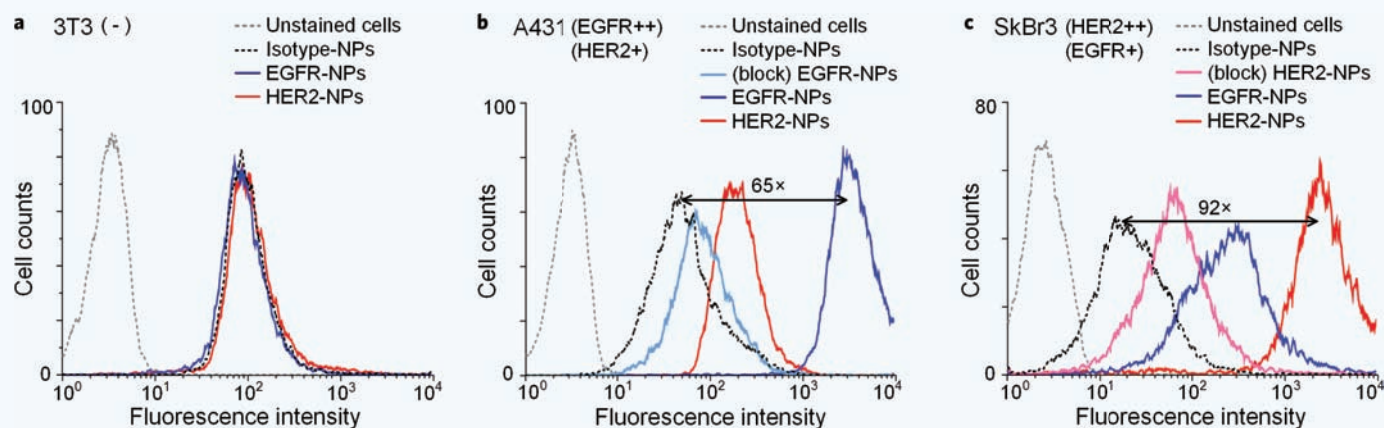


Figure 1 Flow cytometry validation of conjugated NPs with cultured cells. NPs were conjugated with either anti-EGFR, anti-HER2 or isotype control monoclonal antibodies. These NPs were individually used to stain (a) 3T3 (-), (b) A431 (EGFR++, HER2+) and (c) SkBr3 (HER2++, EGFR+) cell lines for 15 min, followed by flow cytometry analysis. Fluorescence histograms from unlabeled cells as well as cells stained with isotype-NPs, EGFR-NPs and HER2-NPs are shown. In addition, the results of competitive-binding experiments are displayed, in which excess mAb is used to block cell-surface receptors prior to NP staining (see text for details).

Fig. 1c support these reports, showing that cells stained with HER2-NPs have a MFI that is 92 times higher than cells stained with isotype-NPs, and that cells stained with EGFR-NPs have a MFI that is 15 times higher than cells stained with isotype-NPs. When competitively inhibited with an excess of mAb, A431 cells (inhibited with anti-EGFR mAb and then labeled with EGFR-NPs) show a 26-fold reduction in MFI, while SkBr3 cells (inhibited with anti-HER2 mAb and then labeled with HER2-NPs) show a 17-fold reduction, suggesting that the EGFR- and HER2-conjugated NPs bind specifically to cell-surface EGFR and HER2 receptors, respectively.

In order to further confirm the specific-binding ability of the NP conjugates, cells from flow-cytometry experiments were transferred to glass slides and observed under a confocal microscope (Leica TCS SP5). By setting a tight detection pinhole to produce a thin optical section, it was observed that the fluorescence signal was primarily localized to the periphery of the cells (Supplementary Fig. 4), indicating that the NPs were bound to the cell surface rather than internalized. To compare the total amount of NPs bound to each cell (Supplementary Fig. 5), the pinhole was enlarged. The same objective (20 \times ; dry), laser power, and detector gain settings were used to image all samples. Supplementary Fig. 5 shows the bright-field and fluorescence images of 9 stained samples, including all combinations of 3 cell lines and 3 types of conjugated NPs.

Ratiometric analysis of multiplexed NPs on cell monolayers

In vitro experiments were performed to demonstrate the ability to rapidly assess molecular expression on cell monolayers *via* multiplexed detection of SERS NPs with a miniature spectral-detection probe (Fig. 2). To simulate the topical application of NPs on tissue surfaces, confluent monolayers of cells were cultured in 96-well plates (see Methods and Supplementary Fig. 7–9). Equimolar mixtures of positive and negative NPs were simultaneously applied to the monolayers for 15 min. Unbound NPs were then removed *via* multiple rounds of rinsing with phosphate-buffered saline (PBS). SERS spectra

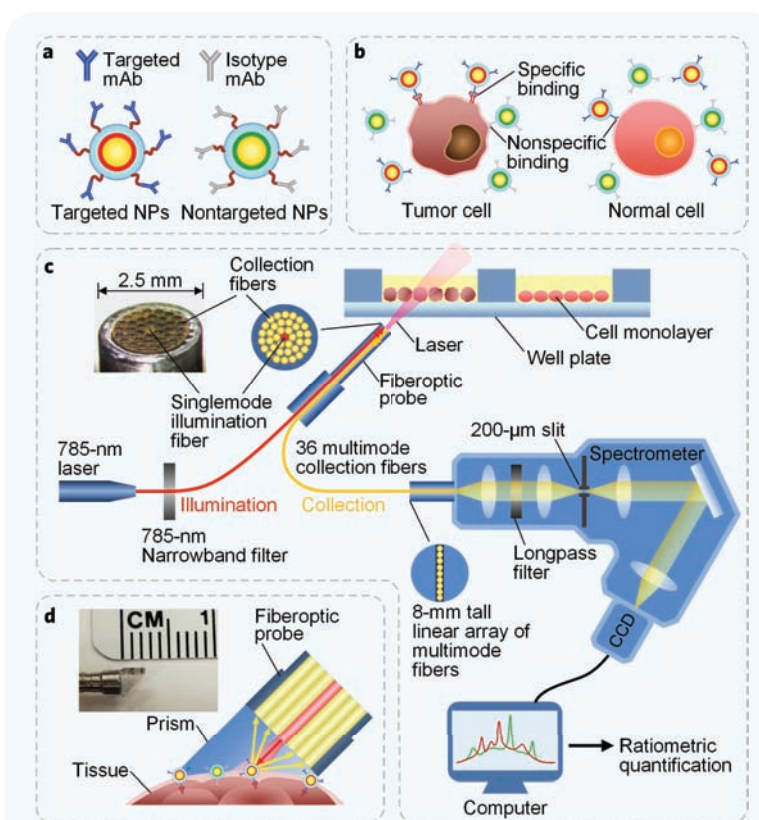


Figure 2 Schematic illustrations of a customized spectral-imaging device to detect bioconjugated NPs on cells and tissues. (a) NPs conjugated with targeted- and isotype-control (nontargeted) monoclonal antibodies (mAb), where a unique flavor of NP is used for each of the targeted and nontargeted contrast agents; (b) A depiction (not to scale) of the specific binding of multiplexed NPs to cell-surface receptors (in the case of targeted NPs) or nonspecific locations (in the case of all NP flavors); (c) Experimental platform to detect multiplexed NPs on cell monolayers; (d) Angled contact probe for *ex vivo* and *in vivo* tissue measurements.

were collected with a miniature Raman detection probe, in which the NP mixtures were illuminated with a single low-power (10 mW) diode laser (see Fig. 2c, Methods, and design optimization in Supplementary Fig. 1). Several approaches have been developed to demultiplex (unmix) a spectrum generated by two or more flavors of SERS NPs in the presence of a competing background^{21,24}. An analysis and comparison between various demultiplexing approaches is beyond the scope of this paper but has been explored by others⁵². Here, a simple and commonly used direct classical least squares (DCLS) demultiplexing method was employed²³, along with a calibrated ratiometric strategy to quantify specific vs. nonspecific binding (see SI). We have shown through validation studies (Supplementary Fig. 6) that the DCLS demultiplexing approach provides accurate quantification of NP concentrations and ratios across our measurement conditions.

In initial experiments, a dual-flavor mixture (150 pM per flavor) consisting of a targeted (positive) NP flavor and a nontargeted (negative) NP flavor, was applied to each well to stain cell monolayers for 15 min, followed by 3 rounds of rinsing to remove unbound NPs (each rinse replaced ~90% of the liquid in each well). Raman spectra from an A431 monolayer stained with NPs, before and after rinsing, are shown in Fig. 3a. The Raman peaks are distinguishable in spite of a competing broadband spectral background, even after multiple stages of rinsing. For each cell line, raw spectra were analyzed (3 wells of each cell line, 3 different positions per well, and 2 acquisitions at each position for a total $n = 18$) to calculate the weight of each flavor, the weight ratio of positive and negative NPs, and the actual concentration ratio of the NPs (+NPs/-NPs). The weight ratios measured before rinsing were used to calibrate subsequent ratiometric measurements, assuming that a 1:1 equimolar ratio of NPs was used in the staining mixtures. The calibrated NP concentration ratios are plotted in Fig. 3b,c (additional details are provided in the SI).

For A431 cell monolayers, EGFR-NPs and isotype-NPs were used for initial 2-flavor staining experiments. As shown in Fig. 3b, after three rinses the concentration ratio (+NPs/-NPs) increases to 2.4 for A431 monolayers but remains near unity for 3T3 monolayers. For SkBr3 cell monolayers (Fig. 3c), the concentration ratio of HER2-NPs/isotype-NPs increases to 2.6 after three rinses, while remaining near unity for 3T3 cell monolayers. After three rinsing steps, the approximate concentration of the isotype-NPs decreases to 4–6 pM. Since this is near the limit of detection (LOD) of our spectral-detection probe (see Supplementary Fig. 6), data with further rinsing steps are not shown.

In vitro experiments with three multiplexed NP flavors (two positive and one negative control) were carried out to demonstrate the ability to simultaneously assess the expression of multiple cell-surface biomarkers (EGFR and HER2). Here, a mixture of EGFR-NPs, HER2-NPs and isotype-NPs (150 pM per flavor) were applied to stain cell monolayers, followed by 3 rounds of rinsing. Figure 3d–f show the concentration ratios between positive and negative NPs (both the EGFR/isotype ratio and the HER2/isotype ratio) on the different cell monolayers ($n = 18$ as previously described). These results are consistent with the two-particle staining experiments and demonstrate the utility of ratiometric analyses of multiplexed NPs to rapidly identify the expression of one or more cell-surface proteins.

Ratiometric analysis of multiplexed NPs on tumor explants (*ex vivo*)

Ex vivo experiments were performed to demonstrate the ability to detect biomarker expression in fresh tissues with topically applied SERS NPs and rapid detection with a miniature probe (see Fig. 2d, Methods, and design optimization in Supplementary Fig. 1). A431 and SkBr3 tumor xenografts (5–10 mm diameter) as well as normal tissues (Fig. 4a) were

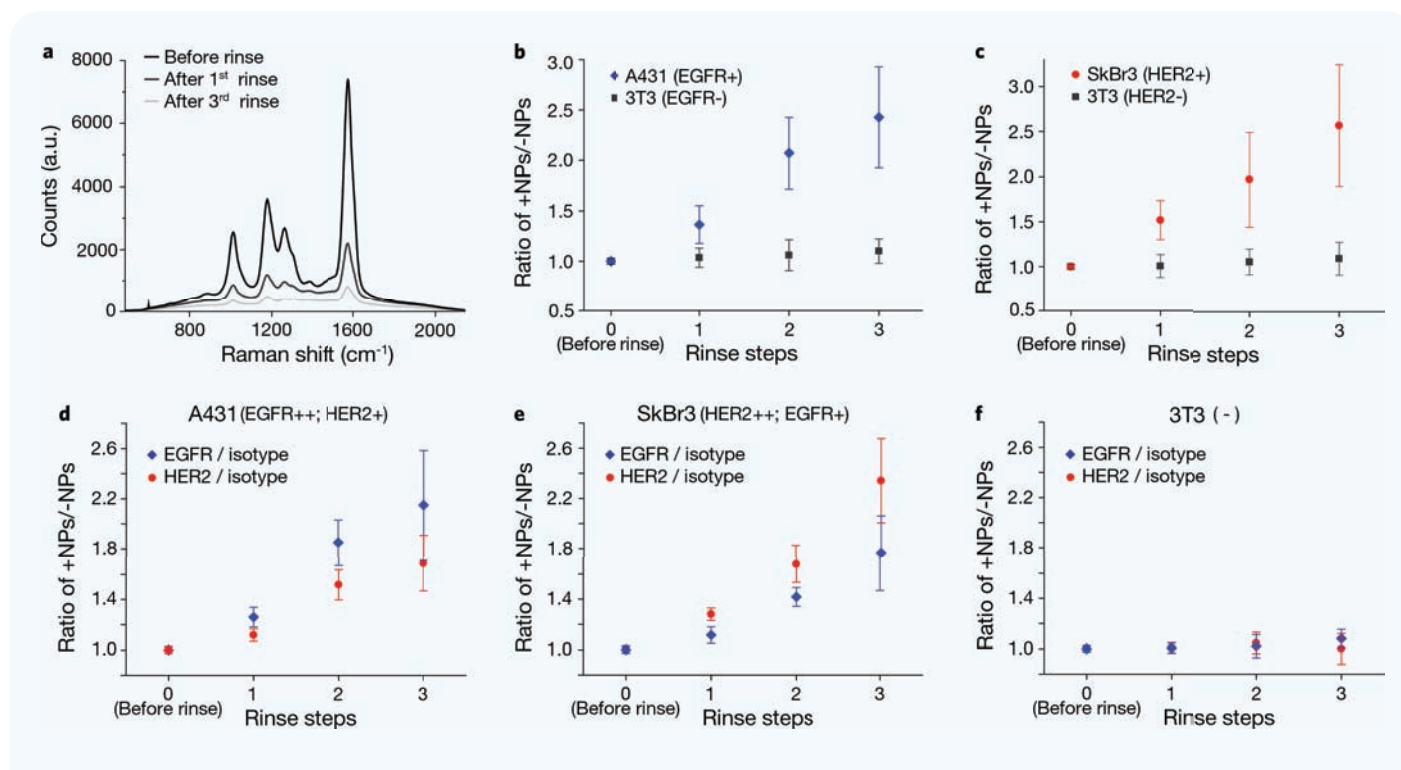


Figure 3 Demultiplexing analysis of the concentration ratio of multiplexed NPs on cell monolayers. Each cell monolayer was simultaneously stained with two (a–c) or three (d–f) flavors of NPs and then rinsed with PBS. (a) Raw spectra before and after rinses (from an A431 cell monolayer); (b) Concentration ratios of EGFR-NPs/isotype-NPs on A431 and 3T3 cells; (c) Concentration ratios of HER2-NPs/isotype-NPs on SkBr3 and 3T3 cells; concentration ratios of EGFR-NPs/isotype-NPs and HER2-NPs/isotype-NPs on (d) A431, (e) SkBr3 and (f) 3T3 cell monolayers.

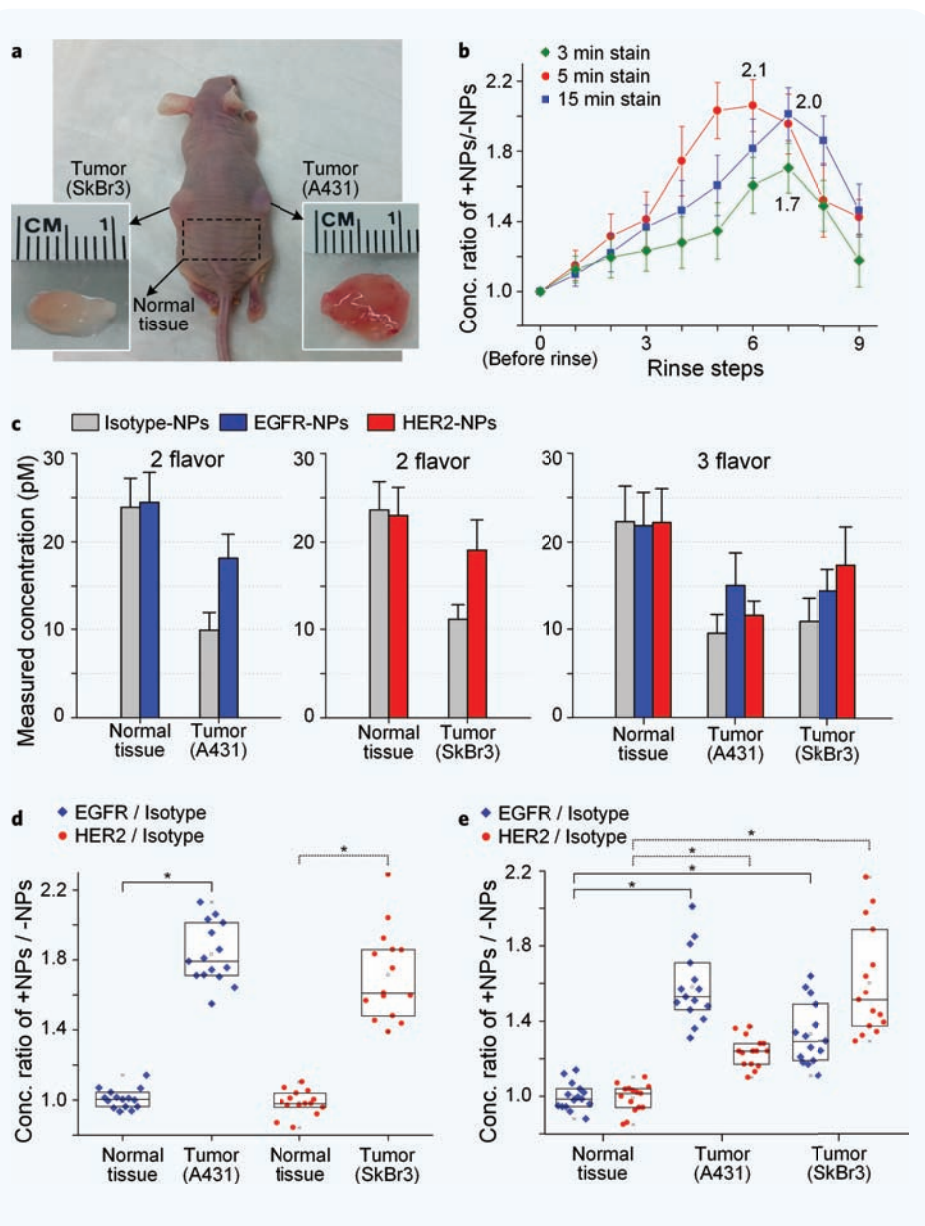


Figure 4 *Ex vivo* experiments with xenograft tumors. (a) Nude mouse with xenograft tumors 14 d after subcutaneous injection of 1×10^6 A431 cells and 5×10^6 SkBr3 cells (the lower insets show the explanted tissues). (b) Experiment to optimize the staining of tissues with SERS NPs showing the concentration ratio of a targeted vs. nontargeted NP (EGFR-NP/isotype-NP) after each rinse step (3 s per rinse) following either a 3-, 5-, or 15-min staining duration. (c) Measured absolute concentrations of NPs on tumor and normal tissues after 18 s of rinsing (equivalent to 6 rinse steps) for 2-flavor and 3-flavor multiplexed experiments, showing elevated nonspecific accumulation in normal tissues vs. tumors. However, plotting the concentration ratio with (d) 2-flavor and (e) 3-flavor multiplexing ($n = 15$) allows for the unambiguous quantification of specific vs. nonspecific binding. * P -value < 0.001 .

resected and cut into similarly sized pieces (30–60 mm³) for the measurement of NP concentrations and ratios.

The staining and rinsing protocols were first optimized to maximize the concentration ratio of +NPs/-NP. For optimization experiments, A431 tumor specimens were immersed into a 50- μ L equimolar mixture of EGFR-NPs and isotype-NPs (200 pM per flavor), and allowed to incubate without agitation for 3, 5 or 15 min (3 pieces per staining condition). The tissues were then washed 9 times by submerging them into 20 mL of PBS (with gentle agitation) for 3 s per wash. SERS spectra were collected with the contact probe (integration time: 0.25 s) after each rinse step. Note that before staining, a 4- μ L droplet of an equimolar NP mixture was placed on

the contact surface of the probe to calibrate subsequent concentration and ratiometric measurements (see SI). **Figure 4b** plots the calibrated NP concentration ratio after each rinse ($n = 6$). A 5-min staining duration resulted in the highest NP concentration ratio with the least number of rinse steps and was therefore used for subsequent *ex vivo* and *in vivo* experiments.

Results from *ex vivo* experiments with two multiplexed NP flavors (1 targeted NP and 1 nontargeted control NP) or three multiplexed NP flavors (2 targeted NPs and 1 nontargeted control NP) are shown in **Fig. 4c–e**. Under identical staining (5 min) and washing conditions (18 s), stronger Raman signals were detected from normal tissues than from tumor tissues (**Fig. 4c**), indicating that there is enhanced nonspecific retention of topically applied NPs in normal tissues vs. the tumor xenografts. However, **Fig. 4d,e** show that the concentration ratio between positive and negative NPs (EGFR/isotype and HER2/isotype) on tumors was significantly elevated, indicating preferential binding of targeted NPs to overexpressed cell-surface biomarkers. In normal tissues, the NP ratios remained near unity due to the negligible expression of EGFR and HER2 in these tissues.

Rapid ratiometric analysis of multiplexed NPs on tumor implants (*in vivo*)

In vivo experiments were performed with a 0.1-s detector integration time (see **Supplementary Fig. 10** for a noise analysis) using a flexible version of the spectral-detection device (**Fig. 5a**) to demonstrate rapid ratiometric quantification of NPs in live mice. **Figure 5b** shows the reference spectra of the SERS NPs and tissue background (no NPs) that were acquired for these *in vivo* experiments. **Figure 5c** provides example spectra to illustrate the DCLS demultiplexing algorithm, including a raw spectrum from a 3-flavor mixture of NPs on tumor tissues acquired with a 0.1-s integration time, a best-fit curve using the DCLS algorithm, the combined spectrum of the NPs after removing the tissue background component, and the demultiplexed spectra from individual NP flavors (blue: EGFR-S440, 17.9 pM; red:

HER2-S420, 13.1 pM; gray: isotype-S421, 10.4 pM). **Figure 5d,e** show the concentration ratio between positive and negative NPs (EGFR/isotype and HER2/isotype) on the tumors and normal tissues of 5 mice ($n = 15$). The results are similar to the *ex vivo* results: the ratios on normal tissues remain near unity, but are significantly elevated on tumors. Interestingly, the measurement uncertainty for the *in vivo* results is less than for the *ex vivo* results, possibly due to reduced mechanical perturbation and measurement artifact when probing *in vivo* tissues compared with surgically excised tissues. In order to demonstrate that our demultiplexing algorithm and results are insensitive to any specific choice of NP flavor, we altered the combination of NP flavors that were conjugated to each of the three mAbs (isotype control,

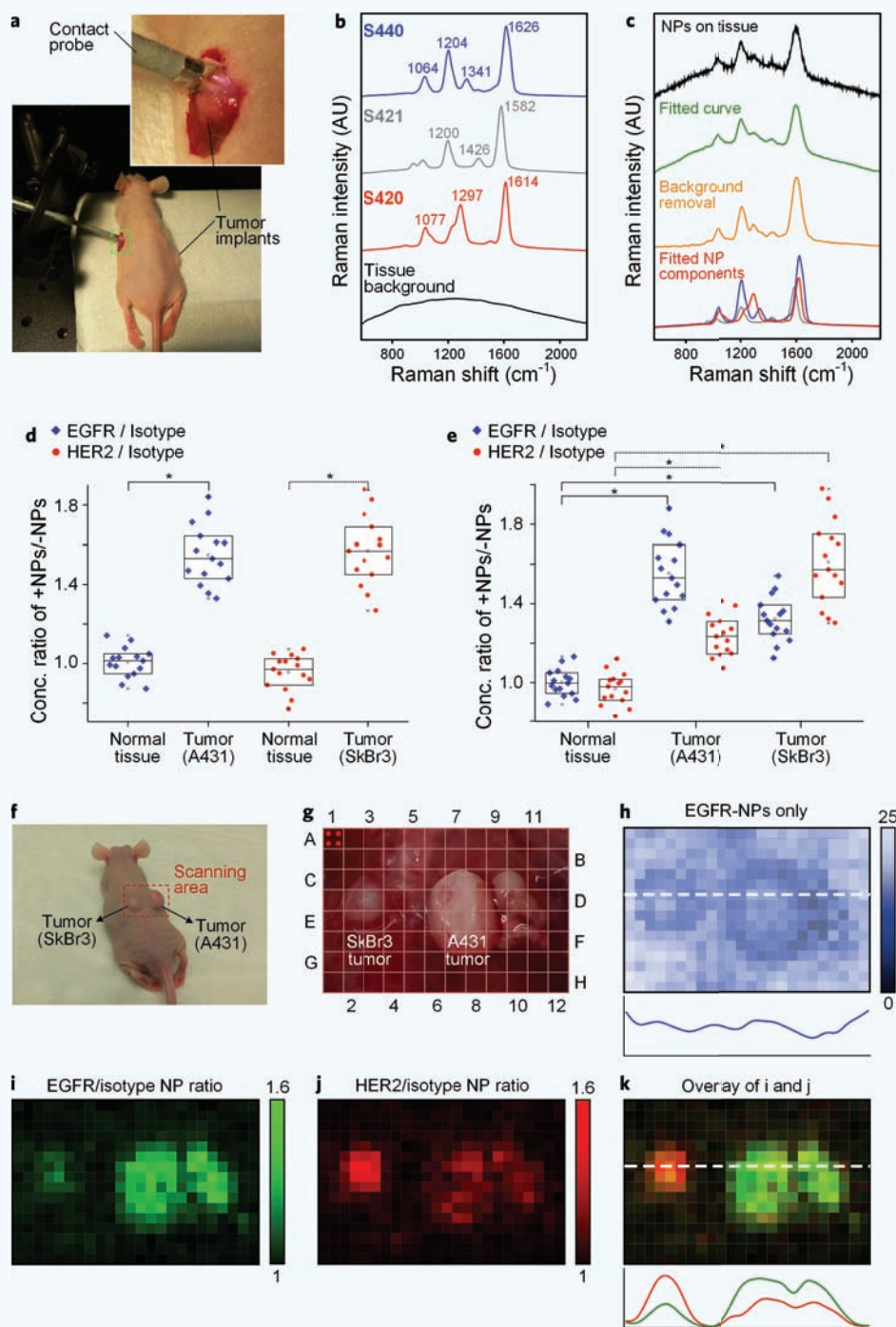


Figure 5 *In vivo* ratiometric analysis of multiplexed NPs on tumor implants. (a) Mouse with surgically exposed tumors; the inset provides a magnified view of the 2.5-mm diameter flexible Raman probe. (b) Reference Raman spectra of pure SERS NPs (red: S420, gray: S421 and blue: S440) and tissue background with no NPs (black). (c) Raw spectra of NPs applied on tissue acquired with a 0.1-s integration time (black), best-fit curve using a DCLS algorithm (green), spectra of NPs on tissue after tissue-background removal using a DCLS algorithm (orange), and the DCLS-demultiplexed NP spectra (blue: EGFR-S440, red: HER2-S420, gray: isotype-S421). The concentration ratio of targeted and nontargeted NPs typically applied on exposed tumors and normal tissues is plotted for (d) 2-flavor and (e) 3-flavor multiplexed NPs ($n = 15$). * P -value < 0.001 . (f–k) Image-grid experiment. (f) Mouse with two adjacent tumor xenografts. (g) Photograph of stained tissue with a virtual grid of detection locations. Four positions were probed within each grid coordinate (e.g. refer to the red dots in coordinate A1) such that 24×16 positions were detected in total; (h) Map of the absolute concentration (pM) of EGFR-NPs; (i) Map of the concentration ratio of EGFR/isotype NPs. (j) Map of the concentration ratio of HER2/isotype NPs. (k) Overlay of EGFR/isotype and HER2/isotype maps.

EGFR-targeted, and HER2-targeted). *In vivo* experiments were repeated with two mice (see **Supplementary Fig. 11**) and yielded results that are consistent with what is presented in **Fig. 5e**.

To further demonstrate the utility of the ratiometric analysis of multiplexed SERS NPs, we scanned a grid of positions over the two tumors (A431 and SkBr3) and surrounding normal tissues on one mouse and mapped the concentration ratios of targeted/nontargeted NPs (**Fig. 5f–k**). For this experiment, a mouse with A431 and SkBr3 tumors implanted adjacent to each other was utilized (**Fig. 5f**). Before detection, a mixture of EGFR-NPs, HER2-NPs and isotype-NPs (200 pM per flavor) was topically applied to stain a $4.2 \times 2.8 \text{ cm}^2$ tissue area for 5 min, followed by a PBS rinse step (20 s). A total of 24×16 positions were detected with a 0.1-s integration time at each point (see **Fig. 5g**). In **Fig. 5h**, the absolute concentration of EGFR-NPs is mapped to simulate a non-ratiometric imaging strategy using a single tumor-targeted contrast agent. Consistent with our *ex vivo* results (**Fig. 4c**), the normal tissues in **Fig. 5h** show higher absolute intensities compared with the tumors due to enhanced nonspecific accumulation of the NPs, which is potentially misleading because of the apparent inverse contrast. **Figure 5i,j** are maps of the concentration ratio of EGFR-NP/isotype-NP and HER2-NP/isotype-NP, respectively. Both maps demonstrate improved contrast between tumor (color) and normal tissues (black) and provide quantification of specific binding to molecular targets. These results are consistent with the previous flow cytometry, *in vitro* monolayer and *ex vivo* tissue measurements, in which A431 tumors express high levels of EGFR and moderate levels of HER2, whereas SkBr3 tumors express high levels of HER2 and moderate levels of EGFR.

DISCUSSION

Interest in SERS-coded NPs has been accelerating in recent years due to their potential for highly multiplexed detection with negligible photobleaching when illuminated with a single low-power laser source. The relatively large size of these NPs ($\sim 120 \text{ nm}$) allows them to be used for the detection of protein biomarkers expressed at the surface of epithelia and surgically exposed tissue with minimal systemic uptake and toxicity^{39,53}. However, to date, there has

not been a demonstration of the ability to rapidly quantify the specific binding of multiplexed SERS NPs *ex vivo* and *in vivo* following a brief topical application and rinse-removal protocol. Hence, we used a miniature spectral probe and a calibrated ratiometric analysis method to investigate the performance of SERS NPs under clinically relevant conditions, demonstrating the ability to accurately quantify molecularly specific vs. nonspecific NP accumulation on tumor and normal tissues with a spectral integration time of 0.1 s after a 5-min topical application and 20 s of rinsing.

We present *in vitro*, *ex vivo* and *in vivo* data to illustrate the progressively more challenging nature of these experimental conditions. For example, cell suspensions are easily stained and rinsed, yielding specific-to-nonspecific binding ratios of nearly two orders of magnitude in flow cytometry experiments (Fig. 1). These results show an unprecedented ability of the large NPs to bind to cell-surface receptors and were achieved with a short staining duration of only 15 min. Monolayers of adherent cells are less efficient to stain and rinse, and yield specific-to-nonspecific binding ratios of 2- to 3-fold when measured with a miniature probe (Fig. 3). Finally, *ex vivo* and *in vivo* tissue xenografts are the most challenging to stain with exogenous molecular contrast agents (Fig. 4,5), as they exhibit nonspecific permeability and retention effects as well as a host of nonspecific chemical targets that reduce detection contrast. For example, Figs. 4c and 5h demonstrate how measuring the absolute concentration of targeted NPs can be misleading due to a larger uptake and retention in normal tissues vs. tumors (*i.e.* inverse contrast). Additional sources of misleading contrast include variations in probe working distance and illumination power, as well as uneven NP application and rinse removal at different tissue locations.

In light of the potentially misleading nonspecific sources of contrast mentioned in the previous paragraph, calibrated ratiometric detection with multiplexed positive- and negative-control contrast agents is a valuable strategy for accurately quantifying specific vs. nonspecific signals⁴³⁻⁴⁵. SERS NPs are particularly well-suited for ratiometric detection, not only because of their high multiplexing capability but also due to the ability of multiple SERS NP flavors to be excited at a single illumination wavelength, ensuring that all NP flavors in a single measurement are interrogated identically in terms of illumination intensity, spot size and effective excitation depth (see SI). In addition, while it is possible under idealized experimental conditions to perform calibrated measurements of absolute NP concentrations (Fig. 4c), these measurements are potentially not as accurate as measurements of concentration ratios. This could be due, for example, to slight differences in sample conditions and measurement geometries between the calibration measurements and the actual experiments. Ratiometric measurements are much less sensitive to these differences, and therefore more accurate, due to the fact that the effects of experimental variations tend to cancel out in the ratios (see SI for additional discussions).

A surprising and significant result of this study is the demonstration that accurate *in vivo* molecular detection is optimally achieved with a topical application of only 5 min in duration. We hypothesize that the nonspecific transport and binding of these NPs within tissues at longer staining durations impedes the ability to easily remove unbound NPs *via* rinsing. While brief staining durations are ideal for the translation of these technologies into clinical settings, the binding affinity of the targeted NPs must be very high in order to achieve significant levels of specific binding at such short time scales. Fortunately, the molecularly targeted NPs developed in this study exhibit sufficiently high binding affinities under brief *in vitro* staining conditions (Fig. 1) that they may also be effectively utilized for the rapid *ex vivo* and *in vivo* detection of biomarkers in fresh tissues.

For rapid clinical diagnostics, additional work is necessary to further optimize the binding affinity of targeted SERS NPs, as well as to minimize nonspecific binding. In addition, while the brightness of the SERS NPs used here allows for the detection of low-pM concentrations of NPs at 0.1-s integration times, brighter NPs are needed to apply these

contrast agents in a greater variety of biomedical detection and imaging applications. Furthermore, while studies have shown the feasibility of detecting and demultiplexing large panels (5 – 10) of nontargeted SERS NPs in animals and *ex vivo* human tissues, further work is needed to demonstrate the ability to quantify a large panel of protein biomarkers with targeted NPs^{21,22,32-36}. While there are many challenges that must be resolved to improve SERS NPs for molecular diagnostics, this study optimizes across multiple disciplines — including a bioconjugation protocol, portable-measurement device, quantitative-detection assay, and a topical-staining protocol — to demonstrate the feasibility of rapidly and quantitatively detecting a panel of cancer biomarkers on *ex vivo* and *in vivo* tissues with multiplexed SERS NPs under time-constrained conditions. These tools have the potential to greatly improve our ability to investigate the mechanisms of disease progression as well as for early detection, surgical guidance, and for guiding/monitoring personalized therapies in the clinic.

METHODS

Fluorophore and antibody conjugation to SERS NPs

The NPs (Cabot Security Materials Inc, Mountain View, CA) utilize a gold core and are encapsulated in silica. The gold-core diameters are ~60 nm and the overall dimension is ~120 nm. Additional details may be found in the literature⁹. The silica surface of the NPs are functionalized with thiols to allow conjugation to a variety of targeting molecules (Fig. 6a). Three “flavors” of NPs were used, identified as S420, S421 and S440, each of which emits a characteristic Raman spectrum (Fig. 5b).

Stock concentrations of SERS NPs (800 pM in water) were diluted in 10 mM MOPS (Sigma-Aldrich, part No. M1254) buffer, pH 7.25, at a volume ratio of 1:1 (*e.g.* 200 μ L buffer to 200 μ L NPs in water), and

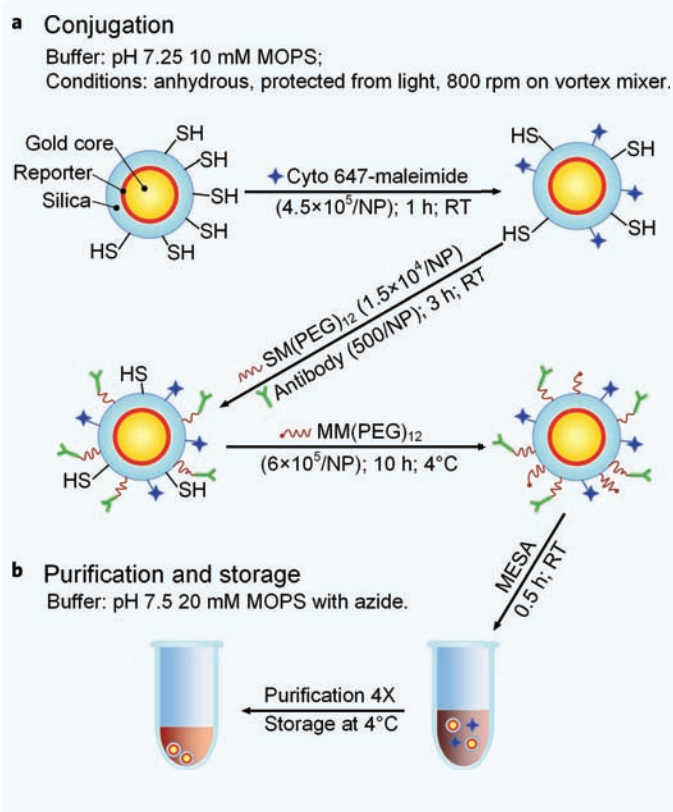


Figure 6 Schematic illustration of the preparation of SERS NPs conjugated to monoclonal antibodies and a fluorophore for flow cytometry. (a) Conjugation and (b) purification.

then reacted with Cyto 647-maleimide (Cytodiagnosics Inc, part No. NF647-3-01), at 4.5×10^5 molar equivalents per NP, at room temperature for 1 h to conjugate Cyto 647 fluorophores to the NP surface (Fig. 6a). Three different monoclonal antibodies (mAb) were used for conjugation: anti-EGFR “panitumumab” (Vectibix, NDC 55513-954-01), anti-HER2 (Thermo Scientific, MS-229-PABX), or an isotype control mAb (Thermo Scientific, MA110407). These mAbs were first eluted over a desalting spin column (Thermo Scientific, 89882) and buffer-exchanged into the MOPS buffer to remove preservatives such as sodium azide. All mAbs were purchased free of protein stabilizers such as BSA or gelatin. Antibodies were added to fluorescent NPs at 500 molar equivalents per NP, along with the heterobifunctional PEG cross linker SM(PEG)₁₂ (Thermo Scientific, 22112), at 1.5×10^4 molar equivalents per NP, and incubated at room temperature for 3 h. Following the primary conjugation reaction, MM(PEG)₁₂ (Thermo Scientific, 22711), at 6×10^5 molar equivalents per NP, was then added to the NPs and reacted over night at 4°C to block residual thiols on the NPs. Note that all reagents were degassed with dry argon gas prior to use and all reactions were conducted under anhydrous conditions in light-protected amber vials (Fisher Scientific, 03-391-36&03-391-17) on a vortex mixer (Fisher Scientific™ Microplate Vortex Mixers, 02-216-101) set at 800 rpm. Finally, the NPs were reacted at room temperature with 2-mercaptoethanesulfonate (Sigma-Aldrich, M1511) for 30 min, at 9×10^5 molar equivalents per NP, and the reaction mixture was then diluted at a volume ratio of 1:1 with storage buffer (20 mM MOPS at pH 7.5 with 0.1% BSA (Jackson Immuno, 001-000-162) and 0.05% sodium azide (Sigma-Aldrich, S2002)). The diluted reaction mixture was purified four times *via* centrifugation (1000 g for 10 min), in which the supernatant was removed and replaced with storage buffer after each round of centrifugation. The conjugated NPs were stored at 4°C and protected from light. UV-VIS spectrophotometry was used to measure the final concentration of the conjugated NPs (Supplementary Fig. 2).

Customized spectral-imaging device

A miniature spectral-imaging device was developed to quantify the concentration and concentration ratios of SERS NPs applied on cells and tissues. The detection probe is cylindrical with an outer diameter of 2.5 mm. A singlemode fiber at the center of the probe is used to illuminate the sample, and 36 multimode fibers (200-micron core) surrounding the singlemode fiber are used to collect optical signals including Raman-scattered light (additional details about the detection system have been described previously²³). A low power 785-nm diode laser (10 mW, well below ANSI safety limits) is used to illuminate the tissue with a spot size of ~200 μm. In order to minimize specular reflections, the probe is angled at 20 to 45 deg with respect to the direction normal to the sample surface (Fig. 2 and Supplementary Fig. 9). A custom spectrometer (Bayspec Inc.) is used to disperse the collected signal onto a cooled deep-depletion spectroscopic CCD (Andor Newton, DU920P-BR-DD). Detector integration times ranging from 0.1 to 2 s were used. The linearity of our device for measurements of 3-flavor NP mixtures (of various ratios) was assessed and is shown in Supplementary Fig. 6.

For monolayer measurements in multiwell plates, a probe holder was used to maintain a constant working distance (~2 mm) and probe angle (20–25 deg) (Supplementary Fig. 9). For *ex vivo* and *in vivo* tissue measurements, a glass prism (Tower Optical Inc., 4531-0006) was adhered to the probe tip to create a fixed-angle and fixed-working-distance contact surface with the tissue (refer to Fig. 2d and design optimization in Supplementary Fig. 1).

Cell culture, flow cytometry, and cell monolayers

A431 and 3T3 cells were cultured in DMEM (Lonza, 12-604F) and SkBr3 cells were cultured in Mccoy's 5A (Lonza, 12-688F), both of which were supplemented with 10% fetal bovine serum (FBS, Thermo Scientific, SH3008803) and 1% penicillin-streptomycin (Lonza, 17-602E). Cells were

cultured at 37 °C at a CO₂ level of 5.0%. Trypsin EDTA 1X (Mediatech, MT25051CI) was used to detach adherent cells for flow cytometry.

For flow cytometry, 50-μL cell suspensions (0.2 million cells) were reacted with 50 μL of 300-pM antibody-conjugated NPs and incubated for 15 min at room temperature protected from light under gentle agitation with a vortex mixer (Fisher Scientific™ Microplate Vortex Mixers, 02-216-101) at 300 rpm. After staining, the cells were purified three times *via* centrifugation (400 g for 5 min) and supernatant-replacement (500 μL per rinse) with FACS buffer (20% FBS in PBS), and then stored in 500 μL of FACS buffer. Competitive binding experiments were also performed to verify the specificity of the targeted SERS NPs. For these experiments, A431 and SkBr3 cells (0.2 million in 50 μL) were incubated 4 h prior to NP staining with 50 μL of 10 mg·mL⁻¹ panitumumab and anti-HER2 mAb to competitively inhibit the cell-surface receptor targets (EGFR or HER2).

For *in vitro* experiments, a 96-well plate with a 200 μm-thick glass bottom was used to culture cells into monolayers. For each cell line, cells were counted and seeded into four wells (1×10^5 cells in 50 μL of media per well for A431 and SkBr3, and 0.5×10^5 cells in 50 μL of media per well for 3T3). After incubation for 24 hours, a similar confluent cell monolayer was observed in all wells (Supplementary Fig. 7). For Raman measurements, 1 of the 4 wells for each cell line was used to acquire a background measurement in the absence of NPs (Supplementary Fig. 8). For the three remaining wells of each cell line, an equimolar mixture of targeted (anti-EGFR or anti-HER2) and nontargeted (isotype control) NPs was added for ratiometric analysis of specific vs. nonspecific binding of the NPs. Both 2-flavor and 3-flavor mixtures were used at a staining concentration of 150 pM per flavor and a total staining volume of 100 μL. Staining was performed for 15 min, followed by up to three rounds of rinsing with PBS. Spectral measurements and ratiometric analyses were performed before and after each rinse step.

Xenograft tumor model

All animal work was approved by the Institutional Animal Care and Use Committee (IACUC) at Stony Brook University. Five male nude mice at 7–9 weeks of age (Taconic Farms Inc, model NCRNU-F) were subcutaneously implanted with tumor cells in their flank under isoflurane inhalation anesthesia (Baxter, 1001936060). 1×10^6 A431 cells suspended in 50% matrigel (BD biosciences, 354234) (200 μL mixture) were injected into one flank and 5×10^6 SkBr3 cells in 50% matrigel (300 μL mixture) were injected into the contralateral side. After 1–4 weeks, the injected cells developed into tumors of approximately 5 to 10 mm in diameter.

For *ex vivo* imaging, the mice were anesthetized and euthanized by cervical dislocation, followed by the surgical removal of tumors and normal tissues of a similar size. The excised tumors and normal tissues were cut into small pieces (30–60 mm³) and hydrated in FACS buffer for background acquisition and NP staining. For *ex vivo* staining, pieces of fresh tumor and normal tissue were submerged for 5 minutes in an equimolar mixture of 2 or 3 flavors of conjugated NPs (200 pM per flavor). After incubation, the pieces were washed up to 7 times by submerging them for 3 s per wash, with gentle agitation, in a large volume of PBS (20–50 mL).

For *in vivo* imaging, the mice were anesthetized *via* intraperitoneal injection of ketamine and xylazine and then the tumors were surgically exposed. An equimolar mixture of 2 or 3 flavors of conjugated NPs (200 pM per flavor) was topically applied to tumor and normal tissues for 5 min. After rinsing off unbound NPs with PBS, the tumor and normal tissues were scanned using the contact probe (Fig. 2d). After imaging, the mice were euthanized.

Statistics

For *ex vivo* and *in vivo* experiments, a two-sample *t*-test was performed to determine the significance of the difference in the positive/negative NP ratio (EGFR/isotype or HER2/isotype) between tumors and normal tissues. A *p*-value < 0.001 was considered significant. Data points represent

average values with error bars denoting one standard deviation from the mean. For the box plots in **Figs. 4d,e** and **5d,e**, the bottom and top of the box represent the 1st and 3rd quartiles of the dataset, respectively, and the band inside the box represents the median (2nd quartile) of the data.

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AUTHOR CONTRIBUTIONS

Y.W. designed experiments, performed nanoparticle conjugations and experiments, analyzed data, and wrote the manuscript. A.K. set up the Raman hardware, completed the demultiplexing algorithm and Labview software. M.S., D.W., Y.C., S.Y.L., and D.M. provided assistance and training for various experiments. P.Z.M. and B.C.W. provided training and support for nanoparticle conjugations. J.T.C.L. initiated the project, designed experiments, provided training and support for various experiments, analyzed data, supervised and coordinated all investigators on the project, and wrote the manuscript.

SUPPLEMENTARY INFORMATION AVAILABLE

Demultiplexing SERS spectra using direct classical least squares; ratio-metric quantification of specific vs. nonspecific binding; optimization of a miniature hand-held probe for contact detection of SERS NPs on tissues (**Supplementary Fig. 1**); calculating the concentration of NP conjugates via UV-VIS spectrophotometry (**Supplementary Fig. 2**); optimization of conjugation and staining parameters (**Supplementary Table 1**); flow cytometry comparison of different control NPs with cultured cells (**Supplementary Fig. 3**); observation of NP distribution on cells via confocal microscopy (**Supplementary Fig. 4**); confocal microscopic images of different cells stained with conjugated NPs (**Supplementary Fig. 5**); limit-of-detection (linearity) test for a 3-flavor mixture (**Supplementary Fig. 6**); microscopic images of confluent cell monolayers of 3T3, A431 and SkBr3 in one well of a 96-well plate (**Supplementary Fig. 7**); schematic of cell-monolayer experiments on a 96-well plate (**Supplementary Fig. 8**); experimental platform for the spectral detection of cell monolayers on a well plate (**Supplementary Fig. 9**); coefficient of variation of the measured concentration of NPs on tissues under different integration times (**Supplementary Fig. 10**); effect of NP flavor on ratiometric analysis (**Supplementary Fig. 11**); Raman signal of NPs in different buffers (**Supplementary Fig. 12**). This material is available free of charge.

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SUPPLEMENTARY INFORMATION

Demultiplexing SERS spectra using direct classical least squares (DCLS)

This study employs a demultiplexing method described previously¹. In a realistic spectral measurement, pure spectra from single- or multi-flavor particle mixtures are mixed with varying magnitudes of broadband background signal (mainly due to laser background and autofluorescence background) and zero-mean Gaussian-distributed white noise (includes shot noise and other stochastic noise sources such as detector readout noise and dark counts). Other than sources of noise, it is assumed that each measured spectrum consists of a weighted sum of fixed nanoparticle spectra (F_n) and broadband background signals (B). Based on the assumption that the combination is linear, we employ a linear least-squares algorithm to compute the relative nanoparticle weights (w_n). A third-order polynomial is included to account for broadband background signals that are not captured by the reference spectrum (B).

$$S = \sum_n w_n F_n + kB + \sum_m a_m P_m + R \quad (1)$$

where:

- S = measured spectral data
- w_n = weight of SERS flavor n
- F_n = known reference spectrum of SERS nanoparticle flavor n
- k = scaling factor for background signal magnitude
- B = known reference spectrum of broadband background
- a_m = weight of m th-order polynomial term
- P_m = m th-order polynomial term (for baseline correction)
- R = residual (minimized by least-squares algorithm)

In order to process raw spectral data, we first subtract the constant detector background that is obtained with the laser illumination turned

off. This background is due to detector noise, which is primarily electronic read noise and a smaller amount of dark thermal noise (dependent upon integration time and the level of detector cooling). The spectra are cropped and only the range 700–2000 cm^{-1} (Raman shift) is used for demultiplexing analysis.

Reference spectra are obtained for pure NP flavors, F_n , as well as for the broadband background, B . In order to obtain a NP reference spectrum, a 3–5 μL drop of NPs (800 pM concentration) is placed on the surface of the spectral-detection device, or at a fixed working distance away from the device. A spectrum is recorded from the NP sample. Next, another spectrum is recorded from a drop of the buffer in which the NPs were suspended (generally water), under the exact same conditions (*e.g.* droplet volume, detection geometry, detector settings). These two spectra are subtracted from each other to obtain the pure reference spectrum of the NPs. In order to obtain a background reference spectrum, B , the sample (cells or tissue) is stained with an appropriate buffer solution (generally PBS) that is devoid of NPs. A spectrum of the buffer-stained sample is acquired under the exact same conditions (*e.g.* laser and detector settings, measurement geometry, etc.) as is used for measuring NP-stained samples. In certain cases, multiple spectra are acquired to obtain an average background reference spectrum, B . Alternatively, a principle component analysis (PCA) may be performed to determine the most appropriate reference spectrum or spectra (if more than one background reference spectrum is desired).

Ratiometric quantification of specific vs. nonspecific binding

For ratiometric quantification of specific vs. nonspecific binding, it is assumed that negative-control NPs exhibit identical nonspecific behavior to targeted NPs. In this study, targeted NP flavors are conjugated to monoclonal antibodies (anti-EGFR or anti-HER2) and a negative-control NP flavor is conjugated to an appropriate isotype-control antibody. NP characteristics such as charge, size, and overall chemistry are therefore very similar between NPs, except for the antigen-binding sites of the targeted mAbs. With these assumptions, the measured weight, w_n , of each NP flavor, as determined by least-squares demultiplexing, can be modeled as:

$$w_n = g_n \cdot I_n \cdot c_n \cdot \sigma_n \quad (2)$$

where:

- g_n = electronic and system gain (detector quantum yield, optical throughput, etc.)
- I_n = laser power
- c_n = concentration of NP flavor n
- σ_n = average scattering cross section of NP flavor n

The ratiometric strategy seeks to determine the true concentration ratio between NP flavors:

$$\frac{c_1}{c_2} = \frac{w_1}{w_2} \cdot \frac{g_2 I_2 \sigma_2}{g_1 I_1 \sigma_1} = \frac{w_1}{w_2} \cdot \kappa \quad (3)$$

The least squares demultiplexing algorithm yields the weight ratio, w_1/w_2 . We assume that the ratio $g_2 I_2 \sigma_2 / g_1 I_1 \sigma_1$ is a constant (κ) and can be determined *via* a calibration measurement of a known concentration ratio of NPs, typically an equimolar ratio where $c_1/c_2 = 1$.

This assumption is based on the following arguments:

1. Under ideal demultiplexing conditions, in which there is negligible crosstalk between NP flavors in the least-squares algorithm, the measured weights, w_n , are linear and directly proportional to the actual NP concentrations, c_n .
2. A single laser illumination source is used to illuminate all NPs identically within each tissue region, such that $I_n = I_1 = I_2$ for all NP flavors.
3. The electronic and system gain are either constant and identical for each NP flavor (*e.g.* optical throughput) or the ratio of these

are unable to diffuse well within tissues, and are largely confined to the superficial surface when topically applied on tissues².

(Ex vivo and in vivo) optimization of a miniature hand-held probe for contact detection of SERS NPs on tissues

One challenge in using a hand-held probe for NP detection is that the working distance and detection angle (see **Supplementary Fig. 1a**) are difficult to control. In particular, for *in vivo* experiments, uneven tissue surfaces and motion from respiration can cause fluctuations in working distance and detection angle. While our ratiometric quantification strategy is designed to mitigate these effects, large variations in signal strength (SNR) and signal-to-background ratio (SBR) can reduce the accuracy of quantitative measurements of NPs located on tissue surfaces. Therefore, here, we have used contact detection to minimize working-distance and angle-dependent effects. The probe consists of a prism to fix the working distance and detection angle, where experiments were performed to determine the optimal working distance and detection angle. For these measurements, mice were anesthetized *via* intraperitoneal injection of ketamine and xylazine, and a small flat area of tissue was stained with an equimolar mixture of 3 flavors of conjugated NPs (200 pM per flavor).

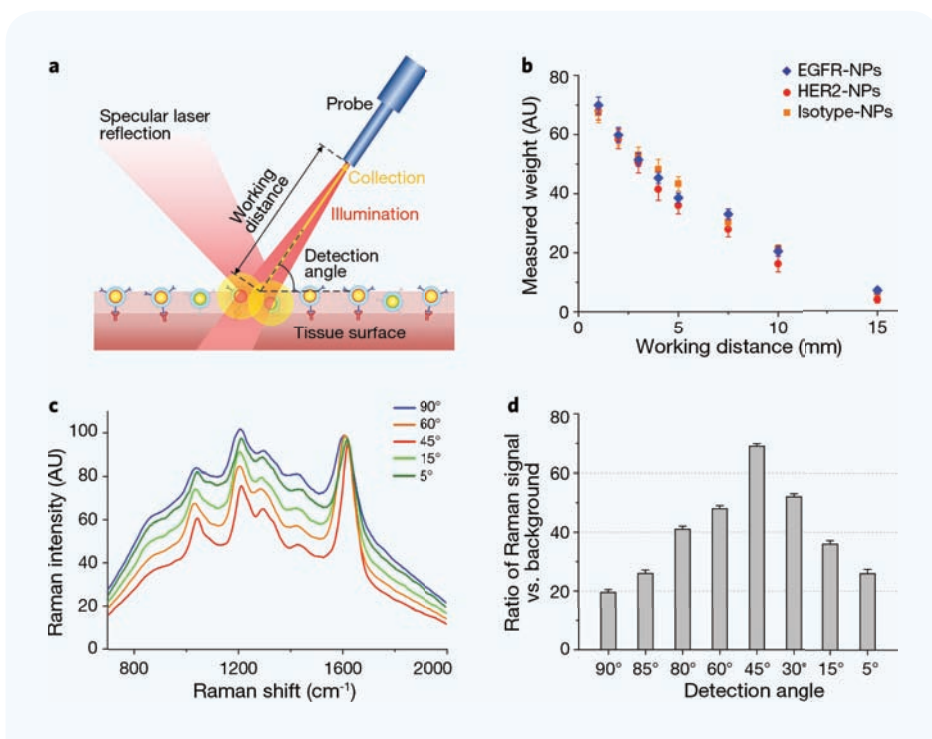
Supplementary Fig. 1b plots the variation of demultiplexed NP weights as a function of working distance (1–15 mm at a 45° detection angle), showing that signal strength degrades with working distance and that a contact probe (zero working distance) yields the highest detection sensitivity. These results are expected due to the reduction in the numerical aperture (NA) of collection with working distance. **Supplementary Fig. 1c** shows raw spectra collected under different detection angles and **Supplementary Fig. 1d** plots the ratio of Raman signal versus tissue background. The data suggest that detection at 45° provides an optimal SBR, likely due to the reduced collection of specularly reflected laser light. Based on these results, a 2.5-mm 45° prism was adhered to the probe tip to fix the detection angle at ~45° for *ex vivo* and *in vivo* experiments.

Optimization of conjugation and staining parameters

Supplementary Table 1 Flow-cytometry fluorescence ratio of HER2-NPs vs. isotype-NPs on SkBr3 cells under different conjugation and staining conditions.

Antibody/ NP ratio	MM(PEG) ₁₂ amount	Staining conc. on cells (pM of NPs)	Staining time on cells (min)	FI ratio
350	0	40	15	9
350	6 × 10 ⁶ /NP	40	15	11
350	6 × 10 ⁶ /NP	100	15	21
350	6 × 10 ⁶ /NP	100	60	14
350	6 × 10 ⁶ /NP	200	15	49
420	6 × 10 ⁶ /NP	200	15	63
500*	6 × 10 ⁶ /NP	200	15	92

*For antibody/NP ratios greater than 500, aggregation of the conjugated NPs becomes increasingly apparent.



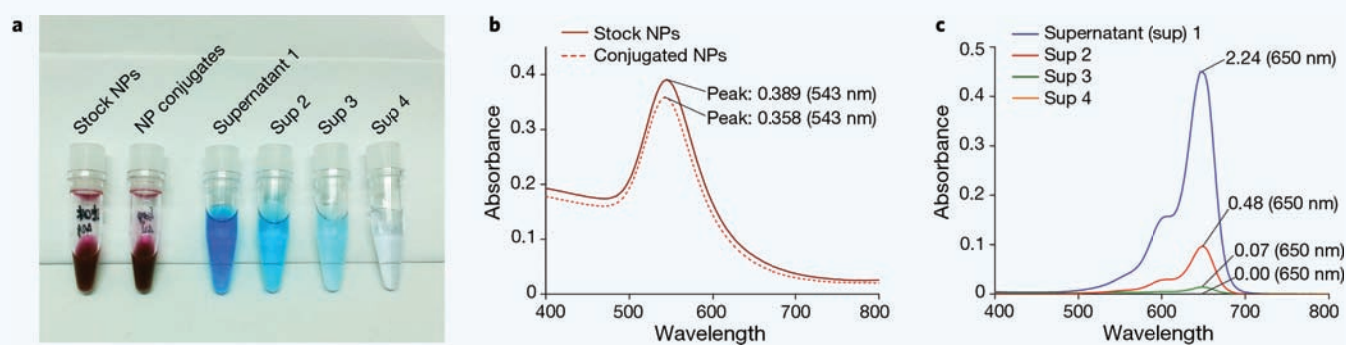
Supplementary Figure 1 Optimization experiments of a miniature hand-held contact probe for Raman detection. **(a)** Schematic illustration of the detection of NPs on tissue surfaces. **(b)** Effect of working distance on demultiplexed NP weights (signal intensity) with the handheld probe ($n = 5$). The probe has a detection angle of 45° with a variable working distance ranging from 1 mm to 15 mm. **(c)** Raw spectra as a function of the detection angle. All spectra are normalized by the magnitude of the main peak at $\sim 1620 \text{ cm}^{-1}$. **(d)** Ratio of Raman signal (from NPs) versus background signal (from tissue autofluorescence and laser reflections).

constants (g_2/g_1) are constant due to a linear behavior of these constants with respect to perturbations (*e.g.* fluctuations in detector quantum yield due to temperature changes, variations in signal due to changes in detector integration time, and/or variations in signal due to changes in measurement geometry). In practice, all calibration measurements are performed with the exact same detector and system settings as the actual experiments, and the individual gains g_n can be treated as constants. However, this is not strictly required, since a single detector and optical detection path are used to record all NP signals and the ratio g_2/g_1 remains constant even under variable gain settings.

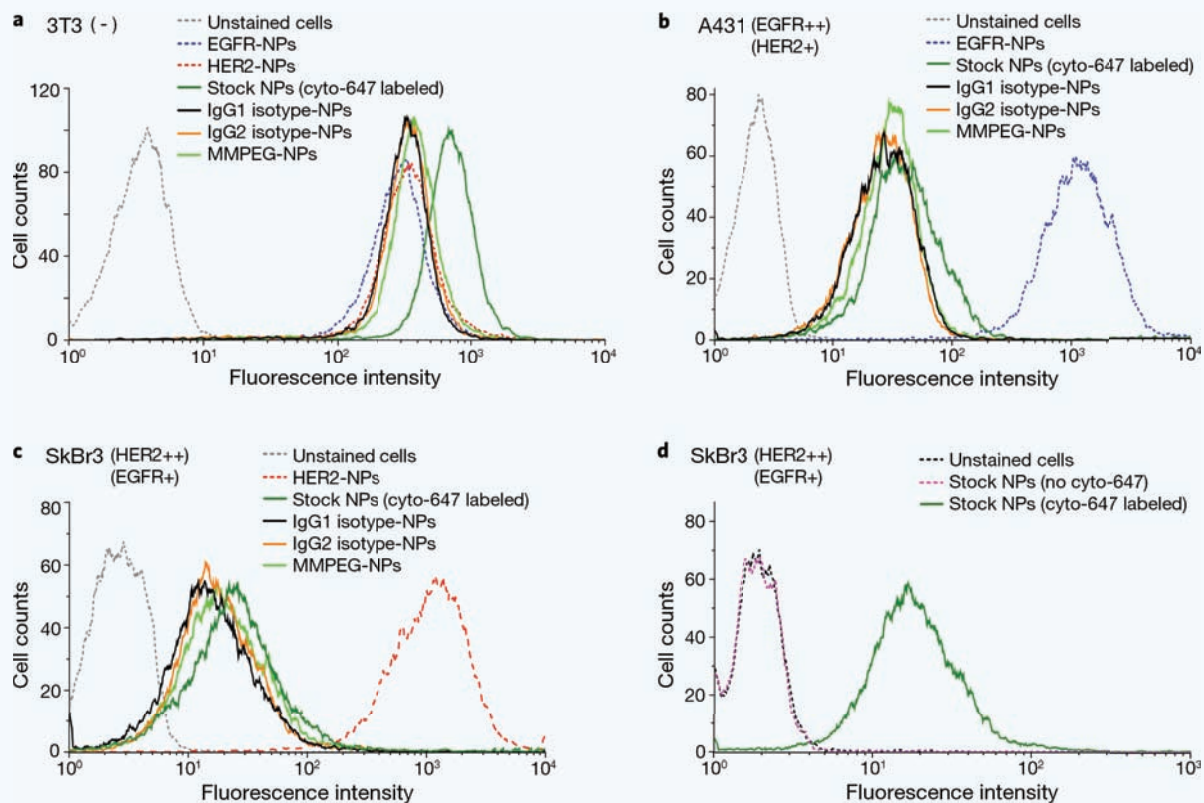
- The ratio of the scattering cross sections σ_2/σ_1 is a constant due to the fact that the Raman-active layer in the SERS NPs is encapsulated in silica and shielded from environmental factors. Any perturbation in the scattering cross sections — due to optical, thermal, or other effects — is assumed to affect all NP flavors in a linear fashion such that σ_2/σ_1 remains constant. In practice, all calibration measurements are performed under the same optical and environmental conditions as the actual experiments and the individual cross sections σ_n can be treated as constants. However, this is not strictly required.

Note that with targeted SERS NPs, it is possible that as the NPs diffuse through tissues, the differential binding between various NP flavors may alter the initial ratio of the NP mixtures within the staining “reservoir” of NPs. Therefore, the initial 1:1 ratio between all NP flavors in the staining solution may not be true deeper within tissues. However, due to the large size of our SERS NPs ($\sim 120 \text{ nm}$ in diameter), the NPs

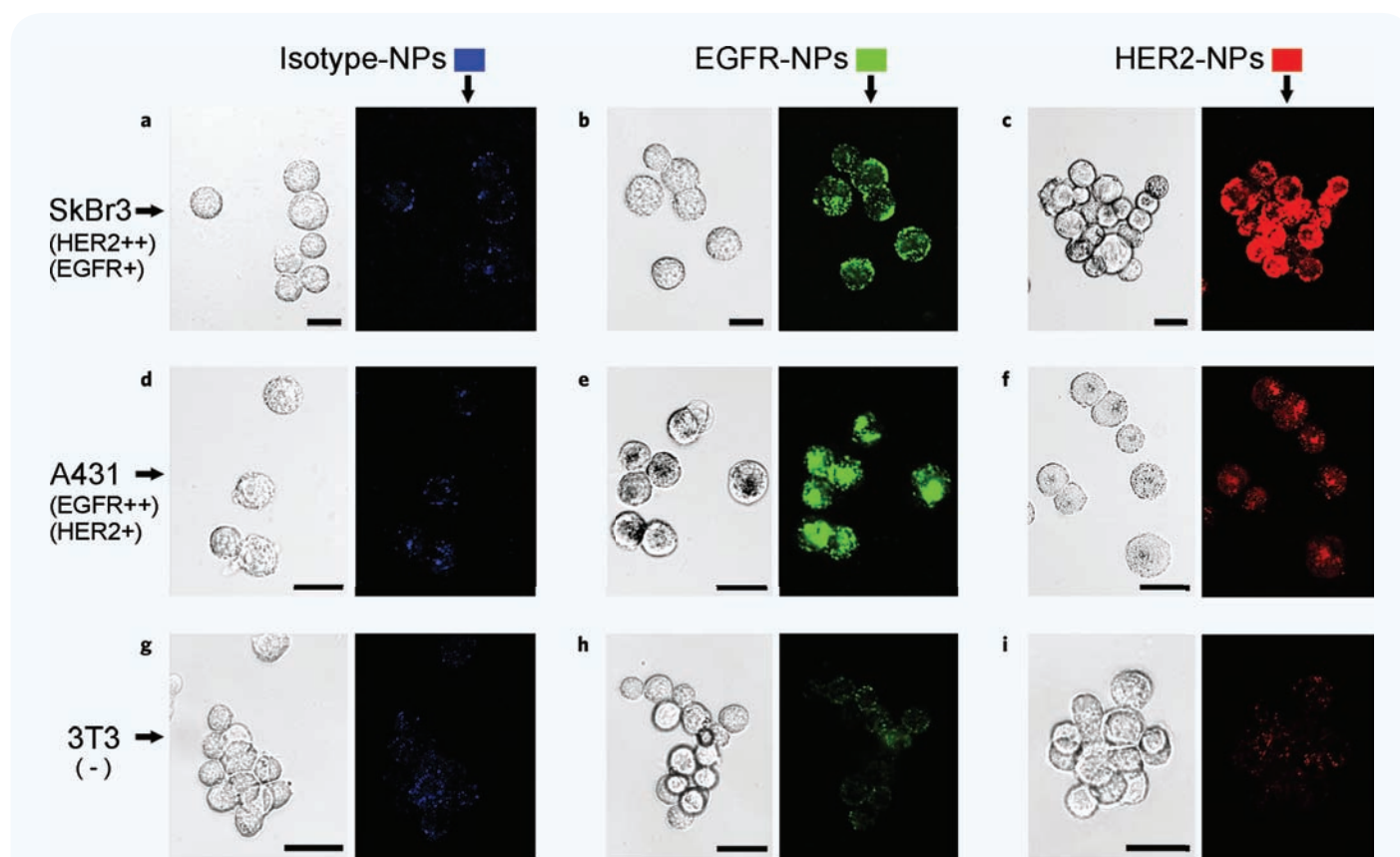
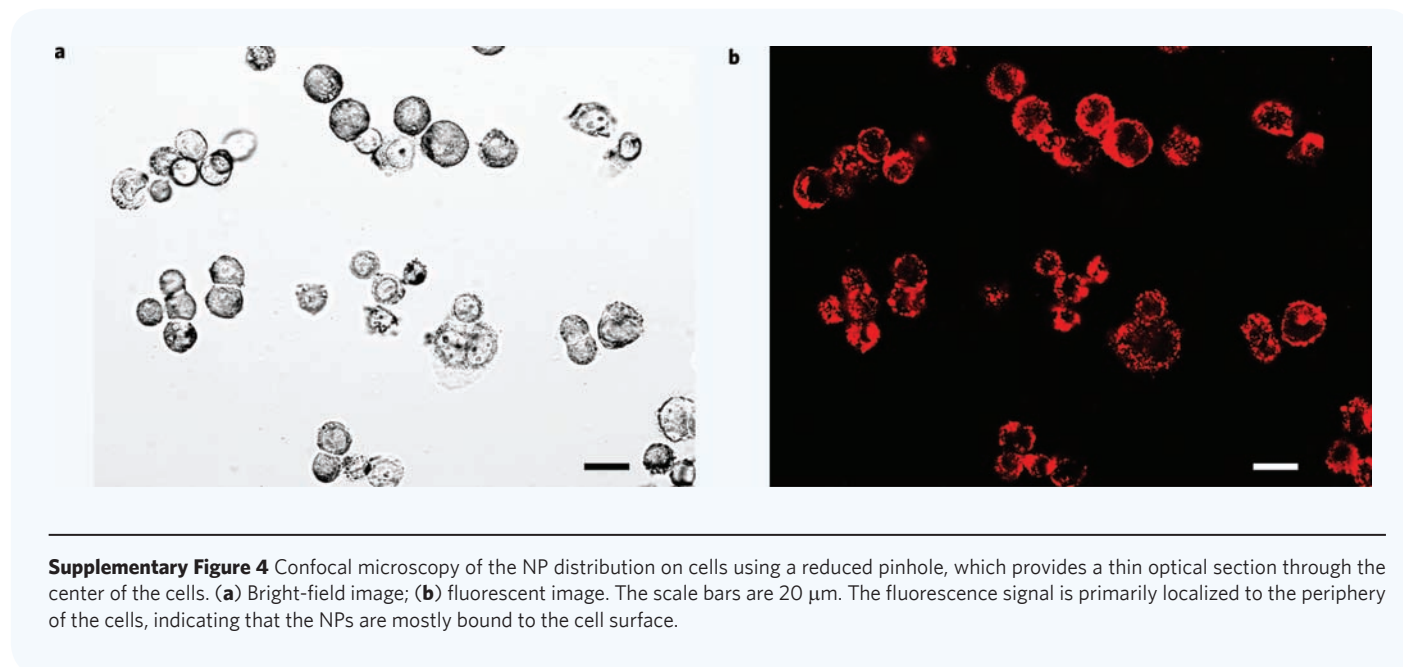
Calculating the concentration of NP conjugates via UV-VIS spectrophotometry

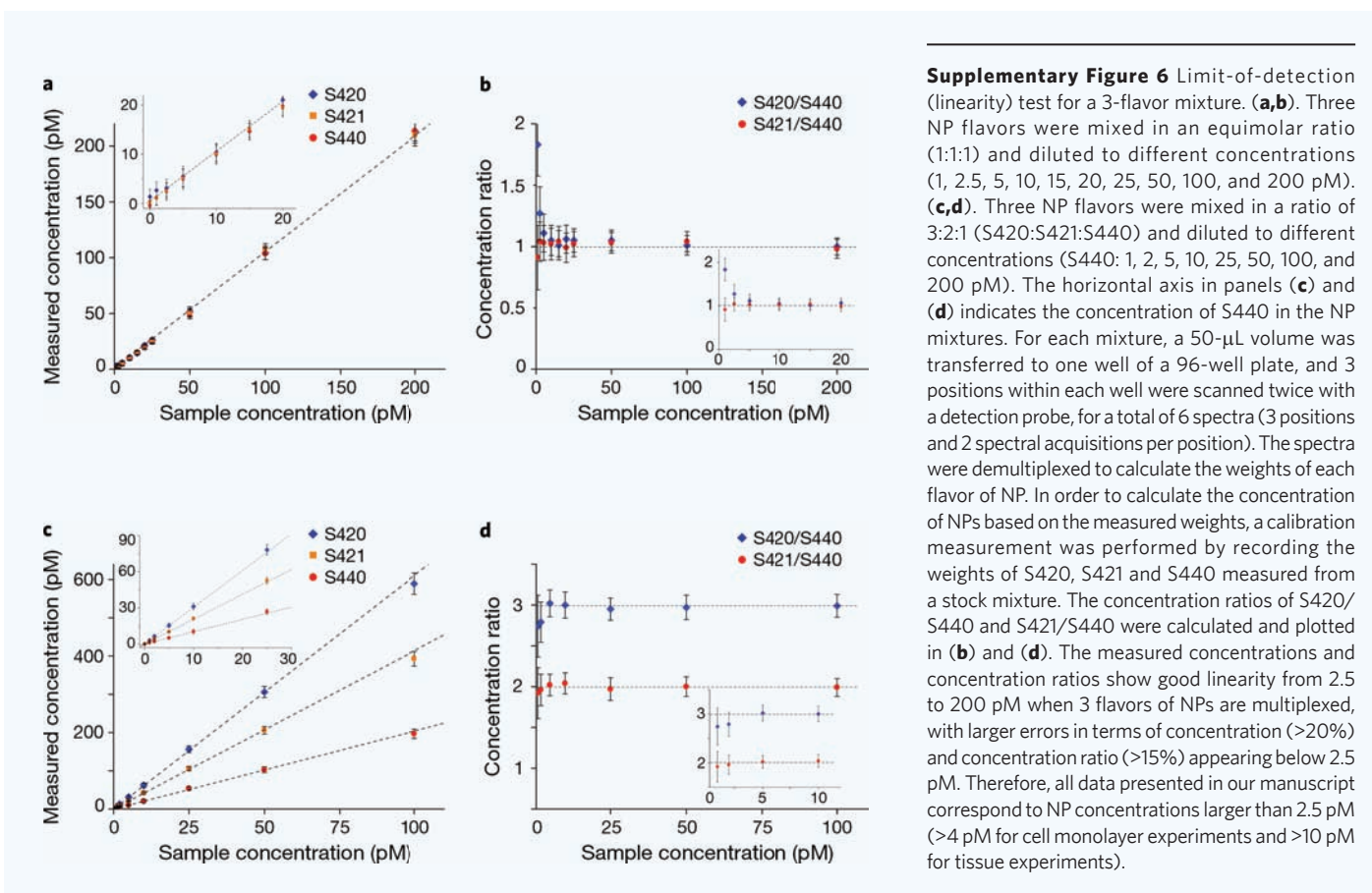
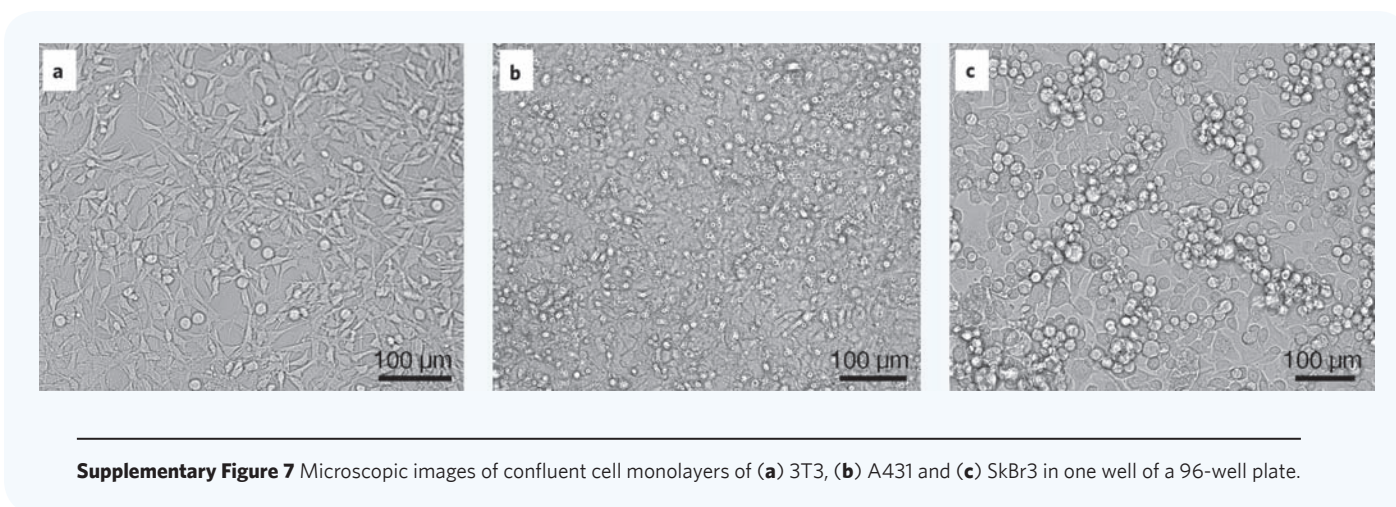


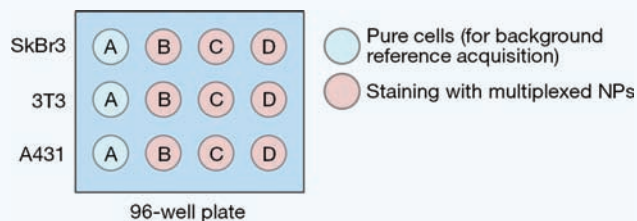
Supplementary Figure 2 UV-VIS spectrophotometry for purifying and calculating the concentration of NP conjugates. (a) The color of NPs and supernatants during purification; (b) absorbance spectra of stock NPs and conjugated NPs for calculation of NP concentration (50 \times dilution); (c) absorbance spectra of supernatant (3 \times dilution; the main component is cyto 647) from 4 rounds of purification.



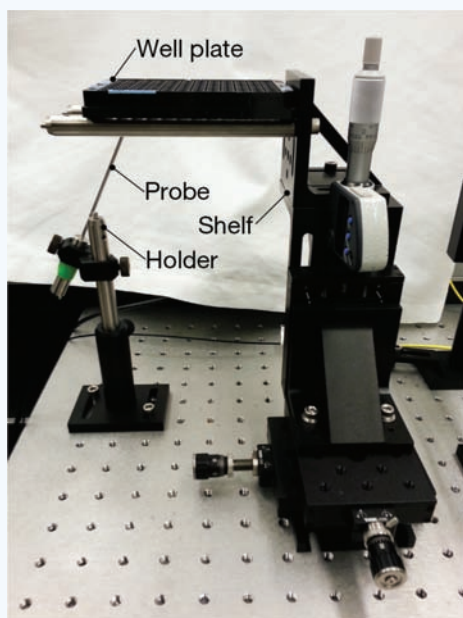
Supplementary Figure 3 Flow cytometry comparison of control NPs with cultured cells. For (a-c), four types of control NPs, labelled with the fluorophore cyto-647, were prepared for this optimization experiment, including: (1) stock NPs that were only labelled with cyto-647, (2) MMPEG-NPs that were conjugated with an excess of MM(PEG)₁₂ to block thiols on the NP surface, and (3) IgG1 and (4) IgG2 isotype-NPs that were conjugated to mouse IgG1 isotype mAb and human IgG2 isotype mAb, respectively, along with MM(PEG)₁₂ (refer to the conjugation protocol). NPs conjugated with anti-EGFR and anti-HER2 monoclonal antibodies were also prepared with the equivalent fluorophore: NP ratio and antibody: NP ratio as the isotype-NPs. The various NPs were individually used to stain (a) 3T3 (-), (b) A431 (EGFR++, HER2+) and (c) SkBr3 (HER2++, EGFR+) cell lines for 15 min (200 pM), followed by flow cytometry analysis. The flow experiment shown in panel a used a higher laser power to compare the behaviors of different control NPs. Although previous reports describe the use of pure (stock) SERS NPs or PEG-modified NPs^{3,4} as controls, the data in panel a suggest that IgG1 and IgG2 isotype-NPs more accurately mimic the nonspecific behavior of EGFR-NPs and HER2-NPs on the control cell line (3T3). In addition, panels b and c show the fluorescence ratio of targeted NPs vs. various control NPs for the tumor cell lines A431 and SkBr3. (d) Fluorescence from unstained cells and NP-stained cells. Non-fluorescent stock NPs (S421) and cyto-647 labeled stock NPs (S421) were individually used to stain SkBr3 cell lines for 15 min, followed by flow cytometry analysis. The results show that since the NPs are encapsulated in silica, there are no surface-enhanced fluorescence effects and that the fluorescence signals are only due to the cyto-647 fluorophores that are conjugated to the NPs.

Observation of NP distribution on cells *via* confocal microscopy

(In vitro) limit-of-detection test**(In vitro) cell monolayer experiments with 96-well plate**

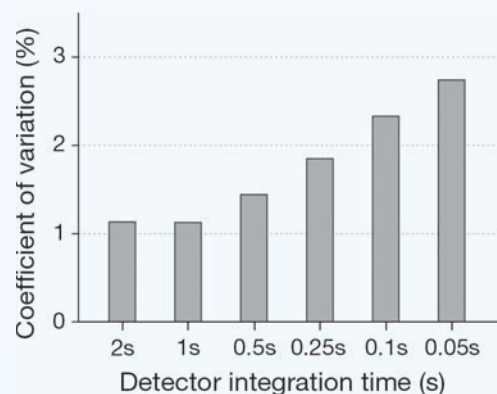


Supplementary Figure 8 Schematic of cell-monolayer experiments on a 96-well plate. For each cell line, cells were counted and seeded into four wells A-D (1×10^5 cells in 50 μ L of media per well for A431 and SkBr3, and 0.5×10^5 cells in 50 μ L of media per well for 3T3), followed by incubation for 24 h to obtain similar cell monolayers (about 2×10^5 cells). Prior to experiments, media was removed and the cells were washed with PBS. In well A, PBS was added to allow for the acquisition of a background measurement in the absence of NPs. For the 3 remaining wells of each cell line (B-D), an equimolar mixture of targeted (anti-EGFR and/or anti-HER2) and nontargeted (isotype control) NPs was added to allow for ratiometric analyses of specific vs. nonspecific binding of the NPs. Both 2-flavor and 3-flavor mixtures were used at a staining concentration of 150 pM per flavor and a total staining volume of 100 μ L. Staining was performed for 15 min followed by up to three rounds of rinsing with PBS. Spectral measurements and ratiometric analysis was performed before and after each rinse step.



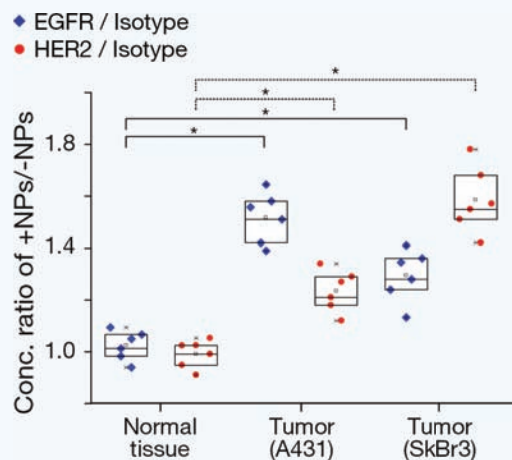
Supplementary Figure 9 Experimental platform for the spectral detection of cell monolayers on a well plate. A probe holder and a shelf were used to maintain the working distance and the illumination and detection angles (probe angle) during measurements.

(In vivo) Effect of detector integration time on measurement uncertainty



Supplementary Figure 10 Coefficient of variation of the measured concentration of NPs on tissues under different integration times. For each data set, 50 acquisitions were taken.

(In vivo) Effect of NP flavor on ratiometric analysis



Supplementary Figure 11 The concentration ratio of targeted and nontargeted NPs typically applied on exposed tissues and measured *in vivo*. Tumors and normal tissues were topical stained with a mixture of 3 NP flavors (three measurements on each of two mice for a total $n = 6$). In these experiments, the NP flavors utilized for each targeted agent are different from those in Fig. 6e. For the experiments in Fig. 6e, S420 NPs were conjugated to an anti-HER2 mAb and S421 NPs were conjugated to an isotype-control mAb. Here, S421 NPs are conjugated to an anti-HER2 mAb and S420 NPs are conjugated to an isotype-control mAb. *P-value < 0.001.

SUPPLEMENTARY REFERENCES

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