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Qualifying label components for effective biosensing by advanced high-throughput SEIRA methodology

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Electronic Supplementary Material available.

The need for technological progress in bio-diagnostic assays of high complexity requires both fundamental research and constructing efforts on nano-scaled assay recognition elements that can provide unique selectivity and design-enhanced sensitivity features. Nanoparticle induced sensitivity enhancement, and its related application to multiplexed capability Surface-Enhanced InfraRed Absorption (SEIRA) assay formats are fitting well these purposes. The potential of diverse fluorophore-antibody conjugates, being chemisorbed to low-cost gold nanoparticulate SEIRA substrates, has been explored with respect to their spectral discriminability. These novel biolabels deliver molecular SEIRA fingerprints that have been successfully analyzed by both uni- and multivariate analyzing tools, to discriminate their multiplexing capabilities. We show that this robust spectral encoding via SEIRA fingerprints opens up new opportunities for a fast, reliable and multiplexed high-end screening in biodiagnostics.

Surface-Enhanced InfraRed Absorption (SEIRA) spectroscopy, bioassays, multiplexing, fluorophore conjugates, principal components analysis (PCA)

### 1. Introduction

Nanotechnology has an increasing impact on both the design and functionality of biomedical and bio-diagnostic devices for reliable high-performance analysis. High-throughput capabilities require the simultaneous detection of various analytes combined with appropriate bioassay components. Additionally, reduction of the biosensors' cost, using for instance low cost nanoparticles (NP) instead of artificial nanostructures is another important aspect. Various bioassay configurations may be used, as the immobilization of targeting elements such as antibodies on glass slides coated with various targeting ligands, as microarrays. To design antibodies assays that function as molecular probes, a selective binding event has to be optimized with respect to the over whole physicochemical properties which may influence the successful readout of signals. This constitutes the core of the first design steps to elaborate multi-purposes bio-assay platforms. Bio-assays are standardized tests usually implemented to support multiple intended uses such as the detection of pharmaceuticals, antigens, hormones, or toxins within a complex sample. However, there is usually a long way to the establishment of a bio-assay setup, which necessitates preliminary steps such as (i) the development of performance-based methods, (ii) the analysis of biological assays and (iii) their validation. This calls for advanced methodologies and readout procedures. Surface-Enhanced InfraRed Absorption (SEIRA) in combination with statistical tools addresses the challenges of accuracy, specificity, precision, and robustness by providing highly resolved vibrational information of complex molecular compounds. Surface-Enhanced InfraRed Absorption (SEIRA) constitutes an ideal platform to isolate vibrational spectroscopic signatures of targeted and active molecules (e.g. antibodies) of the bio-assay, and challenges Surface-Enhanced Raman Spectroscopy (SERS) readout <sup>1, 2</sup>. Dissimilarly to SERS providing sensitive access to Raman-active modes in molecules, SEIRA allows for efficient detection of IR-active modes, thus providing complementary speciation information. Enhancement processes (see comparison in the Supporting Information) of chemisorption to physisorption for SERS and SEIRA, can modify the similarity of the effective speciation sensitivities of both methods <sup>3-5</sup>.

The present work addresses one of the design steps of an assay platform: the targeted biolabels consisting in our case of fluorophore-labeled antibodies chemisorbed onto gold nano-aggregates (Au NA), acting as nano-antennas for biocompatible, low-cost and amplified SEIRA detection <sup>6</sup>, and efficient fast discrimination for proper read-out in tunable and pre-selectable spectral ranges <sup>3</sup>.

Systematic analyses were performed on model antibody Goat Anti-Mouse IgG and fluorophore-conjugated Goat Anti-Mouse IgG such as fluorescein isothiocyanate (FITC), rhodamine(B)isothiocyanate (RITC) and sulphorhodamine B dye (SULFO) fluorophores. These biolabels adsorbed at/nearby NP-functionalised (silanised) surfaces present additional and enhanced spectral signatures due to plasmonic modifications that were compared with those issued from non-chemisorbed gold NA films simply incubated with the label components.

Extensive areas of dried sample films were analyzed with IR/SEIRA spectroscopy and the resulting complex hyperspectral datasets were submitted to automated statistical analysis, namely, principal components analysis (PCA). Relationships and dependencies between chemical functional groups of the various antibody-fluorophore conjugations systems were determined. Consequently, this paper presents profound insights on multivariate discrimination capabilities by direct visualization of derived structural clusters among various biolabels' configurations.

### 2. Material and methods

For further details, please confer to the Electronic Supplementary Material.

# 2.1 Synthesis of Au NP SEIRA substrates and immobilization of Au NP on a MirrIR low-e-substrate

Au NP were synthesized according to a modified protocol of Lee and Meisel<sup>7</sup>, yielding an estimated concentration of  $1 \times 10^{-9}$  M (ca.  $7 \times 10^{14}$  NP/L) and characterized by UV/Vis-NIR spectroscopy (see Figure S-1, Electronic Supplementary Material). For the preparation of SEIRA substrates, the immobilization of Au NP was performed according to the protocol illustrated in ref.<sup>8</sup>, using mid-infrared (MIR) low-e-slides.

### 2.2 Synthesis of fluorophore-labeled antibody conjugates

Different fluorescent dyes that are widely used for biomedical applications were studied. The conjugation of the fluorophore–antibody was performed by covalent linkage between the free residual amino groups of the antibody <sup>9</sup> and the thiocyanate groups and sulfonyl groups of the fluorophores, respectively <sup>10</sup> and further characterized by MALDI-TOF-MS (Figure S-2, Electronic Supplementary Material).

### 2.3 Atomic Force Microscopy

Au NP modified low-e-slides were studied with a NeaSNOM (Neaspec GmbH, Germany) in AFM mode for accessing the surface topography of the functionalized substrates and the existence of immobilized Au NP. Sample areas of 8  $\mu$ m × 8  $\mu$ m and 4  $\mu$ m × 4  $\mu$ m were scanned with a corresponding pixel number of 400 pixels referring to a resolution of 20 nm (8  $\mu$ m-scans) and 300 pixels referring to about 13 nm resolution (4  $\mu$ m-scans). A platinum iridium V coated monolithic silicon AFM tip with a tip radius of curvature with less than 25 nm (NanoWorld AG, Swiss) was used for probing the substrate surface. Image processing (median alignment, pixel column correction) was performed with Gwyddion 2.2.6 (CMI, Czech Republic). The surface topography depicts Au NA and single NP to be immobilized on the glass slide. The Au NP showed spherical morphologies and were in the size range of about 40 ± 3 nm.

### 2.4 SR-based FTIR micro-spectroscopy

### 2.4.1 Instrument and Data acquisition

The measurements were carried on at the IR beamline of the Metrology Light Source (MLS) storage ring with a Vertex-80v FTIR spectrometer <sup>11</sup> coupled to an IR microscope Hyperion 3000 (Bruker Optics GmbH, Germany) equipped with a 128 × 128 pixels FPA (Focal Plane Array) detector (pixel size:~  $3 \mu m$  at 15× magnification).

For FTIR micro-spectroscopy measurements, the SR source ( $\sigma_x$ =670 µm,  $\sigma_y$ =183 µm, beam current ~170 mA) was focused through an IR cassegrain objective of 15 fold magnification onto the sample films. MIR-spectra (from 3900 cm<sup>-1</sup> to 900 cm<sup>-1</sup>) were acquired in transflection mode by co-adding 128 scans at 16 cm<sup>-1</sup> resolution. Background scans were collected before each sample measurement from a region free of sample and rationed against the sample spectrum. The use of synchrotron light has many advantages thanks to its high brilliance, non-invasiveness, coherence and chemical sensitivity compared to conventional IR sources <sup>12</sup> and this study illustrates its applicability to the acquisition of large spectral datasets with FPA.

### 2.4.2 Sample preparation

For the biolabels, fluorophores and antibody-fluorophore conjugates, 2  $\mu$ l of the fluorophoreantibody conjugate solution was added to 20  $\mu$ l of suspended Au NP, resulting in a final concentration of ca. 0.1 mg/ml, respectively. After an incubation time of five minutes, the conjugate solution was dropped-coated onto a low-e-slide, then dried in a desiccator. For comparison, the same amount of conjugate solution without NP was dropped-coated. The NP (20  $\mu$ l) were exposed to 2  $\mu$ l of 10<sup>-4</sup> M of FITC, RITC and SULFO solution respectively, resulting in a final concentration of 10<sup>-5</sup> M. Fluorophores were prepared in DMF and further diluted in sodium bicarbonate buffer (0.1 M, pH~9) to a final concentration of 10<sup>-5</sup> M. The antigen Mouse Anti-Goat IgG was pre-dissolved at a concentration of 0.5 mg/ml in borate buffered saline (100 mM, pH~8.2) and mixed with the colloidal suspensions to a final concentration of 0.05 mg/ml. For Au NA-based SEIRA substrates, FTIR spectra were acquired in absorption mode in the 22000 – 900 cm<sup>-1</sup> spectral range using a MCT (Mercury Cadmium Telluride) detector and InGaAs photodiode (16 cm<sup>-1</sup> spectral resolution, 128 scans). As reference for immobilized Au NP, a slide coated with APTMS for each discrete concentration was measured.

### 2.4.3 Data analysis: uni- and multivariate analysis

Each FPA measurement dataset comprised 16384 spectra, and due to the limited area illuminated by the bright SR beam, contained regions with no MIR signal allowing for correcting the pixel noise and therefore enhancing the signal to noise ratios. The datasets for each of the 8 sample conditions (Goat Anti-Mouse IgG, Goat Anti-Mouse IgG with Au NA, fluorophore, fluorophore with Au NA, fluorophore conjugate, fluorophore conjugate with Au NA, Mouse IgG antibody with/without Au NA) were reduced into matrices of 550 spectra and loaded into Origin 8G (OriginLab Corporation, USA) for polynomial baseline-correction. Each of these spectral dataset is issued from a sample region of ca. 37  $\mu$ m × 96  $\mu$ m, which is adequate to evaluate the potential of a FPA FTIR set-up as bioassay direct readout configuration. Indeed, the dimension of the surface probed here corresponds to biodiagnostically relevant areas delivered by sample systems spotted onto microarrays which range typically between 10  $\mu$ m up to 500  $\mu$ m<sup>13</sup>.

Principal Component Analysis (PCA): The second derivatives of the full dataset of 4400 baseline-corrected absorbance spectra were calculated and consecutively submitted to Savitzky-Golay algorithm (5 smoothing points) and vector-normalization before PCA using MatlabR2011a (The Mathworks Inc., Natick, USA) and PLS toolbox (Eigenvector Research, Inc., USA). The principle components (PCs) were calculated from PC1 up to PC4 and illustrated as scatter plots; such high-order approach been needed for discriminating certain label components.

### 3. Results and discussions

### 3.1 Au NA-immobilization on MirrIR low-e-MIR substrates for SEIRA studies

Self-assembled Au NP multilayers have successfully been implemented as versatile platforms for molecular sensing approaches<sup>14</sup>. Figure 1 displays the spectral features of Au nanoparticulate multilayers that were generated during the drop-coating procedure and analysed in the 22000 - 900 cm<sup>-1</sup> region. Due to aggregation of the Au NP, a red-shift of the plasmon band is induced, and a broader plasmon band appears, extending from the NIR to the

MIR region (compare with Fig. S-1). The position of the extended plasmon band shows that pre-resonant excitation by surface plasmons in the MIR region can be exploited for the SEIRA studies. This is in accordance with previous observations <sup>15</sup>, where a continuum-like resonance for Au nanoshell array samples was observed in the MIR spectral range.

### 3.2 MIR signatures delivered by fluorophore-antibody NP biolabels

Similar to clinical diagnostics, dried films of serum proteins or antibodies are convenient for FTIR investigations. The study of the secondary structure composition, structural dynamics and/or the study of conformational changes (effects of ligand binding, temperature, pH) of the antibodies in liquid phase is not the aim of this work presented here. In the following we discuss the signatures from SEIRA NPs-labeled biolabels (such as antibodies and conjugates) that can be used to visualize and ultimately quantify the bound analyte which is not that obvious or detected in the conventional FTIR configuration. Of notice that it is inherent to the SEIRA technique compared to conventional FTIR is the metal surface influence that may lead to some spectral shapes, shifts and changes in intensities<sup>16</sup>.

### 3.2.1 Goat Anti-Mouse IgG-FITC conjugate

The spectral properties of unlabeled and labeled Goat Anti-Mouse antibody (Fig. 2A and E) were studied. The signature of Goat Anti-Mouse IgG exhibits two strong IR bands at 1535 cm<sup>-1</sup> and 1635 cm<sup>-1</sup> that refer to the Amide II and Amide I bands being characteristic for proteins <sup>17</sup> (Table S-1). In the relatively large wavenumber region (3741cm<sup>-1</sup> to 3278 cm<sup>-1</sup>) the band intensity of the stretching (str.) vibrations corresponds to –NH and –OH groups of amino acid residues <sup>18</sup> which form the secondary structure of the antibody are enhanced in presence of Au NP (see rhombs  $\diamond$ ). This observation corresponds to IR bands that were detected from other protein-Au bioconjugates <sup>19</sup>. The spectrum delivered by the Goat Anti-Mouse IgG incubated with Au NP (Fig. 2B) shows a similar spectral pattern as for 2A. The band at ca. 3278 cm<sup>-1</sup> is attributed to –NH str. vibrations of antibody proteins <sup>20</sup>. The signature of FITC is given in Fig. 2C and exhibits bands referring to str. vibrations (region from 1652 cm<sup>-1</sup> to 1496 cm<sup>-1</sup>) of the xanthene ring moiety and external group modes <sup>21</sup>. The MIR signature delivered by FITC molecules adsorbed onto the Au NP is shown in Figure 2D and comprises str. vibrations of the carbonyl groups of FITC <sup>21</sup> (see asterisks \*). The new occurring band

originate from –C=N and –C=C str. vibrations. This can be explained by different binding orientations which are restricted due to the fluorophore-NP incubation procedure during which the chemisorbed FITC molecules tend to orient. In addition, the origin of SEIRA being observed for Au NP-FITC is due to a formation of large colloidal Au NA, presumably induced by the drop-coating process. The NA chemisorbed FITC molecules orient their dipole moments to be in line with the local electric field of the metal NA, resulting in much larger absorptions than when they are not in line with the oscillating electric field <sup>22, 23</sup>. Such larger absorptions of FITC molecules can be exploited for building up complex conjugates of sensitive multiplexed assay approaches.

An enhancement of the modes in the spectral region between 1304 cm<sup>-1</sup> and 1180 cm<sup>-1</sup>, as well as at ca.1458 cm<sup>-1</sup> can be observed as well (compare Fig. 2C and Fig. 2D). The Goat Anti-Mouse IgG-FITC signature in Figure 2E comprises lines which can be assigned to FITC typical bands (see asterisks \*) and bands between 3720 cm<sup>-1</sup> and 1103 cm<sup>-1</sup> referring to antibody specific signatures. Figure 2F shows the SEIRA spectrum of the antibody-FITC conjugate measured in presence of Au NA. This signature comprises lines which can be assigned to FITC typical bands (Table S-1). For example, in the vicinity of the mode at ca. 1643 cm<sup>-1</sup> (xanthene skeleton str. vibration) of the unenhanced antibody-FITC conjugate now a broad shoulder being located at 1597 cm<sup>-1</sup> appears in the SEIRA spectrum of the antibody-FITC conjugate. This vibration originates from the aromatic –C–C str. vibrations of the xanthenes ring skeleton of FITC molecules which was already observed in the SEIRA spectrum of FITC (compare Fig. 2D). This shows that SEIRA is able to detect with high sensitivity fluorophore molecules in their conjugated states along with their molecular orientation.

Considering the intended use of labeled antibodies in immunoassays, the signatures of the targeting counterpart Mouse Anti-Goat IgG were studied. Figure 2G illustrates antigen specific lines being typical to antibody protein vibrations. The SEIRA signature of the Mouse IgG antibody is displayed in Fig. 2H that shows enhanced modes in the spectral region between 1800 cm<sup>-1</sup> and 900 cm<sup>-1</sup>. Another possible contribution which may explain the observed spectral differences is due to chemical enhancement. Molecules strongly chemisorbed on a (noble) metal surface show a larger enhancement than do physisorbed molecules, suggesting a chemical effect between the molecule and the surface due to strong interaction (Au NA-antibody complexation). This mechanism would describe the case that we discuss and correspond to our antibodies that we detect in presence of Au NA. It is known that the absorption coefficients of chemisorbed molecules are larger than those of condensed

overlayers and therefore interactions between adsorbed molecules may also affect the intensity and thus impact and explain the enhancement of some modes. Charge oscillations between molecular orbitals and the metal surface may enlarge the absorption coefficients of adsorbates <sup>2, 24</sup>.

### 3.2.2 Goat Anti-Mouse IgG-RITC conjugate

In order to study the multiplexing capability of further fluorophore conjugates, RITC, a rhodamine derivative <sup>25</sup>, was used for the coupling. Figure 3B and Fig. 3D display the MIR signatures of RITC and the Goat Anti-Mouse IgG-RITC conjugate spectrum in presence of Au NA. Since both fluorophores FITC and RITC have a quite similar molecular composition their MIR signatures exhibit the same modes, i.e., the bands at 1172 cm<sup>-1</sup> ( $\delta$  C-OH) and at ca. 2941 cm<sup>-1</sup> ( $v_{as}$  CH<sub>2</sub> vibration) are observed in both MIR fingerprints <sup>21</sup>. However, some spectral differences are noticeable. Compared to the spectrum of FITC, the main peak in the RITC spectrum is located at 1589 cm<sup>-1</sup> (Fig. 3A) which can be attributed to aromatic str. vibrations. The enhanced intensities of the vibrational modes referring to RITC are highlighted with rhombs  $\diamond$ . The mode at 1743 cm<sup>-1</sup> can be assigned to a -C(=O)H str. vibration <sup>26</sup>. The SEIRA spectrum of RITC measured on Au NA (Figure 3B) exhibits an enhanced mode at 1589 cm<sup>-1</sup>. It is remarkable that the relative intensity of the str. mode of the xanthene ring moiety at 1342 cm<sup>-1</sup> experiences an enhancement by a factor of five, which proves again the selectivity of revealing one of the most specific molecular groups of the fluorescent dye.

In case of RITC measured on Au NA, further new IR vibrations are observed for instance at 1666 cm<sup>-1</sup>. Further, the mode at 1520 cm<sup>-1</sup> disappears which is caused by a change in molecular orientation. The antibody-RITC conjugate spectrum (Fig. 3C) exhibits RITC-typical modes which are also observed in the bare fluorophore spectrum (Fig. 3A). For instance, the mode at 1589 cm<sup>-1</sup> originates from aromatic –C–C str. vibrations of RITC molecules <sup>26</sup>. In presence of Au NA, the conjugate signature (Fig. 3B) delivers one broad band at 1589 cm<sup>-1</sup> and the mode at 1647 cm<sup>-1</sup> is observed as a shoulder being located at ca. 1639 cm<sup>-1</sup> (Table S-2).

### 3.2.3 Goat Anti-Mouse IgG-SULFO conjugate

Figure 4B and 4D shows the signatures of the SULFO and SULFO-labeled Goat Anti-Mouse antibody, in presence of Au NA. Compared to the unenhanced signature of SULFO (Fig. 4A), the mode referring to the aromatic –C–C str. vibration <sup>27</sup> at 1581 cm<sup>-1</sup> is enhanced, as the line at 1597 cm<sup>-1</sup> is shifted to shorter frequencies. Further modes originate from –C–O str. vibrations of the carbonyl group and –C–C str. vibrations nearby the xanthene ring moiety <sup>28</sup>. The SEIRA signature of SULFO (Fig. 4B) shows structural differences in terms of band shifts and its spectral behaviour in proximity of Au NA is quite similar to FITC (compare Fig. 2D). Further, new bands appear in the long frequency region, for example lines at 3186 cm<sup>-1</sup> referring to –NH str. vibrations at 3325 cm<sup>-1</sup> can be observed along with str. vibrations at 3780 cm<sup>-1</sup> (–OH) <sup>29</sup>. In the antibody-SULFO conjugate spectrum the band at 1597 cm<sup>-1</sup> is red-shifted. However, conjugate typical bands appear at 1412 cm<sup>-1</sup> along with modes at 1034 cm<sup>-1</sup>, 1242 cm<sup>-1</sup> originating from SULFO. The spectrum of the antibody-SULFO conjugate measured in presence of Au is very similar to the conjugate spectra measured without Au NA (cf. to Table S-3).

The univariate study of these different antibody-fluorophore conjugate signatures shows that the electric field enhancing capabilities differ between Au NP and the type of antibodyfluorophore conjugate which justifies the systematic approach on the spectral properties of such fluorophore-antibody NP complexes. Additionally, this shows that for the bioassay design, SEIRA induced by NP provides different vibrational information from fluorophoreanibody NP labels that can be used for building up spectral databases.

Summarising the results, FTIR spectroscopy combined with SEIRA are important analytical tools for studying the biochemical composition of proteins such as antibodies. Bands of high interest for biological purposes and labeling procedures can be found in the Amide I/II and ester region. The Amide I region is traditionally exploited for probing the proteins secondary structures, as an indicator band for monitoring spectral changes due to alterations in molecular structure, usually due to external perturbations. Our studies show that complex biolabels that are readout by SEIRA deliver distinct spectra that clearly differ from their unenhanced counterparts: the conventional FTIR spectra (please confer to Figure S-3, and Tables S-1–S-3, Supplementary information).

### 3.3 Multivariate discrimination of the signatures of label components

PCA as a powerful chemometric method for finding patterns in complex FTIR datasets enables fast identification and separation of labeled antibodies which are designed for multianalyte detection <sup>30 31</sup>. Figure 5 shows scores plots (PC1-PC4) that entail spectral differences. All PCA procedures displayed in Fig. 5A-C (and Fig. 6b) were performed by repetitive experiments (data not shown here, 95% reproducibility).

The PCA displays datasets of the Goat Anti-Mouse IgG that are well separated from the antigen data in PC1 against PC2. Results from PCA confirm that averagely 85 % of the variance of the spectral datasets, i.e. the discrimination capability up to the third order, could be explained for the fluorophores. Additionally, we decided to put on display all components up to the fourth order to illustrate the options for discriminating different label and/or label components. The scores (PC1 against PC2) of Goat Anti-Mouse IgG are positive and lie in the second quadrant of the variance-weighted room. In PC1 against PC3, the Goat Anti-Mouse IgG antibody data are well separated from the unenhanced spectra. The spectral differences observed in the SEIRA spectrum of FITC (see Fig. 2D) are also reflected by PC1 against PC2. Figure 5A illustrates that in PC1 against PC2 FITC is separated from the FITC-labeled antibody. With respect to the design of multiplexed immunoassay formats, this shows that the enhancement capabilities of different Au NA labels provide MIR signatures with different spectral patterns. PCA for studying the multiplexing capability of the other fluorophore conjugates containing RITC molecules was also performed successfully. Similar to the PC scores illustrated in Figure 5A, a discrimination of the RITC MIR signatures measured on Au NA in PC1 against PC2 was feasible. A variance of about 80 % is captured by the components PC1-PC3. The Goat Anti-Mouse IgG-RITC conjugate cluster has quite similar distance towards its unlabeled counterpart as the FITC-antibody conjugate, which reflected by the good separation in PC1 against PC2.

Compared to all other fluorophore-antibody conjugates discussed so far, the SEIRA signature of the FITC-labeled Goat Anti-Mouse IgG is most similar to the unlabeled antibody sample; for SULFO the distance in PC1 against PC2 is the highest which shows the increase of discrimination capability of the conjugated fluorophores.

The multivariate analysis on spectral features is an important scope for the design and development of bioassay platforms, as functionalization with biolabel components may require several preparation steps before the establishment of a functionally completed assay setup.

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# 3.4 MIR signatures of the antibody-FITC conjugate measured on the Au NP modified low-e-substrate

Figure 6a shows the extinction spectrum and surface topography of Au NP being immobilized on an APTMS-functionalized substrate (1% APTMS, time of incubation: 24 h, see Fig. 6a, A-C). A second extended plasmon band is observed in the MIR region and proves the existence of Au NA <sup>32</sup> as for the unmodified glass substrate. This is important in case of the enhancement capabilities of immobilized NP as large NA reaching microscale size dimensions of bulk materials might drastically lose their enhancement properties.

The PCA results confirm that averagely 85 % of the variance of the spectral datasets up to the third order could be explained. PCA of the Mouse Anti-Goat IgG signatures shows a better separation from the enhanced ones (PC1 against PC2, PC3 in Fig. 6b). The MIR signatures are illustrated in Figure 6c. Compared to the FITC signature, spectral differences in the enhanced counterpart can be observed, i.e., the band at 1738 cm<sup>-1</sup> (compare with Fig. 3) is split into lines at 1726 cm<sup>-1</sup> and 1752 cm<sup>-1</sup>. The dependence on the substrate configuration and modification is very striking when comparing the antibody-FITC conjugate being adsorbed on the Au NA on the APTMS-modified slide: the signature exhibits FITC typical bands along with antibody modes (Fig. 6c, C). Compared to the conjugate signature in Fig. 2 where a mode at ca. 1643 cm<sup>-1</sup> can be observed, now a new band at 1681 cm<sup>-1</sup> (-C=O str.) appears after APTMS modification. This shows that the SEIRA signature of FITC and its antibody conjugate differs from the signatures measured with the APMS-modified substrate and the assay configuration itself. It should be noted that the vibrational modes in the 1600 cm<sup>-1</sup> to 1300 cm<sup>-1</sup> range can be assigned to both aminopropyl and free aminogroups as well. However, no peak at 1491 cm<sup>-1</sup> associated with NH<sup>3+</sup> in the SiO<sup>-</sup>–H–NH<sup>3+</sup> group of APTMS residues <sup>33</sup> is noticeable which shows that this substrate enables the overlap-free detection of amino-group modes delivered by the targeting units itselves.

A tremendous impact of this substrate with respect to its enhancement capability is shown for the antigen Mouse Anti-Goat (Fig. 6c, F) in the region from 1460 cm<sup>-1</sup> to 1300 cm<sup>-1</sup>. In particular, the mode at 1641 cm<sup>-1</sup> is extremely intensified by a factor of about 8. This is in accordance to the calculated dimension of the average enhancement factor for the MIR spectral region as described in ref. <sup>34</sup>.

### 4. Conclusions

We demonstrate the efficiency of the SEIRA technique to probe the molecular characteristics of different antibody-fluorophore NP conjugations considered in this study. Our measurements show that the spectral signatures of the native antibody Goat Anti-Mouse IgG and the conjugated fluorophores are preserved in the proximity of Au NPs but enhanced by the latter. This implies that the elaboration of specific assay platforms spectral databases can be envisaged with SEIRA. Coupled with PCA analysis, this method allows clear discrimination of the functionalized systems, which ascertain their potential for multiplexed bioassay applications and biosensor development of high efficiency with customized responses. Our work highlights the prospective use of NP-assays for biodiagnostics <sup>35, 36</sup> and as catalysts for enhancing selectively bio-physiological responses of cells or bacteria. Consequently SEIRA applications <sup>37</sup>, can be expected to show even higher performance characteristics than the one of conventional / established methods such as SERS <sup>38</sup>.

In such, a logic extension of our work was to confront the potentialities of SEIRA to discriminate biolabels' spectral features in a real biological environment, which is subject of another communication.

Altogether, this paper evidences the availability in the complex bio-diagnostic micro-assays toolbox of a novel mean to identify the biolabel' systems suitability along with their discrimination potential and their multiplexing capability. Enhancing functionality performances is the key point to obtain higher and targeted screening performance, ultimate goal of complex bio-diagnostic assay platforms <sup>39</sup>.

### Acknowledgements

We thank Ralf Bienert (BAM Federal Institute for Materials Research and Testing) for use of the DLS apparatus. The machine-physicists team of the Helmholtz-Zentrum Berlin (HZB) is gratefully acknowledged for tuning the MLS according to our experimental needs. We thank Arne Hoehl of the MLS team of the PTB for helpful discussions and use of the beamline instrumentation. Sincere thanks are given to Sören Selve (ZELMI, TU Berlin) for the preparation of the TEM images.

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Absorbance spectrum of Au NA showing an extended plasmon band in the mid-infrared region used for the SEIRA experiments. Analysis was performed in the spectral range from 22000 cm<sup>-1</sup> to 900 cm<sup>-1</sup>. 125x87mm (300 x 300 DPI)





MIR spectra of (A) Goat Anti-Mouse IgG, (B) Au NA incubated with Goat Anti-Mouse IgG, (C) FITC, (D) Au NP and FITC, (E) Goat Anti-Mouse IgG-FITC, (F) Au NP and Goat Anti-mouse IgG-FITC and the antigen (G) Mouse Anti-Goat IgG and (H) Au NP and Mouse Anti-Goat IgG measured with SR. Vibrations of FITC are marked with asterisks \*. Enhanced vibrations are marked with rhombs  $\diamond$ . Further details: cf. to Table S-1. 190x320mm (300 x 300 DPI)



MIR spectra of (A) RITC, (B) Au NP and RITC, (C) Goat Anti-Mouse IgG- RITC, and (D) Au NP and Goat Antimouse IgG- RITC measured with SR. Vibrations of RITC are marked with asterisks \*. Enhanced vibrations are marked with rhombs ◊. Further details: cf. to Table S-2. 122x124mm (300 x 300 DPI)





MIR spectra of (A) SULFO, (B) Au NP and SULFO, (C) Goat Anti-Mouse IgG- SULFO, and (D) Au NP and Goat Anti-mouse IgG- SULFO measured with SR. Vibrations of SULFO are marked with asterisks \*. Enhanced vibrations are marked with rhombs ◊. Further details: cf. to Table S-3. 131x124mm (300 x 300 DPI)



PCA illustrated from PC1 to PC4. PCA was performed with vector-normalized second derivative spectra over the spectral range from 3900 cm<sup>-1</sup> to 900 cm<sup>-1</sup>. PCA clusters are color-coded as Goat Anti-Mouse IgG (violet), Au NP and Goat Anti-Mouse IgG (red), fluorophore (green), Au NP and fluorophore (cyan), Goat Anti-Mouse-fluorophore (magenta), Au NP and Goat Anti-Mouse-fluorophore (black), Mouse Anti-Goat IgG and NP (yellow) and Mouse Anti-Goat IgG with Au NP (blue). 293x564mm (300 x 300 DPI)



(a) Absorbance spectrum of Au NP immobilized on an APTMS-functionalized substrate (cAPTMS~1%, incubation time: 2 h) and corresponding surface topography (scale bars for (A): 0.1  $\mu$ m, (B) and (C): 0.05  $\mu$ m). Au NA are visible as spherical shaped clusters. (b) PCA analysis (3900 cm<sup>-1</sup> – 900 cm<sup>-1</sup>). (c) MIR signatures of (A) FITC, (B) FITC on the APTMS-substrate, (C) Goat Anti-Mouse IgG-FITC and (D) Goat Anti-mouse IgG-FITC on the APTMS-substrate and the antigen (E) Mouse Anti-Goat IgG and (F) Mouse Anti-Goat IgG on the NP-modified APTMS substrate measured with SR. Vibrations of FITC are marked with asterisks \*. Enhanced vibrations are marked with rhombs  $\diamond$ . 177x152mm (600 x 600 DPI)