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Towards improved quantitative analysis using surface-enhanced Raman scattering incorporating internal isotope labelling†

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Raman spectroscopy has attracted considerable interest during the past two decades as a vibrational technique used for the molecular characterisation of different molecules. Whilst the Raman effect is known to be generally weak, it is also known that this can be greatly improved using surface-enhanced Raman scattering (SERS). Indeed, in recent years, the power of SERS for rapid identification and quantification of target analytes in a wide range of applications has been repeatedly demonstrated in multiple studies. Moreover, the application of SERS in combination with an isotopically labelled compound (ILC), as an internal standard, has also very recently shown promising results for quantitative SERS measurements, by improving both its accuracy and precision. This is due to the ^{12}C and ^{13}C or ^1H and ^2H (D) having similar physicochemical properties. The use of these internal standards results in the reduction of any influences due to the number of nanoparticles within the analysis zone and fluctuations in laser fluence. Thus, in this study we have employed SERS for quantitative detection of tryptophan (Trp) and caffeine. These have been chosen because Trp is readily available as the deuterated form and caffeine is available in both ^{12}C and ^{13}C . Quantum chemical calculations based on density functional theory (DFT) have been utilized to determine the vibrational characteristics of the target analytes. For SERS analysis incorporating isotopologues of tryptophan three independent experiments were conducted with three different batches of nanoparticles over a 12 month period; our results show that the use of this internal standard improves quantification of this target molecule. In particular for the independent test sets (i.e., samples not used in quantitative partial least squares regression (PLSR) model construction) we observed improvements in the linearity for test set predictions, as well as lower errors in test set predictions, when isotope internal standards were used during SERS for both deuterated tryptophan as well as ^{13}C caffeine. This work is an extension of and a natural progression from our earlier studies. By exploring additional analytes of interest, allowing for the assessment of the different types of stable isotopes as internal standards, and demonstrating the transfer/robustness of isotopologues for use with SERS, we believe this approach could be readily extended to other biologically-relevant compounds.

Introduction

Raman spectroscopy is a powerful technique providing biochemical fingerprints of the sample under investigation.^{1,2} The Raman effect has generally been associated with weak signals, which can be said to have been the driving force behind the continued development of Raman techniques such as resonance Raman spectroscopy (RRS) or surface-enhanced

Raman scattering (SERS) to improve the signal response of conventional Raman scattering. Thus, issues concerning low sensitivity and reproducibility are being overcome by these new techniques and approaches. It is generally accepted that SERS improves the Raman signal through the adsorption (or very close proximity) of species onto a roughened metal substrate.³⁻⁵ There is a perception within the literature that SERS suffers from poor reproducibility (as reviewed in ref. 6). For colloidal-based SERS, this may be due to the number of nanoparticles within the collection volume and unavoidable fluctuations in laser power (fluence). In this study isotopic labelling (IL) strategies are employed to improve this quantitative method further in terms of efficiency, reproducibility and linearity. The IL approach is based on the addition of a known constant quantity of an isotopically-labelled analog (isotopologue or spike) of the target analyte to the original sample

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as an internal standard, prior to any further treatment. As the isotopolog is chemically identical (with the obvious exception of the isotope substitution) the IL-SERS approach combines high sensitivity and molecular specificity with the asset of reliability, accuracy and repeatability through quantitative analysis. Although isotope labelling mass spectrometry (ILMS) has been applied in different areas of analytical chemistry, as a quantitative detection method to provide high accuracy reference measurements,⁷⁻⁹ compared to IL-SERS this approach requires complex sample preparation, while the instrumentation cost is also relatively high. The IL approach has also been carried out by Zakel and Stosch to develop a quantitative method for measuring creatinine as a biomarker in human serum.¹⁰ This study concluded that the combination of SERS and IL had proved to be a valuable approach for clinical diagnostic measurements.

Tryptophan (Trp) is an essential amino acid required for the biosynthesis of proteins and functions as a precursor of active substances such as serotonin.¹¹ However, a reduction in the level of Trp may cause pellagra, which is niacin-Trp deficiency disease with symptoms of dermatitis, diarrhea, dementia, and mood-alterations. The change in brain Trp levels also leads to changes in brain serotonin synthesis.¹² A recent study has also shown significant changes in Trp levels concomitant with aging.¹³ Trp is comprised of two active functional groups; carboxylic (-COOH) and amino (-NH₂), the interaction of carboxyl and amino groups on SERS has been investigated previously, as strong vibrational enhancement is generated from these two groups.¹⁴ Several studies have also investigated the SERS activity and the interaction of different metal surfaces with amino acids.¹⁵⁻¹⁷ These studies have also demonstrated that Trp molecules may interact with the surface through COO and NH₂ groups.

Caffeine (1,3,7-trimethylxanthine), is a typical alkaloid related to the purine alkaloids, which is also a natural compound widely found in coffee beans, tea leaves, cola nuts and several plants, it is also added to soft drinks as a flavouring agent¹⁸ and many pharmaceutical products. The international Olympic Committee (IOC) has classified caffeine as an abuse drug when present in urine at level of >12 mg mL⁻¹.^{19,20} The overconsumption of caffeine can lead to adverse effects on human health such as increase of gastric-acid secretion, heart disease (cardiac arrhythmia), kidney malfunction and disorder of the nervous system.²¹ Therefore, accurate quantitative analysis of drugs is important in order to establish long-term abuse of illicit compounds, as well as establish accurate drug dosing for legal therapeutics. However, analysing the levels of drugs found in human bio fluids requires methods that are highly sensitive and reproducible, as well as the additional potential for portability and point-of-care diagnostics.

Detection and quantification of Trp and caffeine has improved through the use of multiple analytical techniques such as chromatography coupled with mass spectrometry.^{19,22,23} Whilst these methods generate a high level of qualitative and quantitative results, they are not, to date,²⁴ generally considered field portable, and require specialist operatives as well incurring a high analysis cost. By contrast, recent studies have

demonstrated the application of SERS for the detection and quantification of Trp and caffeine with high levels of sensitivity and reproducibility.^{25,26} Alharbi et al. concluded that SERS can be used to identify long-term abuse of illicit materials as well as determine accurate drug dosing for legal therapeutics. Furthermore, Chen and colleagues reported the interaction of caffeine molecules with different colloids at various concentrations.²⁷ Finally, in a very recent study, we have also demonstrated the effect of isotopic labelling (using deuterated codeine) for improved quantitative detection of codeine in human plasma using SERS.²⁸

The current work is a follow on from the mentioned work above, that was recently published in *Analyst*,²⁸ where in the present study we aim to investigate stable isotopes of hydrogen and carbon: the isotopolog for Trp was deuterated (Trp-d₅) and for caffeine all three methyl groups were replaced with ¹³CH₃. In this study we show that SERS can be successfully developed as a highly quantitative method for the detection of Trp and caffeine via adoption of the IL-SERS principle combined with multivariate statistical analysis techniques. In this study we also perform repeat measurements where the laser power is 'degraded' to show transfer/robustness of the isotopologue concept. Our results clearly demonstrate the potential applications of this strategy as a quantitative approach for different compounds, which could provide increased accuracy and precision and which have the potential to benefit a wide range of analytical fields.

Experimental

Chemicals

Gold(III) chloride trihydrate (99.9%), trisodium citrate, silver nitrate (99.9%), high purity Trp (99%) and caffeine were purchased from Sigma Aldrich (Sigma-Aldrich, Dorset, UK). Isotopically labelled Trp-d₅ (98%) and caffeine (trimethyl-¹³C₃, 99%) were purchased from Cambridge Isotope Laboratories, Inc. (Tewksbury, MA, USA). Chemicals were used as supplied and all solvents used were of analytical grade.

Nanoparticle synthesis and characterisation

The Turkevich method was used to synthesise gold nanoparticles.²⁹ A 100 mL volume of 50 mg HAuCl₄ solution was added to 850 mL of boiling water, and while the solution was vigorously stirred 50 mL of trisodium citrate was added. After 30 min the solution had changed to a deep red colour and was left to cool.

The Lee and Meisel citrate reduction method was used to produce silver nanoparticles,³⁰ and to remove any trace metals from the glassware, aqua regia (nitric acid : hydrochloric acid (1 : 3, v/v)) was used for cleaning. The vessels were washed using cleaning detergent and rinsed with water prior to being oven dried at 50 °C for 25 min. After dissolving AgNO₃ (90 mg) in 500 mL of water, this was heated to boiling point. Then 1% trisodium citrate (10 mL) solution was added, and stirred gently at a steady boil for 15 min. When the solution started to change

to a milky green colour, the end point of the reaction was reached.

See ESI† for more details of UV-vis and SEM instrumentation. UV-vis spectrophotometry was used to characterize the nanoparticles size distribution to allow comparison of several batches and data generated were similar to those data collected previously by our group.^{31–33} The λ_{max} of nanoparticles were identified to be 534, 563 and 527 nm for gold and silver respectively (Fig. S1a and b†). Moreover, scanning electron microscopy (SEM) was used to assess the nanoparticles in terms of their morphology (Fig. S1c and d†), and the average size of the nanoparticles for AuNP and AgNP was found to be 16.12 nm and 83.26 nm respectively.

Stock solutions and internal standard preparation

Aqueous stock solutions of natural isotopes containing Trp and caffeine (10 mL each), along with the correspondingly labelled internal standard compounds, Trp- d_5 or ^{13}C caffeine, were prepared in water. Mixture dilutions of the unlabelled compounds were prepared over a range of concentrations for Trp and caffeine from 0–90 mM. The final concentrations of the internal standards spiked into unlabelled samples were 20 mM. Calibration curves for each of the analytes were constructed using the above mixtures.

Colloidal-based SERS in solution

Spectra were collected using a DeltaNu Advantage benchtop Raman spectrometer (DeltaNu, Laramie, WY, USA), equipped with a 633 nm HeNe laser with a power output of 3 mW on the sample. Spectra were collected for 30 s over a range of 200–3500 cm^{-1} . Samples were placed in an 8 mm diameter glass vial and subjected to laser irradiation once loaded into the sample cell attachment. The instrument was calibrated using a poly-styrene internal standard as supplied by the instrument manufacturer to determine the ideal distance from the laser to the glass vial. SERS sample preparation and analysis was performed as follows: 200 mL of colloid was added to a glass vial followed by 200 mL of Trp – Trp- d_5 or caffeine – ^{13}C caffeine, along with 50 mL aggregating agent. The vial was then vortexed for 2 s and inserted into the sample cell attachment where a spectrum was acquired immediately.

Data analysis

The molecular geometry and vibrational frequency calculations were carried out using the Gaussian 09 package.³⁴ Structures were constructed by hand and optimised using density function theory (DFT) with the B3LYP functional^{35,36} utilising the cc-pVTZ basis set.³⁷ Vibrational frequencies and Raman intensities were calculated at the same level of theory for geometry optimisation. The assignments of Raman bands for Trp and caffeine with their labelled molecules were based on gas phase calculations.³⁸ Visualisation of vibrational modes and generation of Raman spectra was carried out using Gaussview 5.³⁹

For chemometric analysis Raman spectra were analysed using MATLAB software R2013a (The Math Works Inc, Natick, USA). First, a set of univariate regression models were

established by the regression of the integrated areas of the discriminatory peaks of interest, against the known concentration of the target analytes (Trp and caffeine). A further set of univariate regression models was then established by replacing the integrated peak areas of the analytes with the peak area ratios of the analytes to the internal standards (analyte/internal standard).

Finally, a series of multivariate regression models were also constructed using partial least square regression (PLSR).^{40,41} These three models used the entire Raman spectra for quantification of the two analytes with, and without, the Trp- d_5 or ^{13}C caffeine isotopes as internal standards. The generated PLSR models were validated by a projection approach,^{42,43} where 60% of the data were used as the training set to generate the model (concentrations 0, 20, 40, 50, 70, and 90 mM Trp or caffeine) while the remaining 40% were projected as a test set (10, 30, 60 and 80 mM) for validation purposes. A set of calibration figures of merit were calculated for the models based on errors: root-mean-square error of cross validation (RMSECV) for the training set and root-mean-square error of prediction (RMSEP) for the test set. In addition, coefficients of determination were calculated for both the training sets (R^2) and test set (Q^2).

Results and discussion

The application of IL coupled with conventional Raman spectroscopy has focussed considerable attention on the classification⁴⁴ and quantification of bacterial metabolism,⁴⁵ as well as being utilised for the characterization of various bioprocesses.⁴⁶ This study is an extension and natural progression of our earlier study, where we focused only on the deuterated molecule codeine spiked into the human plasma.²⁸ In addition, our previous study only used a single batch of nanoparticles, whereas the current study is investigating the effects of isotope standards using either deuterium (tryptophan) or the ^{13}C labelled molecule caffeine, and also includes multiple batches of SERS substrates which were run on different days (a year apart).

Preliminary Raman analysis (see ESI† for more details of the conventional Raman measurements) of Trp and Trp- d_5 and ^{12}C and ^{13}C caffeine were carried out to identify vibrational spectral shifts due to changes in reduced mass of various functional groups containing either D (^2H) or ^{13}C . Comparison of the Raman spectra of labelled and unlabelled Trp (Fig. S2a†) revealed the main bands affected by the labelling process to be peaks at 844, 1050, 1189 and 1597 cm^{-1} , corresponding to phenyl ring CD bending, indole ring CD (NH) bending, phenyl ring breathing and indole ring CD (NH) bending respectively.⁴⁷ These band assignments have been based on DFT calculations and Table 1 contains a full list of all other detected peaks and their corresponding band assignments. A similar approach was also employed to identify and compare the differences between Raman spectra of labelled and unlabelled caffeine (Fig. S2b†). The high intense peak at 561 cm^{-1} is attributed to CH bending, CH_3 symmetric bending and CH_3 rocking vibrations which displayed a noticeable shift in the Raman spectrum of labelled caffeine to 548 cm^{-1} . All other detected peaks and their respective

Table 1 Experimental and calculated Raman and SERS bands (in cm⁻¹) with vibrational assignments for Trp and Trp-d₅

Raman-Trp	Calculated	Raman-Trp-d ₅	Calculated	SERS-Trp	SERS-Trp-d ₅	Vibrational assignment
595	599	583	586	—	—	Indole out-of-plan bend, indole ring CH (NH) out-of-plane bend, CH ₂ wag
754		702	714	753	—	Indole ring bend, phenyl ring CH bend, NH ₂ wag, CH ₂ rock
871	893	820	837	862	850	CH ₂ rock, NH ₂ wag, indole ring CH
1006	1033	844	863	1044	—	Phenyl CH bend, phenyl ring breathing
1118	1135	1050	1064	1122	1047	Ind ring CH (NH) bend, pyrrole ring CN str
1207	1218	1153	1164	—	—	Phenyl ring CH bend, pyr ring NH bend
1231	1245	1189	1199	1228	1188	Phenyl ring breathing, pyrrole ring CN str, indole ring CH (NH) bend, CH ₂ wag
1357	1371	1334	1356	1350	—	Indole ring bend, indole CH (NH) bend, NH ₂ rock, CH _{alkyl} bend
1423	1450	1406	1424	1441	—	Indole ring bend, indole CH (NH) bend, CH ₂ rock
1575	1583	1551	1553	—	—	Indole ring CC str, indole ring CH (NH) bend
1616	1655	1597	1634	1600	1588	Indole ring CH (NH) bend, phenyl ring CC str

vibrational assignments are listed in Table 2.^{27,48,49} As caffeine is only labelled through the CH₃ functional groups, the Raman peaks related to the indole ring were not significantly affected by the labelling process. However, this could perhaps be investigated in more detail by labelling the nitrogen atoms of the ring in future studies.

Optimisation of SERS parameters is generally necessary to reach the optimal conditions and reproducibility and achieve signal enhancement.^{6,50} It is well known that the sensitivity of SERS depends on the binding of molecules with a surface as

well as the presence of an aggregation agent.⁵¹ Therefore, in this study four different colloids including silver citrate (SC), gold citrate (GC), hydroxylamine silver citrate (HASC) and sodium borohydride-reduced silver (SBRS), have been examined and compared. In general, batch-to-batch variation and reproducibility of SERS is a key discussion point within the Raman community. Therefore, multiple batches of nanoparticles (n = 3) were prepared and used separately over a 12 month period, in order to investigate the efficiency of the isotopic labelling approach towards the quantitative detection of the analyte of

Table 2 Experimental and calculated Raman and SERS bands (in cm⁻¹) with vibrational assignments of unlabelled and ¹³C labelled caffeine

Raman unlabelled caff	Calculated	Raman labelled caff	Calculated	SERS unlabelled caff	SERS labelled caff	Vibrational assignment
447	445	445	441			CH ₃ sym. bend + CH ₃ rock
487	483	482	482	506	496	Imidazole ring def + O]C-N def + CH bend
561	555	548	546			CH bend + CH ₃ sym. bend + CH ₃ rock
643	647	643	642	646	646	Imidazole ring def + O]C-N def + CH ₃ sym. bend
740	751	737	744	746		Imidazole ring def + CH ₃ sym. bend + CH ₃ rock
803	813	803	811			Imidazole ring def + C]O def + CH ₃ rock
931	939	926	932	940		Imidazole + CH ₃ rock
1026	1032	1021	1028	1006		CH ₃ rock + C-N str + C-N bend
1079	1088	1073	1081	1075		H-C]N bend
1140	1155	1128	1143	1169	1131	N-CH ₃ asym. bend
1241	1265	1239	1259	1250	1250	C-N str
1289	1299	1287	1297	1328	1325	C ₂ -N ₃ sym. str + CH ₃ rock
1332	1354	1328	1348			Imidazole trigonal ring str
1365	1383	1359	1379	1372	1363	C-N str + CH ₃ sym. bend
1412	1451	1406	1438	1416	1413	CH ₃ sym. bend + C ₁₄ -N ₇ sym. str
1556	1571	1556	1570	1584	1575	CH ₃ sym. bend + CH ₃ bend + C-N str
1603	1619	1603	1617			C]C sym. str + C-N sym. str + CH ₃ sym. bend
1657	1670	1657	1668	1681		Out-of-phase C]O str + C]C sym. str
1700	1722	1700	1720			In-phase C]O str

interest. The results indicated that gold citrate and silver citrate provided the highest reproducibility and stability for Trp (data not shown) and caffeine (Fig. S3†) respectively. Moreover, potassium nitrate, sodium chloride and potassium sulfate solutions (0.5 M) were also examined in order to identify the optimal aggregating agent for the analytes used.

The highest SERS intensity for Trp was observed when KNO_3 was added to Au-citrate (Fig. S4†), which was used for all further studies. Furthermore, in order to establish the influence of the aggregation time, Trp peak area at 1122 cm^{-1} (C–H vibration) both with and without the IL internal standard was monitored as a function of time and recorded through 78 time-points for 39 min. We observed a rapid initial aggregation after addition of the salt (KNO_3) which gave the optimum SERS response

(Fig. S5a†). Moreover, mixtures of Trp with Trp-d5 remained unchanged for a period of time 15 min compared to Trp alone, which supports our assumption that the isotopologue associates with the metal nanoparticle surface with the same efficiency as its unlabelled equivalent. The intensity is approximately half because the concentration of Trp has been diluted to 50% when spiked with the isotope (CH/CD) (Fig. S5b†).

SERS spectra of unlabelled and labelled Trp obtained using gold colloids are displayed in (Fig. 1a). These spectra show successful binding of Trp and Trp-d5 to the surface and are similar to other previous studies that investigated unlabelled Trp.^{25,52} It is clear that some of the bands are shifted to lower wavenumbers due to the presence of the heavier isotopes of Trp-d5. Peaks at 1122 , 1228 and 1600 cm^{-1} , assigned to phenyl

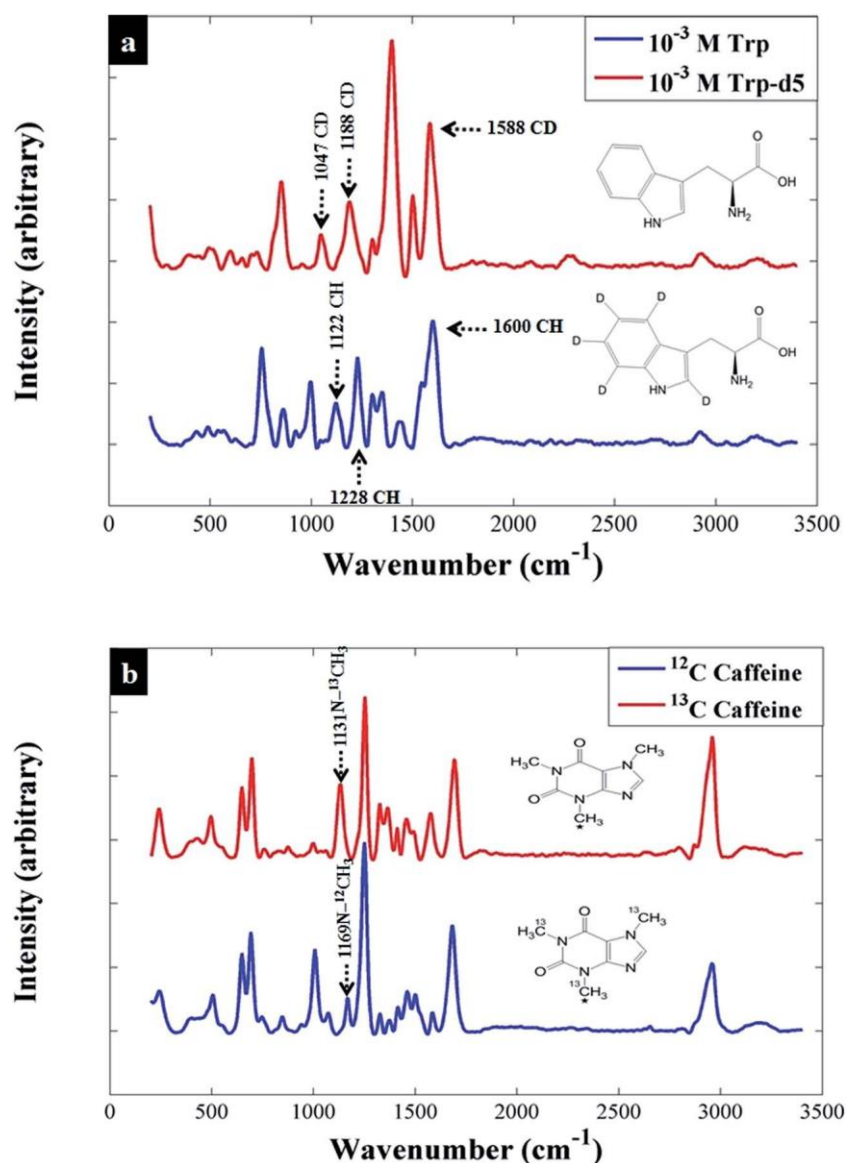


Fig. 1 Baseline-corrected SERS spectra of unlabelled (blue) and labelled (red) analytes: (a) Trp and Trp-d5 with gold citrate and 0.5 M KNO_3 aggregation agent; (b) ^{12}C 'unlabelled' caffeine and ^{13}C labelled caffeine with silver citrate and 1 M NaCl aggregation agent. Key vibrations are highlighted (see Tables 1 and 2 for band assignments). In addition, the structures of tryptophan and caffeine are shown along with the position of the ^2H or ^{13}C isotope label.

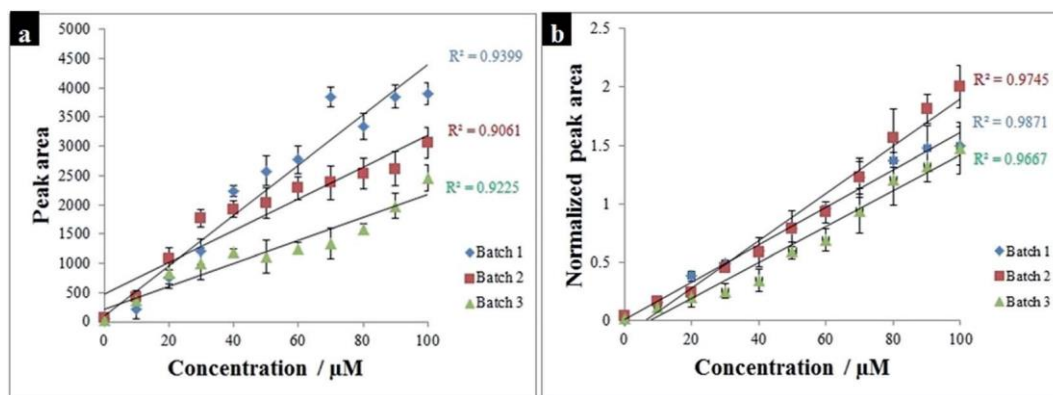


Fig. 2 Plots of characteristic peak for Trp: (a) represents peak area of Trp at 1122 cm^{-1} versus Trp concentration; (b) shows normalized peak area using the IL internal standard at 1047 cm^{-1} for correction. The error bars denote standard deviation of the mean of five measurements.

CH bending vibration, phenol ring breathing and indole ring CH (NH) bending, exhibited a shift to lower wavenumbers at 1047 , 1188 and 1588 cm^{-1} . These band assignments are supported by our DFT results, which confirm these peak assignments. A list of the main bands and their corresponding assignments are presented in Table 1.^{16,47,53}

According to Fig. 1b, the most significant SERS spectral band induced by ^{13}C labelling of caffeine is the peak at 1169 cm^{-1} , assigned to the $\text{N-}^{13}\text{CH}_3$ asymmetric bending vibration, which shifts to 1131 cm^{-1} .⁵⁴⁻⁵⁶ Once again DFT calculations were used in order to obtain the vibrational assignments, and these band assignments for unlabelled and labelled caffeine are provided

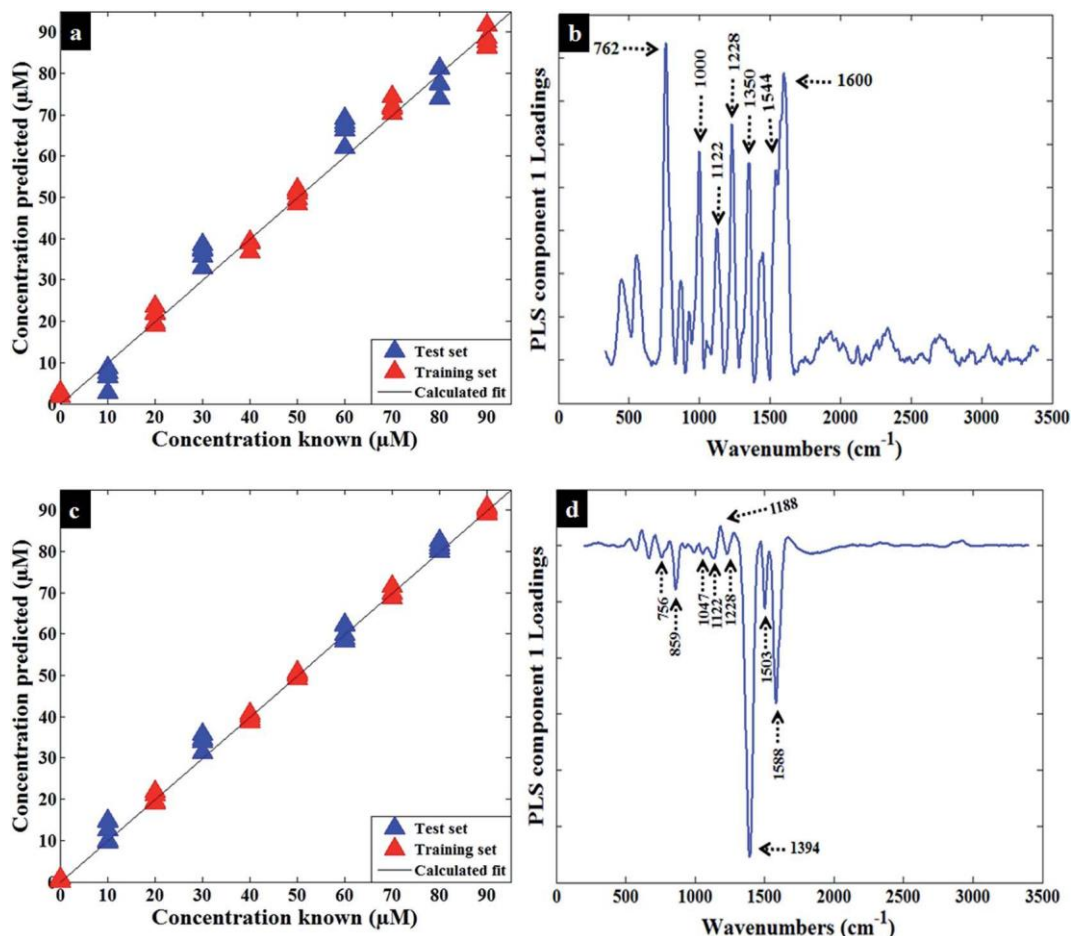


Fig. 3 Representative typical PLSR plots from batch 1 of predicted concentration versus known concentrations of (a) Trp and (c) mixtures of Trp + Trp-d₅. Also shown are the corresponding PLSR loading plots for (b) Trp alone and (d) after including the Trp-d₅ isotopic labelled analyte.

in Table 2. Calibration curves were plotted to establish the linearity of analyte specific peak areas as a function of Trp (Fig. 2a) and caffeine (Fig. S6a†) concentrations. In addition, the ratios of normalised peak areas of the analytes to their IL internal standards were also plotted versus the concentration range (Fig. 2b and S6b†). The linear ranges of these plots were used to calculate the limit of detection (LOD) of Trp and caffeine to be 3.86 mM and 0.1 mM respectively. Noticeably, there is an offset in the normal calibrations (Fig. 2a), which could be due to batch-to-batch variation from repeat productions of the NPs, as well as differences in the size of nanoparticles, their concentrations, and the relative enhancement which may result in poor linearity and accuracy.

However, normalization of the peak of interest at 1122 cm^{-1} to the IL internal standard at 1047 cm^{-1} have clearly overcome this issue and minimised this offset. On closer inspection of the corrections after the use of the isotopologue it can be observed that the three calculated best fit lines are not wholly congruent. It is possible that within the separate nanoparticle batches that this is a result of slight differences in orientation of the tryptophan molecules on the three surface types, due to the

different geometries of the colloid. This means that for each batch of colloid a new calibration curve would need to be generated and this is usual when any detector is calibrated for absolute quantification. We do see that the UV-visible absorbance spectra of all three batches of Au colloid are indeed different (Fig. S1a†). This may then lead to minimal changes in the relative ratios of the C–H and C–D vibrations in tryptophan (from the substitution of ^1H with ^2H on the indole ring in this molecule) that are observed between the batches. As discussed above, the IL internal standard is used as a reference to compensate for experimental errors and as can be seen in Fig. 2b and S6b,† there was an improvement in both accuracy and reproducibility upon normalisation to these IL internal standards. In particular, the linearity of the signal with respect to analyte concentration was improved as the regression results increased from $R^2 = 0.9399$ to $R^2 = 0.9871$ for Trp from batch 1 (see the regression results of batch 2 and 3 in Fig. 2) and from $R^2 =$

0.8394 to $R^2 = 0.9669$ in caffeine (Fig. S6b†).

In addition to the univariate linear regression, we also employed multivariate calibration using the quantitative PLSR models, conducted as described previously in the methods

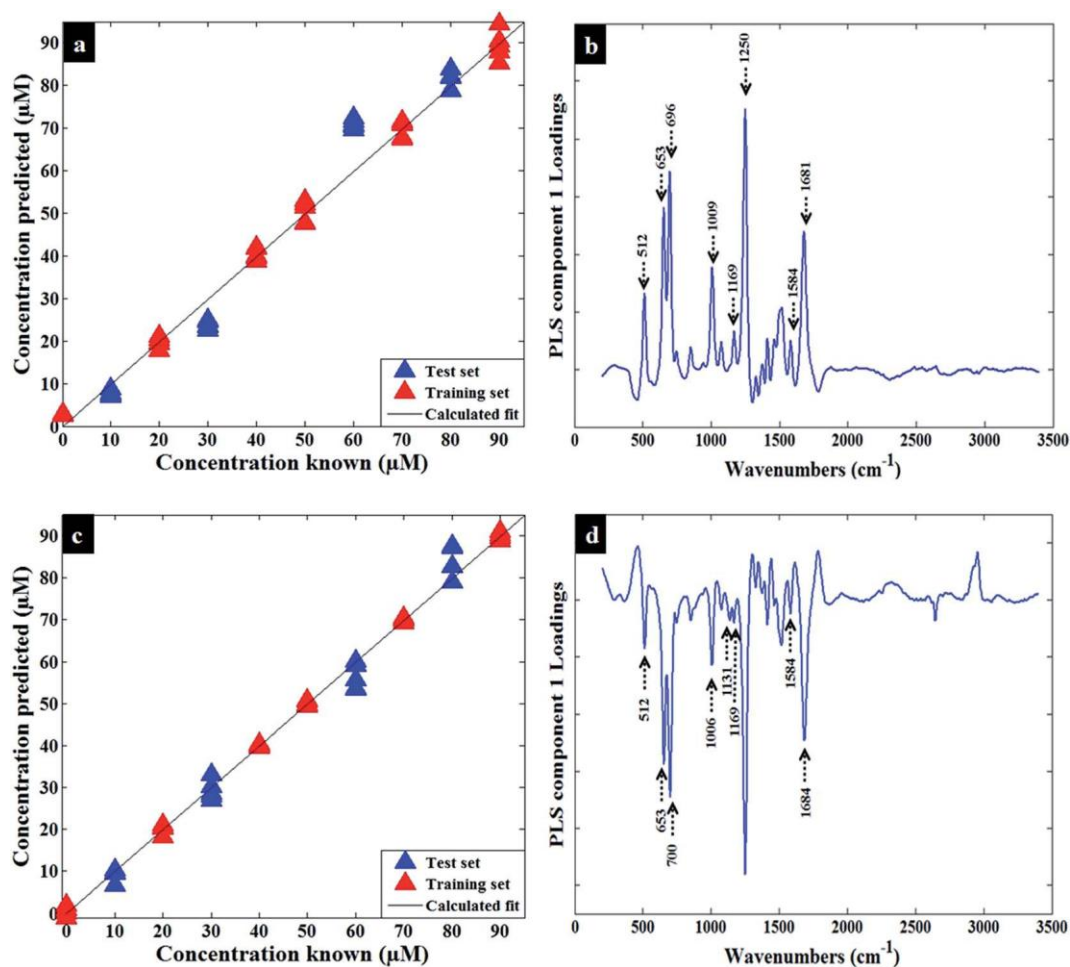


Fig. 4 Representative typical PLSR plots of predicted concentration versus known concentrations of (a) caffeine and (c) mixtures of ^{12}C caffeine + ^{13}C caffeine. Also shown are the corresponding PLSR loading plots for (b) ^{12}C caffeine alone and (d) after including the ^{13}C isotopically labelled caffeine.

Table 3 Summary of the statistics for prediction of Trp using multiple batches of nanoparticles and caffeine with and without spiking with the isotopically labelled equivalent chemical^a

Metric	Batch 1		Batch 2		Batch 3		Batch 1	
	Trp	Trp + Trp-d ₅	Trp	Trp + Trp-d ₅	Trp	Trp + Trp-d ₅	Caffeine ¹² C	Caffeine ¹² C + caffeine ¹³ C
Factors	3	6	3	6	2	3	5	7
R ²	0.9945	0.9993	0.9630	0.9969	0.9313	0.9598	0.9952	0.9993
Q ²	0.8152	0.8542	0.3857	0.7454	0.5385	0.7958	0.7703	0.8490
RMSECV	7.7490	6.8831	16.8798	9.2835	15.1905	10.1286	7.0048	0.7974
RMSEP	6.0296	2.8857	7.3547	6.0944	17.9969	10.7889	2.0788	0.1124

^a Factors specify the number of latent variables used in PLSR; RMSEP, root-mean-squared error of prediction (from the test set); RMSECV root-mean-squared error of cross validation (from the training set); R² and Q² show linearity for the training and test sets predictions, respectively.

section. The PLS models showed improvements via the inclusion of the IL internal standards for the analysis of Trp (Fig. 3a, c, S7a, c, S8a, and c†) in batches 1 to 3 respectively, and caffeine (Fig. 4a and c). Furthermore, the PLS predictions showed good agreement between the predicted concentrations of the target analytes and their known concentrations in general, and again, the models using the data with IL added significantly improved the linearity of the test set predictions (Q²) of Trp from 0.81 to 0.85 from batch 1 and caffeine from 0.77 to 0.85, with a decrease in test set errors (RMSEP) from 6.03 to 2.86 for Trp and from 2.08 to 0.11 for caffeine. The comparison between these two analytes using multiple batches of nanoparticles in terms of higher reproducibility and accuracy is detailed in Table 3. Furthermore, the PLS component loadings were plotted to confirm the above findings, which revealed that the peaks at 1047, 1188 and 1588 cm⁻¹ were the most significant variables affected by Trp-d₅ (Fig. 3b, d, S7b, d, S8b, and d†) from batches 1–3 respectively, and the band at 1131 cm⁻¹ was the peak most distinctly influenced by ¹³C caffeine (Fig. 4b and d).

Conclusions

This study, which was a natural progression from and extension of our previously published work, has clearly demonstrated the potential application of SERS combined with isotopologues (IL-SERS) for the quantitative detection of Trp and caffeine. The unique features of SERS spectra resulting from the added isotopes improved the quantification methods based on the ratio of (analyte/IL internal standard) and this was the case for both ²H for tryptophan and ¹³C for caffeine. The C–D stretch in the tryptophan isotope is clearly identifiable and can be used as the internal standard vibration for the C–H stretch from non-deuterated tryptophan (i.e. the natural isotope abundance). In order to demonstrate the reproducibility of our approach, we chose to repeat the quantitative analysis of this molecule with three different batches of colloid over a 12 month period. The results reported here clearly demonstrate that this isotopologue improves the level of quantification, both from univariate analysis as well as via the multivariate method of PLSR. Thus the application of the IL approach, has clearly improved the accuracy of the PLSR models while minimising errors resulting

from batch-to-batch variations of nanoparticles. This approach may also be useful when the SERS band shifts due to the isotopes overlap with other vibrational features in the spectra (which would mean that ratios of bands would be difficult to compute). The recorded Raman and SERS spectra of Trp and caffeine and their labelled compounds were assigned based on literature assignments as well as theoretically modelled spectra from quantum chemical calculations based on density functional theory. For our SERS analyses we determined the LODs of Trp and caffeine to be 3.86 mM and 0.1 mM respectively, with indication of high reproducibility, accuracy and stability of the spectra that forwards IL-SERS as a practicable strategy with potential for various analytical applications. Future studies will be conducted to extend this method for analysis of target determinants present in complex biological samples such as blood, plasma and urine.

Conflicts of interest

The authors have no conflicts of interest.

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