

## REVIEW

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Cite this: *RSC Adv.*, 2018, 8, 25888

# Challenges in application of Raman spectroscopy to biology and materials

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Raman spectroscopy has become an essential tool for chemists, physicists, biologists and materials scientists. In this article, we present the challenges in unravelling the molecule-specific Raman spectral signatures of different biomolecules like proteins, nucleic acids, lipids and carbohydrates based on the review of our work and the current trends in these areas. We also show how Raman spectroscopy can be used to probe the secondary and tertiary structural changes occurring during thermal denaturation of protein and lysozyme as well as more complex biological systems like bacteria. Complex biological systems like tissues, cells, blood serum etc. are also made up of such biomolecules. Using mice liver and blood serum, it is shown that different tissues yield their unique signature Raman spectra, owing to a difference in the relative composition of the biomolecules. Additionally, recent progress in Raman spectroscopy for diagnosing a multitude of diseases ranging from cancer to infection is also presented. The second part of this article focuses on applications of Raman spectroscopy to materials. As a first example, Raman spectroscopy of a melt cast explosives formulation was carried out to monitor the changes in the peaks which indicates the potential of this technique for remote process monitoring. The second example presents various modern methods of Raman spectroscopy such as spatially offset Raman spectroscopy (SORS), reflection, transmission and universal multiple angle Raman spectroscopy (UMARS) to study layered materials. Studies on chemicals/layered materials hidden in non-metallic containers using the above variants are presented. Using suitable examples, it is shown

Received 25th May 2018

Accepted 9th July 2018

DOI: 10.1039/c8ra04491k

rsc.li/rsc-advances

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include Raman spectroscopy, SERS, development of novel Raman based instrumentation towards detection of materials, air quality management, catalysis, carbon based materials, adsorption and nanoparticles.



how a specific excitation or collection geometry can yield different information about the location of materials. Additionally, it is shown that UMARS imaging can also be used as an effective tool to obtain layer specific information of materials located at depths beyond a few centimeters.

## Introduction

'VidyaViniyogaVikasaah' (a Sanskrit phrase) means progress comes from proper application of knowledge. This statement is true for the research and development in the field of Raman spectroscopy, which is considered as an evolving technology towards a direct impact to society. Although the discovery happened in 1928, the number of publications remained low until the 1960s; the web of science data indicates that there was an upsurge in the number of publications in the decade following the invention of lasers followed by a steady growth in the number of publications. The 90's saw a revival in Raman research both in theory as well as on instrumentation. Its potential applications were understood immediately as this is a straightforward technique without much need for sample preparation and almost any material, in any state under extreme conditions, can be probed. In addition, due to its non-destructive nature, it can be used as a technique for *in situ* measurements. Moreover, Raman microscopy has made it possible to probe very small quantity of materials with high spatial resolution. With the advent of techniques like Scanning Near-Field Microscopy (SNOM) and Tip-Enhanced Raman Spectroscopy (TERS), resolution of the order of few nanometres can also be achieved. This advantage makes it an essential and unique technique in the field of biology and materials science to explore the molecular basis of structure and function.<sup>1-4</sup>

Progress in the field of Raman spectroscopy can be largely attributed to the recent advances made in the areas of instrumentation.<sup>5</sup> Historically, analytical applications involving Raman spectroscopy was thought to be difficult, however, developments in the sources such as diode lasers, charge-coupled devices (CCDs), small spectrometers, optics along

with interfacing optical fibres with Raman have made it a novel method for analytical applications. Confluence of nanotechnology and Raman spectroscopy which led to the discovery of Surface Enhanced Raman Spectroscopy (SERS) has opened new vistas for detection of analytes at very low concentrations. It is possible to obtain Raman signals of biomolecules, chemicals, explosives, drugs *etc.* at parts per billion (even at single molecule level) with great sensitivity and specificity.<sup>6-17</sup>

Raman spectroscopy owing to its molecular specificity can be a boon, however, in the case of biological samples, it complicates the analyses. Firstly, all biological systems are composed of biochemicals such as lipids, proteins, nucleic acids, carbohydrates *etc.* Therefore, vibrations from all the modes comprising the chemicals shall be manifested in the Raman spectra. To an untrained eye, spectra of a cell, tissue and a bacterium may appear very similar. Therefore, analyses of such convoluted complex spectra and assignments can be a challenging endeavour. Despite this, these complex spectra can yield a treasure of information pertaining to the structure and dynamics of the sample under evaluation. Once the assignments are made, the whole world of biochemical processes, such as metabolic pathways or dynamics can be unearthed. The second challenge towards obtaining Raman spectra of biomolecules is the occurrence of peaks from the matrix. For instance, paraffin fixed tissue may show a similar peak to a C-H stretch.<sup>18,19</sup> Differentiating the actual spectra from the matrix therefore, becomes an equally important part before analyses. Thirdly, understanding the system under study and making an informed judgement based on the experiments and correlating it with the available data is crucial. Finally, signal processing of a large number of data set can be a contributing factor towards understanding of the system.



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The objective of this article is based on the review of our work to demonstrate the challenges and potential of Raman spectroscopy for applications in biology and materials. Emphasis has largely been placed on the challenges of using Raman spectroscopic techniques for studying biomolecules and non-invasive, depth sensitive identification of materials. This laboratory has been working on various facets of Raman spectroscopy for the last three decades. Only a few results pertaining towards research on biological systems and materials have been included in this review which is organized in the following manner:

(1) Discussion on the challenges of probing biomolecules using Raman spectroscopy. Various examples on Raman spectroscopic study of biomolecules such as proteins, lipids *etc.* have been discussed. Furthermore, examples of Raman spectroscopy for disease diagnosis have been presented. In particular, identifying and assigning Raman spectral bands to specific molecular structures in relation to multitude of complex biomolecules present in a system.

(2) Discussion on depth sensitive Raman measurements such as spatially offset Raman spectroscopy, universal multiple angle Raman spectroscopy towards depth sensitive detection of chemicals (explosives) and transmission Raman spectroscopy. Recent developments and the relative advantages have also been presented.

## Results and discussions

### (I) Raman spectroscopy towards biological applications

The following sections review how fundamental studies on different types of biomolecules can be carried out and also the complications involved in Raman band assignments using Raman spectroscopy. We also discuss some of the very recent applications of Raman spectroscopy in translational research, citing examples of how this technique can help in understanding and diagnosing different diseases like cancer, malaria, dengue and others.

**(a) Raman spectroscopy of biomolecules.** Biomolecules are essential building blocks of life. They are central to all life processes and are involved in carrying out important metabolic reactions and maintaining the overall biochemistry of a living organism. Different kinds of biomolecules like nucleic acids, proteins, lipids and carbohydrates are involved in maintaining the functional homeostasis of a cell. Any biological material (cell, tissue *etc.*) is made up of different composition of various biomolecules. Therefore, it is important to understand the structure and properties of various biomolecules separately. Interestingly, each of these biomolecules has distinct chemical identities and therefore has unique molecular structures. Consequently, these biomolecules generate a signature Raman spectrum.<sup>20,21</sup>

To demonstrate this aspect, we have recorded Raman spectra of pure biomolecules belonging to different classes-protein (Bovine Serum Albumin (BSA)), DNA (calf-thymus DNA), lipids (cholesterol) and carbohydrates (glucose), as shown in Fig. 1.

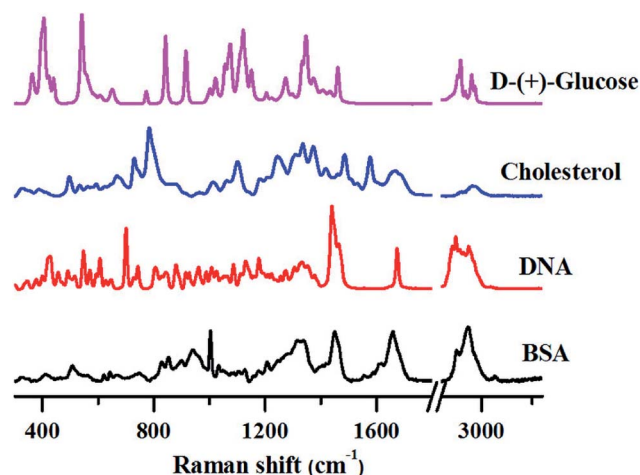


Fig. 1 Distinctive Raman spectra of various biomolecules. All spectra were collected from powdered samples using 785 nm excitation. Spectra were corrected for cosmic rays and baseline corrected prior to analysis.

Although the Raman spectra shown above are quite complex, several unique marker bands can be identified for each of these biomolecules as explained below.

**Proteins.** Proteins have numerous functions like catalyzing many reactions, guiding the flow of electrons in reactions, transmitting information between specific cells, controlling the passage of molecules around the cell membrane or providing the filamentous architecture within cells *etc.* They are characterized by their unique three-dimensional architecture.<sup>22</sup> A small change in the conformation can make them biologically inactive and may even be toxic. Various life-threatening diseases have been associated with such denaturation of proteins for *e.g.*, Alzheimer's disease, Parkinson's disease *etc.*<sup>23,24</sup> Raman spectroscopy is an excellent tool to look at the protein structure. Several vibrational modes can be used for the analysis of peptide structure. Most characteristic bands are associated with the CONH group, referred to as amide A (NH stretching, about 3500 cm<sup>-1</sup>), amide B (NH stretching, about 3100 cm<sup>-1</sup>) and Amide I to VII.<sup>25</sup> Out of all, Amide I & III are frequently used to estimate the secondary structure of proteins. Amide I mode which ranges from 1580 cm<sup>-1</sup> to 1700 cm<sup>-1</sup> arises from stretching vibration of C=O.<sup>26</sup> It is very sensitive to the backbone conformation and not affected by the side chains. Amide I band can be deconvoluted with various sub-bands which directly correlate with various secondary structures. Similarly, Amide III band (1200 cm<sup>-1</sup>–1300 cm<sup>-1</sup>) which arises from C–N stretching coupled with N–H bending vibrations can also give complementary information about the protein backbone conformations.<sup>27,28</sup> Asher's group suggested an empirical relation between Amide III frequency and dihedral angle  $\Phi$ , which paves the way to analyse conformation of proteins using Amide III band of Raman spectra.<sup>29,30</sup>

Apart from the secondary structure, characteristic Raman bands have also been identified for various side chains and disulphide regions as well. George J. Thomas resolved the signature for each of the eight cysteines per 666-residue subunit



in the trimeric tail spike of an icosahedral bacteriophage P22.<sup>31</sup> They demonstrated that a broad range of S-H...X interactions is accessible to the protein. The integration of microscope in Raman spectrometer has enabled studying of protein crystals, which led to genesis of a new technique termed as Raman crystallography.<sup>32</sup> Raman crystallography can identify and follow the conformations of reaction intermediates in single crystals. Helfand *et al.* monitored the reactions between three clinically relevant inhibitors, tazobactam, sulbactam, and clavulanic acid, and SHV beta-lactamase (EC 3.5.2.6) in single crystals using a Raman microscope and observed acyl-enzyme formed between Ser70 and the lactam ring's C=O group as a key intermediate in the reaction pathway.<sup>33</sup> Various characteristic Raman bands for proteins are summarized in Table 1.

**Nucleic acids.** Richard C. Lord and co-workers recorded the Raman spectra of various nucleic acid constituents for the first time in 1960.<sup>47</sup> Further improvements in the instrumentation have enabled studying of various problems concerning the structure and interaction in nucleic acids, nucleic acid hydration, interaction with metal ions, DNA melting, DNA damage *etc.*<sup>48–51</sup> Raman spectra of nucleic acids contain a large number of bands originating from various moieties like sugar, phosphate, bases *etc.* which provide very useful information about backbone conformation, base pairing *etc.*

Very strong Raman peaks are observed by ring breathing modes of various bases in the region of 600 cm<sup>-1</sup> to 800 cm<sup>-1</sup>.<sup>20</sup> Therefore, this region can be used to differentiate various bases. Out of all the bases, thymine has a very distinctive peak at around 1671 cm<sup>-1</sup> originating from C=O stretching which makes it easily distinguishable from rest of the bases.<sup>20</sup> Different conformations of backbone (A-DNA, B-DNA and Z-DNA) can be identified using Raman spectra. J. M. Benevides and co-workers have studied different backbone conformation at various salt conformations. They could qualitatively identify the A, B and Z-DNA using phosphodiester Raman band lying in the region of 750 to 800 cm<sup>-1</sup>.<sup>52</sup> Apart from the above, many

important group vibrations have also been assigned for nucleic acids in vibrational spectra, for example, region from 1800–1500 cm<sup>-1</sup> has been assigned for nucleobase vibrations appear which are extremely sensitive to base pairing interactions and 1500–1250 cm<sup>-1</sup> region have been assigned for vibrational coupling between the base-sugar entities, stacking ligation *etc.*<sup>53</sup> Okotrub *et al.* (2014) quantified DNA in the cell the nucleus using intensity of phosphate mode at 1096 cm<sup>-1</sup> without extraction or dye labelling using Raman spectroscopy.<sup>54</sup> Wood's group (2013) probed DNA double-strand breaks (DSBs) with TERS coupled with Atomic Force Microscopy (AFM) and suggested that the O-C bond in the DNA backbone is most sensitive to UVC radiation and its cleavage causes DSBs most of the time.<sup>55</sup> Deckert's group recorded the TERS spectra from an artificial DNA single strand with alternating blocks of adenine and cytosine. They could resolve adenine and cytosine bases placed at a distance of 2.1 nm by unique Raman spectral features proving that TERS can offer resolutions upto single nucleobase level.<sup>56,57</sup> Various characteristic Raman bands for nucleic acid are summarized in Table 2.

**Lipids.** Lipids, also called fats, are one of the important classes of biomolecules. They have important role in energy storage, cellular signalling, digestion and absorption of food, building of cellular membranes and various tissues, maintaining body temperature *etc.* Raman spectroscopy provides an invaluable tool to estimate various structural properties of lipids like the degree of unsaturation, level of *E*, *Z* unsaturation *etc.*<sup>60</sup> The stretching vibrations from C=O in lipids are observed at around 1749 cm<sup>-1</sup> which are quite distinct and far from that of proteins. Similarly, many other markers bands for lipids like Raman bands at 1430 cm<sup>-1</sup> (CH<sub>2</sub> scissoring), 1443 cm<sup>-1</sup> (CH<sub>2</sub> and CH<sub>3</sub> deformation of lipids and triglycerides), 1453 cm<sup>-1</sup> (C-H bending of lipids) have been used to quantify lipids in oral cancer tissues.<sup>61</sup> Nieva *et al.* (2012) characterized the lipid phenotype of breast cancer cells using Raman spectroscopy.<sup>62</sup> Raman spectra analysed in the range of 2820–3030 cm<sup>-1</sup>, where

Table 1 Peak assignments for proteins<sup>a</sup>

| Contributing group | Frequency (cm <sup>-1</sup> ) | Assignment                                                  |
|--------------------|-------------------------------|-------------------------------------------------------------|
| Backbone           | 1670–1680                     | β turn, random structure, Amide I <sup>26</sup>             |
|                    | 1660–1670                     | β sheet, Amide I <sup>34</sup>                              |
|                    | 1650–1655                     | α helices, Amide I <sup>35</sup>                            |
|                    | 1630–1635                     | β sheet, Amide I <sup>36,37</sup>                           |
|                    | 1550                          | Amide II <sup>25</sup>                                      |
|                    | 1270–1300                     | α helices, Amide III <sup>38</sup>                          |
|                    | 1240–1250                     | Random structure, Amide III <sup>39</sup>                   |
|                    | 1230–1240                     | β sheet, Amide III <sup>39</sup>                            |
| Disulphide         | 505–515                       | GGG conformation <sup>40,41</sup>                           |
|                    | 520–530                       | GGT, GTG conformation <sup>40,41</sup>                      |
|                    | 540–545                       | TGT conformation <sup>40,41</sup>                           |
| Side chain         | 1615, 830, 643                | Tyrosine <sup>42</sup>                                      |
|                    | 850/830                       | Hydrogen bonding state of tyrosine; <sup>43</sup>           |
|                    | 1550, 1010, 750               | Tryptophan <sup>44</sup>                                    |
|                    | 880/1361                      | Ring environment of indole ring of tryptophan <sup>45</sup> |
|                    | 1360/1340                     | Hydrophobicity marker for tryptophan <sup>44</sup>          |
|                    | 1609, 1205, 1003              | Phenylalanine <sup>46</sup>                                 |

<sup>a</sup> α: alpha helices, β: beta sheets, GGG: *gauche-gauche-gauche*, GGT: *gauche-gauche-trans*, GTG: *gauche-trans-gauche*, TGT: *trans-gauche-trans*.



Table 2 Peak assignments for nucleic acids<sup>a</sup>

| Contributing group | Frequency (cm <sup>-1</sup> ) | Assignment                                                                             |
|--------------------|-------------------------------|----------------------------------------------------------------------------------------|
| Base               | 726                           | Ring breathing of A <sup>58</sup>                                                      |
|                    | 668, 746, 785                 | Ring breathing of T <sup>58</sup>                                                      |
|                    | 668                           | Ring breathing of G <sup>58</sup>                                                      |
|                    | 781, 785                      | Ring breathing of C <sup>58</sup>                                                      |
|                    | 785                           | Ring breathing of U <sup>58</sup>                                                      |
|                    | 1257                          | Ring mode (C, T) <sup>58</sup>                                                         |
|                    | 1336                          | Ring mode (A, G); A form of DNA <sup>59</sup>                                          |
|                    | 1485                          | Ring stretching (G, A) A form of DNA <sup>59</sup>                                     |
| Backbone           | 1576                          | Ring stretching (A, G); A form of DNA <sup>59</sup>                                    |
|                    | 807                           | $\nu_{\text{as}}(\text{O-P-O})$ , $\nu_{\text{s}}(\text{O-P-O})$ ; A-DNA <sup>52</sup> |
|                    | 1099                          | $\nu_{\text{s}}(\text{PO}_2^-)$ ; A-DNA <sup>52</sup>                                  |
|                    | 1415                          | df(CH <sub>2</sub> ); A-DNA <sup>52</sup>                                              |
|                    | 784                           | $\nu_{\text{s}}(\text{O-P-O})$ ; B-DNA <sup>52</sup>                                   |
|                    | 830                           | $\nu_{\text{as}}(\text{O-P-O})$ ; B-DNA <sup>52</sup>                                  |
|                    | 1090                          | $\nu_{\text{s}}(\text{PO}_2^-)$ ; B-DNA <sup>52</sup>                                  |
|                    | 1420                          | df(CH <sub>2</sub> ); B-DNA <sup>52</sup>                                              |
|                    | 748                           | $\nu_{\text{s}}(\text{O-P-O})$ ; Z-DNA                                                 |
|                    | 792                           | $\nu_{\text{as}}(\text{O-P-O})$ ; Z-DNA                                                |
|                    | 810                           | $\nu_{\text{as}}(\text{O-P-O})$ ; Z-DNA                                                |
|                    | 1095                          | $\nu_{\text{s}}(\text{PO}_2^-)$ ; Z-DNA                                                |
|                    | 1425                          | df(CH <sub>2</sub> ); Z-DNA                                                            |

<sup>a</sup>  $\nu$ : stretching (s: symmetric, as: asymmetric), b: bending, t: twisting, df: deformation, A: adenine, T: thymine, C: cytosine, G: guanine, U: uracil, A-DNA: A-form of DNA, B-DNA: B form of DNA, Z-DNA: Z form of DNA.

total fatty acid content (TFA) and total unsaturated fatty acids (TUFA) (3015 cm<sup>-1</sup>) bands were quantified using bands at 2845 cm<sup>-1</sup> and 3015 cm<sup>-1</sup> respectively. Popp's group (2015) analysed fungal lipids inside living hyphae using Raman spectroscopy.<sup>63</sup> Raman spectra from single vesicles were collected and analysed for peak intensity ratio  $I(1270 \text{ cm}^{-1})/I(1445 \text{ cm}^{-1})$  from the signals of =CH and -CH<sub>2</sub>/-CH<sub>3</sub> groups, respectively. This ratio is linked to the iodine values of lipids. Raman spectra from lipids can be characterized in seven regions as explained in Table 3 (all assignments have been taken from Czamara *et al.*<sup>60</sup>).

**Carbohydrates.** Carbohydrates are naturally occurring compounds and are essential constituents for living organisms. They have significant functions, like, main source of energy, major constituent of cell wall of plants and microorganisms, molecular agents in cellular recognition *etc.* Carbohydrates can be classified as monosaccharides, oligosaccharides and polysaccharides. Nonetheless, structural investigations on different carbohydrates are difficult as the basic monomeric unit in the

polymeric chains is very similar. Despite this, different nature of H-bonding or the ring configuration gives rise to several marker peaks for carbohydrates. Wiercigroch *et al.* (2017) have compiled Raman and IR features of 14 carbohydrates in the solid state.<sup>64</sup> Arboleda *et al.* recorded Raman spectra of nine anomeric stable monosaccharides in solutions with concentrations as low as 10 mM and volumes as small as 15  $\mu\text{L}$ . It is observed that Raman spectra are sensitive to the configuration of the carbon centres and unique spectra are obtained of all nine monosaccharides.<sup>65</sup> Stuart *et al.* (2006) demonstrated *in vivo* quantification of glucose in animal model for the first time using Surface-Enhanced Raman Spectroscopy (SERS).<sup>66</sup> Raman spectra of carbohydrates can be categorised in five main regions as given in Table 4. All assignments have been taken from Wiercigroch *et al.*<sup>64</sup>

The above work provides extensive evidence that different biomolecules yield distinct Raman spectra. While understanding the absolute structure of any biomolecule can be a confounding task, monitoring changes in the chemical

Table 3 Peak assignments for lipids<sup>a</sup>

|            | Frequency (cm <sup>-1</sup> ) | Assignment                                                                                                                                         |
|------------|-------------------------------|----------------------------------------------------------------------------------------------------------------------------------------------------|
| Region I   | 2800–3100                     | $\nu(\text{CH})$ , $\nu_{\text{as}}(\text{CH}_3)$ , $\nu_{\text{s}}(\text{CH}_3)$ , $\nu_{\text{as}}(\text{CH}_2)$ , $\nu_{\text{s}}(\text{CH}_2)$ |
| Region II  | 1600–1800                     | $\nu(\text{C}=\text{C})$ , $\nu(\text{C}=\text{O})$                                                                                                |
| Region III | 1400–1500                     | b(CH <sub>2</sub> /CH <sub>3</sub> ), sc(CH <sub>2</sub> /CH <sub>3</sub> ), b(CH <sub>2</sub> )                                                   |
| Region IV  | 1200–1300                     | t(CH <sub>2</sub> ), df(CH)                                                                                                                        |
| Region V   | 1050–1200                     | $\nu(\text{C-C})$ , $\nu(\text{P-O})$                                                                                                              |
| Region VI  | 800–1050                      | b(CH), skeletal(C-O-O), $\nu_{\text{as}}\text{N}^+(\text{CH}_3)_3$                                                                                 |
| Region VII | 500–700                       | $\nu_{\text{s}}\text{N}^+(\text{CH}_3)_3$ , cholesterol ring df, df(C=O-O), b(CH <sub>2</sub> ) ring                                               |

<sup>a</sup>  $\nu$ : stretching (s: symmetric, as: asymmetric), b: bending, sc: scissoring, t: twisting, df: deformation, skel: skeletal.



Table 4 Peak assignments for carbohydrates<sup>a</sup>

|            | Frequency (cm <sup>-1</sup> ) | Assignment                                                                        |
|------------|-------------------------------|-----------------------------------------------------------------------------------|
| Region I   | 3100–3600                     | $\nu(\text{OH})$                                                                  |
| Region II  | 2800–3100                     | $\nu(\text{CH})$ , $\nu_{\text{as}}(\text{CH}_2)$ , $\nu_{\text{s}}(\text{CH}_2)$ |
| Region III | 1200–1500                     | $\text{df}(\text{CH}/\text{CH}_2)$                                                |
| Region IV  | 800–1200                      | $\nu(\text{C}-\text{O})$ , $\nu(\text{C}-\text{C})$                               |
| Region V   | 100–800                       | $\text{df}(\text{CCO})$                                                           |

<sup>a</sup>  $\nu$ : stretching (s: symmetric, as: asymmetric), b: bending, sc: scissoring, t: twisting, df: deformation.

structure in response to some perturbation can be efficiently analysed using Raman spectra. Consequently, Raman spectroscopy has been used to study changes in the structure of various biomolecules with external perturbation, like protein folding, protein–protein interaction, DNA–drug interaction, DNA melting *etc.*<sup>67</sup>

To elucidate further, we performed thermal denaturation studies on the protein, lysozyme and compared the heat-induced changes in the Raman spectra with the native form (Fig. 2).

The Amide I peak in a Raman spectrum is a broad band which ranges from 1580–1720 cm<sup>-1</sup>. It can be deconvoluted into various sub-bands corresponding to different secondary structures like helices,  $\beta$ -sheets, random structures *etc.* The band at around 1655 cm<sup>-1</sup> is usually assigned for helices, 1665 cm<sup>-1</sup> for  $\beta$ -sheets and 1675 cm<sup>-1</sup> for random structures. Lysozyme in its native state, has 41% helices and 10%  $\beta$ -sheets,<sup>68</sup> therefore, Amide I for the native state of this protein had a peak centre around 1660 cm<sup>-1</sup>. After heat denaturation, the content of  $\beta$ -sheets and random structure increased at the expense of helices. Hence, the intensity of the band corresponding to the helices ( $\sim 1655$  cm<sup>-1</sup>) decreased and the intensity of the bands for  $\beta$ -sheets ( $\sim 1665$  cm<sup>-1</sup>) and random structures ( $\sim 1675$  cm<sup>-1</sup>) increased. This resulted in a change in the band shape of Amide I and a consequent shift in the peak maxima. This shift in the peak maxima was better understood by taking the second derivative of the spectra.

The changes in the Amide III band ( $\sim 1200$ – $1300$  cm<sup>-1</sup>) also corroborated this fact. There was a decrease in the intensity of the Amide III band around 1300 cm<sup>-1</sup>, assigned for  $\alpha$ -helices. This was coupled with an increase in the intensity of the band between 1200–1260 cm<sup>-1</sup>, which has contributions from  $\beta$ -sheets and random coils.

The modulations in the Amide I and III bands demonstrated how one can probe the changes in the secondary structure of proteins.

The tertiary structure of the protein can also be probed using various Raman marker bands for side-chains. For instance, the ratio of 1360 cm<sup>-1</sup> and 1340 cm<sup>-1</sup> serves as a hydrophobicity marker for tryptophan. As a consequence of thermal denaturation of lysozyme, there was a decrease in this ratio, indicating a change in the environment of buried tryptophan residues from hydrophobic to hydrophilic. Similarly, there are several other bands assigned for side chains of proteins which can be analysed using Raman spectroscopy (see Table 1).

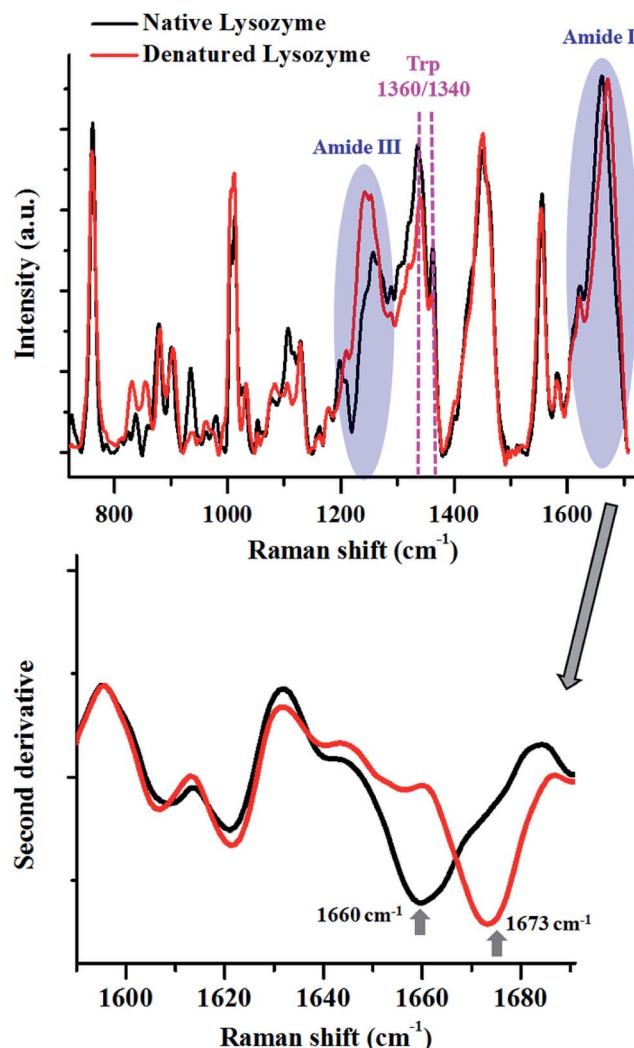


Fig. 2 Raman spectra of lysozyme in its native (black) and thermally denatured (red) state. The major changes in the spectra are highlighted. The shift in the peak maxima of Amide I is shown by the second derivative spectra. Spectra shown in top panel are baseline corrected and vector normalised prior to analysis.

This highlights how Raman spectroscopy has proved to be a valuable technique in analysing conformational changes in the proteins as result of different perturbations.

However, it can become a very daunting task when one wants to analyse complex biological systems like cells, tissues, body fluids *etc.* While biomolecules are ubiquitously present in all biological systems, their relative concentrations can be vastly different. This becomes very evident when the Raman spectra of different tissues and body fluids are compared, since a differential concentration of the biomolecules results in a characteristic Raman spectrum.

We have highlighted this fact by comparing the Raman spectra of liver tissue and blood serum isolated from healthy mice. Raman spectra were recorded from 50  $\mu\text{m}$  thin sections of liver tissue and  $\sim 2$   $\mu\text{L}$  of blood serum isolated from BALB/c mice using 785 nm laser excitation (Fig. 3). The region, 600–1725 cm<sup>-1</sup> was analysed since this constitutes the fingerprint



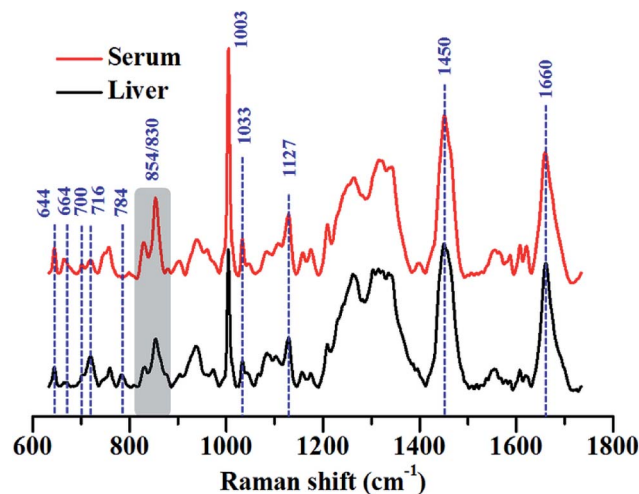


Fig. 3 Average Raman spectra of liver and blood serum isolated from BALB/c mice. The major peaks that were different across the samples have been highlighted using arrows and the different peak ratios that were considered are highlighted using grey boxes. Each spectrum shown is an average of 200 spectra collected from 4 mice. All spectra were baseline corrected, vector normalised and smoothed prior to analysis.

region for most biological samples. Intense generic bands with contributions from proteins and lipids, like 1003, 1451 and 1660  $\text{cm}^{-1}$ , were observed in the Raman spectra of liver and serum.

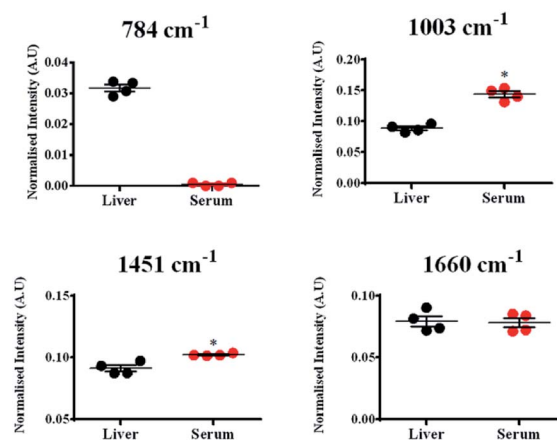
The serum is predominantly composed of different kinds of proteins. It contains up to 80  $\text{mg mL}^{-1}$  of protein along with other constituents like amino acids, salts, lipids and sugars like glucose.<sup>69</sup> The bulk of the protein content of serum comes from albumins, globulins and lipoproteins. This could also be seen in the Raman spectra of serum which was heavily dominated by Raman bands characteristic of proteins, for example, 644, 830, 854, 1003 and 1033, 1127, 1200–1300  $\text{cm}^{-1}$  (Amide III) and 1660  $\text{cm}^{-1}$  (Amide I) respectively. All assignments have been listed in the preceding tables.

While these bands were also present in the Raman spectra of liver, their relative intensities and the ratios varied across both the samples, indicating differences in the relative distribution of biomolecules (Fig. 4(A) and (B)).

The Raman band for DNA at 784  $\text{cm}^{-1}$  was observed only in the spectra of liver and was completely absent in the serum indicating negligible amounts of DNA present in the serum (Fig. 4(A)). Liver, being a tissue is made up of a large number of cells containing nucleic acids like DNA and RNA. On the contrary, blood serum is largely cell-free and thus devoid of any quantifiable amounts of DNA under normal conditions.

Other peaks corresponding to biomolecules such as cholesterol (700  $\text{cm}^{-1}$ ) and phosphatidylcholine (716  $\text{cm}^{-1}$ ) could also be observed in the Raman spectra of both the samples. The Raman bands for glycogen could be seen at 854 and 937  $\text{cm}^{-1}$ .<sup>70,71</sup> These bands were more intense in the spectra of liver, which is quite expected, since liver is the main organ that stores glycogen.

#### (A) Calculation of intensities



#### (B) Calculation of intensity ratio

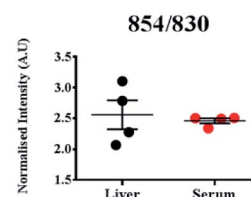


Fig. 4 Calculation of Raman band intensities and intensity ratios pertaining to different biomolecules in liver and serum. Marker bands were chosen for each of the biomolecules; (A) DNA (784  $\text{cm}^{-1}$ ), proteins and lipids (1003, 1451 and 1660  $\text{cm}^{-1}$ ). (B) Ratiometric analysis of two Raman bands – 830 and 854  $\text{cm}^{-1}$ . Each circle in the figure represents an individual mouse. Spectra obtained from a total of 4 mice were used for enumeration. Statistical analysis was performed using Mann–Whitney test where \* indicates  $p < 0.05$ .

Quantifying ratios of Raman bands can be very advantageous as they are least affected by background fluctuations and pre-processing methods.<sup>72</sup> The ratio of two Raman bands at 830 and 854  $\text{cm}^{-1}$  has been shown in Fig. 4(B). These bands have been famously called as the tyrosine Fermi doublet, originating from proteins. The ratio has extensive significance and can be a very good marker for the H-bonding status of the phenolic OH of tyrosine.<sup>69</sup> In serum, these two bands can be assigned to tyrosine since serum is a rich source of proteins. However, in case of liver, the assignment of the band at 854  $\text{cm}^{-1}$  can be ambiguous. In liver, the main contribution to this band comes from glycogen, since it is this organ where glycogen is primarily stored. This point brings to one's notice that most Raman bands can have overlapping contributions from several biomolecules and therefore, understanding the system under study is essential before band assignments are made.

As already mentioned that although the Raman spectra of such biological entities are quite complex, certain marker Raman bands can still be assigned for different classes of biomolecules in such systems.<sup>51,73,74</sup> This can help us monitor the changes induced as a result of any stress or change in the native environment. This point can be proven through an



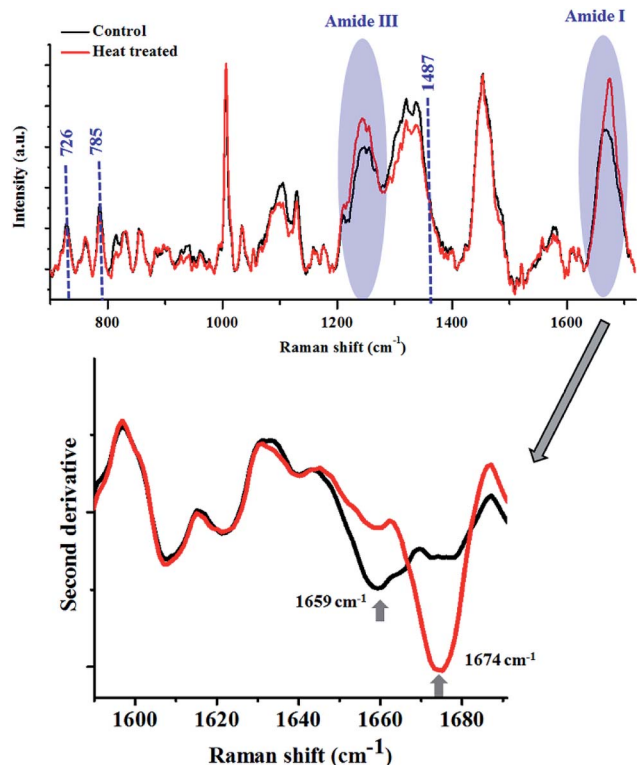


Fig. 5 Raman spectra of bacterial cells before (black) and after (red) heat treatment. *E. coli* WT cells were grown for 8 h in shaking condition at 37 °C and 160 rpm. Bacteria were heat treated at 85 °C for 15 minutes. The shift in the peak maxima of Amide I is shown by the second derivative spectra. All spectra in the top panel were corrected for cosmic rays, background subtracted and vector normalised prior to analysis.

example by monitoring the biomolecular changes induced in bacteria as a consequence of heat treatment.

Heat treatment of the bacterial cells induced several changes in the Raman spectra (Fig. 5), as highlighted in the figure above. The major changes observed were in the Amide I and III bands, similar to what was noted in the lysozyme-denaturation experiment shown in Fig. 2. This emphasises that these Raman bands can be unambiguously assigned to proteins, even in complex systems like bacteria.

After heat treatment, a prominent increase in the intensity of the Raman band at 1674  $\text{cm}^{-1}$  was observed, which indicated the increase in the content of  $\beta$ -sheets/random coils. This was also highlighted by the second derivative spectra. Complementary information was obtained from the analysis of the Amide III band. Accumulation of  $\beta$ -sheet/random coils is highlighted by the increase in intensity of Raman bands in the range of 1200–1260  $\text{cm}^{-1}$  at the expense of helical structure (1300  $\text{cm}^{-1}$ ). Another biomolecule which showed changes as a result of heat treatment was DNA. All the major Raman bands for DNA (726, 785, 1487  $\text{cm}^{-1}$ ) showed a marginal decrease in the heat treated bacterial cells.

These changes showcase how complex systems can also be studied using Raman spectroscopy by analysing the unique features assigned for different biomolecules. In the field of

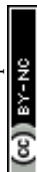
microbiology such information is deemed useful for classification of different bacterial species or understanding biochemical changes in bacteria due to external perturbations in the environment.<sup>73,75</sup> This paves the way for monitoring other perturbations like diseases in more complex living systems like human beings.<sup>76–78</sup> Any disease can also be considered as a perturbation to the native composition of these biomolecules in different tissues and body fluids.

Every disease is associated with a change in biochemistry. This principle is the central doctrine for modern medicine. The biochemical changes can either be a cause for disease manifestation or may be a consequence of the disease itself. Since Raman spectroscopy probes bond vibrations, the spectra generated are very molecule-specific. Any change in the concentration or the conformation of these biomolecules as a result of a disease, however subtle, is reflected in a Raman spectrum. This principle has formed the basis for a Raman-based diagnostic tool. Though Raman scattering is a weak phenomenon, technological advancements over the past few decades have led to shorter acquisition times and improved signal-to noise ratio, making the use of Raman spectrometers in a clinical setting easier. The clinical practicality has also been enhanced by the advent of fibre-optic probes for delivery of light as well as collection of Raman scattered photons. The use of advanced chemometric methods for handling Raman data sets has also made the diagnosis and classification process user-friendly.<sup>79</sup>

Different methods are available that help the user in deriving meaningful information from a complex spectral dataset. Principal component analysis (PCA) is one such technique that reduces the spectra into a defined number of principal components (PCs) that account for spectral variance. A detailed analysis of cluster analysis and PCA in spectral data analysis is demonstrated by Bonnier and Byrne.<sup>80</sup> PCA is an unsupervised method and hence cannot be used for classification and prediction purposes. Hence, supervised methods like Support Vector Machines, Artificial Neural Network, Linear Discriminant Analysis *etc.* are employed. A concise description of different chemometric methods used for Raman spectral data analysis has been given in the review by Butler *et al.*<sup>81</sup>

In the following sections, we discuss examples of how Raman spectroscopy has been used to understand some pathological conditions of the human body like cancer, infection and inflammation.

**Cancer.** Of the several diseases and pathological conditions that have been studied using Raman spectroscopy, cancer tops the list. In 2015, around 8.8 million deaths were due to cancer. As is true for any disease, early and rapid diagnosis of cancer is of utmost need. Traditionally, cancer diagnosis occurs through histopathology which requires a biopsy to be taken from the suspected area. This method of detection is highly subjective and requires the availability of a trained pathologist. Other methods of diagnosis include imaging based modalities like Positron Emission Tomography (PET) scan, Magnetic Resonance Imaging (MRI), Computed Tomography (CT) *etc.* However, these methods are difficult to be included in intra-operative procedures as most of these techniques require



extensive labelling. Cancer, like most other diseases, is accompanied by distinct modulations in biochemistry which are expected to be reflected in the Raman spectrum. Consequently, Raman spectroscopy has been used for the detection of various types of cancers. Kopec *et al.* used a combination of Raman and AFM imaging to determine the biochemical composition as well as the mechanical properties of blood vessels in a tumour mass of human breast tissue. They found an increase in glycogen and lactic acid in breast cancer patients coupled with an increase in the collagen–fibroblast–glycocalyx network.<sup>82</sup> An insight into how Raman spectroscopy can function as a tool for personalised radiation therapy was given by Van Nest *et al.* They used human breast adenocarcinoma xenografts after radiation therapy and found spectral changes linked to different biomolecules like proteins, lipids and nucleic acids on a temporal scale, highlighting the potential of Raman spectroscopy for assessing radiation-responses of tumours.<sup>83</sup> The use of cancerous cell lines as a platform to differentiate invasive *vs.* non-invasive breast cancer on the basis of Raman spectral markers was done by Chaturvedi *et al.* The Raman bands at 1447 cm<sup>-1</sup> (CH<sub>2</sub> bending from lipids), 1003 cm<sup>-1</sup> (phenylalanine) and 876 cm<sup>-1</sup> (tryptophan) were found to be higher in invasive cell lines *vis-à-vis* the non-invasive ones. Their work is in agreement with other reports that have correlated increased lipid levels to more aggressive and metastatic cancers and, therefore, a poor prognosis.<sup>84</sup> The work of Kim S. *et al.*, shows how cancer detection can be improved by coupling Raman spectroscopy to other methods like pH sensing. It is known that pH levels in cancerous tissues are much lower than that of normal tissues. They exploited the chemical information revealed by Raman spectroscopy as well as the metabolic information obtained by the pH sensor for improving the specificity of cancer diagnosis. Their study was done on human breast tissue samples and both the Raman and pH sensors were fibre-optic probes.<sup>85</sup> A recent study undertaken by Winnard *et al.*, has beautifully captured the potential of Raman spectroscopy to characterise organ-specific metastatic lesions at a molecular level, to give insights into metastatic progression. A combinatorial approach of Raman spectroscopy and metabolomics was employed in their study.<sup>86</sup> In another study, stromal adaptations occurring in pre-metastatic lungs that were primed by breast cancer were analysed using Raman spectroscopy. This work was done in mouse models where the mice were xenografted with human breast cancer cells of varying metastatic potentials. Changes in the extra-cellular matrix of pre-metastatic lungs, such as increase in collagen (859 cm<sup>-1</sup>) and proteoglycan (1061 cm<sup>-1</sup>) were identified and this was a direct correlation of the metastatic potential of the breast cancer cells used.<sup>74</sup>

Desroches *et al.*, utilized the higher wavenumber Raman signatures for designing, developing and testing of an *in situ* intraoperative system for brain cancer detection. They engineered this Raman-based system into a commercially available biopsy system for analysis before the tissue was extracted. Using this approach, they could achieve sensitivity and specificity of brain cancer detection in a swine model system, upto 80 and 90% respectively.<sup>87</sup> The reprogramming of lipid biosynthesis

and proteins in high-grade medulloblastoma was demonstrated by Abramczyk and Imiela very recently. Medulloblastoma is a term given to high grade cancers of the central nervous system. Such samples showed a higher content of  $\beta$ -sheets and lower levels of  $\alpha$ -helix when compared to normal tissues. The ratio of the Raman intensities at 2930 and 2845 cm<sup>-1</sup> was also indicative of cancer progression. For all the cancerous samples, this ratio had a value of 1.99 in comparison to the value of 1.45 for healthy tissue.<sup>88</sup>

Murali Krishna's group demonstrated the use of Raman spectroscopy for identification of sites in the oral cavity that have a higher propensity to become cancerous even before the manifestation of clinical symptoms of oral cancer recurrence. This study was performed on 99 patients of oral cancer and spectra were recorded from the cancerous sites as well as contralateral normal mucosa.<sup>89</sup> Nasopharyngeal cancer has also been diagnosed using Raman spectroscopy by Ming *et al.* in 2017. They demonstrated the use of a fibre-optic Raman probe for *in vivo* surveillance of patients suffering from nasopharyngeal cancer. This form of cancer is often very difficult to diagnose owing to the deep anatomical location. This study had grouped patients under three categories – normal, cancerous and post-radiation. All experiments were performed using a 1.8 mm probe, which is currently the smallest probe used for Raman-based diagnosis. A specificity of ~96% was observed in distinguishing normal, cancerous and post-radiation patients.<sup>90</sup>

Almond *et al.* utilised Raman spectroscopy and translated it into an endoscopic tool for the early detection of dysplasia in patients suffering from Barrett's oesophagus. A high-grade dysplasia (HGD) often results into oesophageal malignancy. Tissues from a cohort of 62 patients were taken for the analysis and sensitivity and specificity of 86 and 88% respectively were obtained for detecting HGD and adenocarcinoma.<sup>91</sup>

Another useful application of Raman spectroscopy in cancer research was highlighted by the work of Moradi *et al.*, where they tested the potential of this method to discriminate between human ovarian cancer cells that were sensitive or resistant to cisplatin. An increase in the relative concentration of proteins and glutathione was observed in the resistant cells compared to the sensitive ones.<sup>92</sup> The ability of Raman spectroscopy-based tool to serve an intra-operative device for detecting distant invasive grade 2–4 gliomas was shown by Jermyn *et al.* in 2016. They found that Raman spectroscopy could detect as few as 6 cancer cells per 0.0625 mm<sup>2</sup>, which was much better than traditionally used methods like MRI. All the Raman measurements were performed during brain tumour resections.<sup>93</sup> A detailed review on intra-operative Raman spectroscopy has been given by Brusatori *et al.*<sup>94</sup>

The use of body fluids like blood plasma/serum, urine, tears *etc.* can pave way for a diagnostic method that is minimally invasive. Body fluids can be obtained easily and therefore repeated samplings can be done. Unlike tissues and cells, bio-fluids can be used for routine monitoring as well as under intra-operative conditions. They contain several biomolecules and therefore can be very good indicators of the health status of a person or the underlying disease. The major limitation in using bio fluids for Raman spectroscopy is the relatively low



concentrations of biomolecules when compared to tissues and cells. This problem can be easily circumvented by the use of Drop Coated Deposition Raman Spectroscopy (DCDRS) where the biomolecules are concentrated.<sup>95–97</sup> Harvey *et al.* in 2008 demonstrated the use of urine for the detection of prostate cancer where increased proteins and nucleic acids were associated with the cancerous cells.<sup>98</sup> The work of Corsetti *et al.* demonstrated the efficacy of Raman spectroscopy in distinguishing hormone (androgen)-resistant prostate cancer cells. Bands specific to amino acids like arginine were found to be diminished in the cancerous cells whereas those with contributions from phenylalanine and tyrosine were found to be increased. The androgen-resistant cancer cells also had increased DNA and Amide III bands when compared to the sensitive cells.<sup>99</sup> O'Malley *et al.*, in their study, have highlighted Raman-based lipid quantification to be of diagnostic value for prostate cancer. Increased lipid amounts are directly correlated to the tumour stage.<sup>100</sup>

The use for serum-based cancer diagnosis has also been explored for different types of cancers. Ovarian cancer was studied using a combined approach of ATR-FTIR and Raman spectroscopy on human ovarian cancer patients by Owens *et al.* in 2014.<sup>101</sup> The basis for differentiation using Raman spectroscopy was the spectral shifts in the Raman bands at 711  $\text{cm}^{-1}$  (DNA) and 913  $\text{cm}^{-1}$  (sphingolipids).

**Infectious, inflammatory and other diseases.** While a plethora of information can be found for Raman-based cancer diagnosis, infectious and inflammatory diseases also require significant attention since they constitute a large percentage of the global burden on the health care system.

Infection with the dengue virus claims several lives especially in the developing countries. Current methods for diagnosis include measuring serum IgG and IgM levels and the Ns1 antigen. Mahmood *et al.*, demonstrated the use of Raman spectroscopy for profiling dengue-associated biochemical changes occurring in the blood plasma of 17 infected patients. Changes were seen mostly in the Raman bands associated with lipids (608  $\text{cm}^{-1}$ , 875  $\text{cm}^{-1}$ , 1297  $\text{cm}^{-1}$ , 1719  $\text{cm}^{-1}$  and 1736  $\text{cm}^{-1}$ ) which increased with infection. The peak intensity of the Amide III band at 1239  $\text{cm}^{-1}$  was also higher in the infected samples. An increase in the Raman band at 1409  $\text{cm}^{-1}$  was also observed which can be assigned to IgG. The method showed specificity and sensitivity of 95 and 98% respectively.<sup>102</sup> Dengue-associated spectral changes were also studied by Bilal *et al.* and they found the peaks for lactate at 750, 830 and 1450  $\text{cm}^{-1}$  to be higher in intensity in all 50 dengue patients when compared to 20 healthy controls. They also spiked the serum of healthy controls with lactate in a controlled manner and found that indeed the abovementioned peaks were due to lactate.<sup>103</sup> Ahmed's group from Pakistan has tried to develop a Raman-based diagnostic method for dengue using several multivariate approaches like Support Vector Machines (SVM), random forest method, *etc.*<sup>104</sup>

The same group has also conducted Raman studies on human malarial samples and found 10 distinct spectral signatures that could differentiate malarial samples from healthy controls as well as dengue samples and patients suffering from

non-malarial fever. A multivariate model based on partial least square (PLS) regression was used and the accuracy of detection was 86%.<sup>105</sup>

Malaria-induced changes in the spleen of mice were evaluated using Raman spectroscopy by Frame *et al.* and they found an increase in the heme-associated Raman bands. This directly correlated to the hemozoin detection with peaks at  $\sim 1370$ , 1529, 1588, and 1628  $\text{cm}^{-1}$  which matched closely with the reference spectra.<sup>106</sup> A comprehensive review on the potential of vibrational spectroscopy to diagnose and study malaria and the malarial parasite was written very recently by Perez-Guaita *et al.*<sup>107</sup>

One of the major causes of death in developing countries is tuberculosis (TB). In India alone, around 400 000 people die annually because of TB infections. While tuberculosis is mainly pulmonary in origin, extra-pulmonary TB also exists and is more difficult to diagnose. Tuberculous meningitis is one of the forms of extra-pulmonary TB and requires culturing of *Mycobacterium* for a precise diagnosis. Satyavathi *et al.*, used Raman spectroscopy on cerebrospinal fluid of patients suffering from tuberculous meningitis and found that the silicate Raman signature from *M. tb* positive cases was a very useful indicator of the underlying infection.<sup>108</sup>

Another leading cause of mortality in the Intensive Care Units of hospitals is sepsis. Sepsis can be triggered by infection of any kind and results in an uncontrolled immune response known as the Systemic Inflammatory Response Syndrome (SIRS). SIRS, however, can be a manifestation of other conditions like burns, trauma *etc.* Presently, no reliable marker exists that can efficiently distinguish SIRS from sepsis. In order to achieve this, Raman spectroscopy combined with multivariate approaches was used by Popp's group and they found that SIRS could be distinguished from sepsis with a predictive accuracy of 80%.<sup>109</sup>

Different inflammatory conditions like ulcerative colitis have also been studied using Raman spectroscopy. Ding *et al.* investigated the colon mucosal composition *in vivo* using an endoscope-coupled Raman probe in patients suffering from ulcerative colitis and age and BMI-matched healthy controls. A decrease in the phosphatidylcholine (720  $\text{cm}^{-1}$ ) and total lipids (1303  $\text{cm}^{-1}$ ) was observed in the inflamed colon tissue. The degree of lipid unsaturation, determined by calculating the ratio between Raman bands at 1657 and 1447  $\text{cm}^{-1}$ , was found to be high in the patient group. These differences were also seen when active colitis patients were compared with the quiescent ones.<sup>110</sup> Their results were corroborated by the work of Addis *et al.*, which also showed reduction in phospholipids to be of diagnostic value for ulcerative colitis.<sup>111</sup>

Inflammatory bowel disease (IBD) was also diagnosed by Mahadevan-Jansen's group where a colonoscopy-coupled fibre optic Raman probe was used. IBD includes ulcerative colitis (UC) and Crohn's disease (CD) and the motive in this study was to determine the spectral signatures of both UC and CD.<sup>112</sup>

Inflammation in the oral cavity as a result of periodontal disease was also assessed using Raman spectroscopy by Camerlingo *et al.* They used gingival crevicular fluid from patients with periodontal disease and found an increase in the



carotene content to be diagnostic of the condition ( $1537\text{ cm}^{-1}$ ). They also studied the chemical and structural changes occurring in the periodontal ligament after orthodontic force was applied and found changes in the secondary structures of proteins (Amide I band) to change with application of the force.<sup>113</sup>

In addition to the above mentioned inflammatory diseases, Raman spectroscopy has been extensively used to study and diagnose neurodegenerative diseases like Alzheimer's disease (AD). Till date, there is no cure for such diseases primarily due to their delayed diagnosis which occurs when the extent of brain damage is already quite pronounced. In fact, a definitive diagnosis can only be established post-mortem. It is now known that changes in the brain of such individuals begin much earlier than the appearance of the first symptoms. This can pave way for an early diagnosis which may make therapeutic interventions more successful. Protein misfolding and aggregation are central to most neurodegenerative diseases. Raman spectroscopy is an excellent tool to study protein conformation and therefore, has significant potential in diagnosing such diseases. This has been efficiently shown in the work of Paraskevaïdi *et al.*, where blood plasma was used to successfully discriminate Alzheimer's disease from healthy controls as well as from patients suffering from Dementia with Lewy Bodies (DLB). It is important to highlight that patients suffering from DLB are often misdiagnosed as AD and hence the choice of the patient cohorts is well justified. The same study also showed that patients suffering from early and late stages of AD could be differentiated from healthy controls with a sensitivity of 84%.<sup>114</sup>

In another study, Ryzhikova *et al.*, demonstrated the use of Raman microspectroscopy on blood serum for diagnosis of AD using a cohort of 20 AD patients, 18 from other neurodegenerative diseases and 10 healthy controls with specificity and sensitivity of ~95%.<sup>115</sup>

## (II) Raman spectroscopic studies on materials

Preceding discussions revolved around the use of Raman spectroscopy in the field of biology. It should be emphasized that biomolecules have a low Raman cross-section and as a consequence show weak Raman signals. Probing such weak scatterers has been made possible due to the introduction of Raman microscope in the early 70s.<sup>7-9</sup> The other advantages which Raman spectroscopy offers over other analytical tools are its non-invasive and non-destructive nature. This has resulted in its prolific use in the field of materials such as studies on carbon allotropes, explosives, semiconductors, pharmaceuticals, art, painting and archaeology. Raman spectroscopy has been an indispensable tool for characterization of different allotropes of carbon. Ranging from 3D diamond or graphite to 2D materials such as graphene or 1D materials like carbon nanotubes (CNT), Raman spectroscopy provides information pertaining to lattice vibrations, defects, orientations *etc.*<sup>116,117</sup> For instance, single walled and multi walled CNTs can be distinguished using this technique. The radial breathing modes which are unique to single walled CNT are absent in multi walled CNT.<sup>118</sup> Additionally, various  $\text{sp}^2$  carbon can also be

differentiated.<sup>119</sup> One of the more recent interests on carbon allotropes has been in graphene research. Here, Raman spectroscopy has proved to be an integral tool to determine number of layers of graphene along with orientational information, edges, strains, defects *etc.*<sup>120,121</sup> Enhancement of Raman signals of analytes using carbon materials has also been explored.<sup>122-125</sup> In the field of explosives detection, surface enhanced Raman spectroscopy has been employed for trace level detection.<sup>126,127</sup> Furthermore, studies on feasibility of Raman spectroscopy for stand-off detection have also been explored.<sup>128,129</sup> The subsequent sections provide a different aspect of this technique to demonstrate the usefulness of Raman spectroscopy towards studies on explosives. The first example shows how Raman spectroscopy can be used to monitor melt cast explosives formulations. In the second case, detection of concealed materials in non-metallic containers using depth sensitive Raman spectroscopy such as spatially offset Raman spectroscopy (SORS), transmission Raman and universal multiple angle Raman spectroscopy (UMARS) has been demonstrated.

**(a) Raman spectroscopy as an effective tool in process monitoring of explosives formulation.** The first use of explosive dates back as early as 220 B.C. when some Chinese alchemists were trying to separate gold from silver and met with accident. The material was 'black powder', which is a mixture of charcoal, sulphur and potassium nitrate.<sup>130,131</sup> An explosive in the literature is defined as 'a substance which when suitably triggered, releases a large amount of heat and pressure by way of a very rapid self-sustaining exothermic decomposition reaction'.<sup>130</sup> The temperature of such a reaction is approximately equal to the temperature at the Sun's surface  $\sim 5000\text{ }^\circ\text{C}$ . Secondary explosives are a class of explosives which are comparatively insensitive to heat, friction and impact. They require more energy to initiate the detonation process. Typical secondary explosives are 2,4,6-trinitro toluene (TNT), pentaerythritol tetranitrate (PETN), cyclotrimethylenetrinitramine (RDX), tetrahexaminetetranitramine (HMX) and more recently 2,4,6,8,10,12-hexanitro-2,4,6,8,10,12-hexaazaisowurtzitane (HNIW or CL-20), and mixtures of these explosives in different proportions. Explosives in a nutshell, have been described as a 'devil' during war and considered as an 'evil' during processing, handling, transportation and storage, have proved to be an 'angel' due to their tremendous impact on economy and industries and their innumerable applications in all walks of life.<sup>130</sup> Most of the explosives formulations are composed of binders, oxidizers and explosives. For instance, stabilizers are added to ester based rocket-propellants for the formation of a stable product.<sup>132</sup> Raman spectroscopy technique can be used as an effective tool for non-destructive and rapid evaluation and monitoring of the individual components of the formulation. Additionally, the depletion of the components over time can be monitored during ageing studies of stabilizers. Similarly, different polymorphic forms of explosives can also be determined using Raman spectroscopy.<sup>133,134</sup> Recent trends in the development of high explosives molecules have shifted towards engineering co-crystals of explosives molecules to distinct properties such as greater detonation velocities without compromising with impact or friction sensitivities.<sup>135-137</sup> Raman



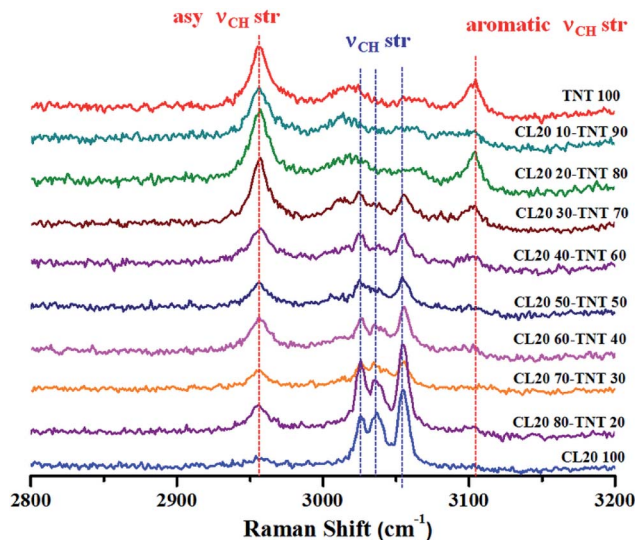


Fig. 6 Raman spectroscopic signatures at higher wavenumber range for CL-20 and TNT melt cast composition at different weight percentages. As the % of one of the component increases, intensity of the corresponding Raman peaks also increases. Data was obtained using 514.5 nm Ar<sup>+</sup> ion laser with 1.25 mW power and 10 s acquisition time.

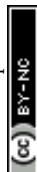
spectroscopic technique can be used to monitor the structures of such co-crystals.<sup>137</sup>

High explosives are used in military applications to achieve the terminal effect *i.e.* destruction of tactical and strategic targets. Typical examples of industrial explosives are blasting gelatine and ammonium nitrate slurries. However, high explosives are useful in civilian applications as advanced demolition devices (ADDs) for demolition of structures with precision and least collateral damage. Most of the ADDs comprise of composition-B (RDX/TNT 60-40) or similar compositions. These are usually manufactured using melt cast technology.<sup>138</sup> TNT forms the work horse for such formulation owing to its low melting point ( $\sim 80^\circ\text{C}$ ) which allows the ease of mixing and homogenization of other explosive ingredients. Therefore, the entire process requires constant monitoring to ensure thorough mixing and formation of a uniform melt. Explosives are aliphatic, aromatic or nitramine based organic molecules which are rich in Raman signatures. Since melt cast compositions are physical mixtures, the structures of the molecules are unaltered (in most of the cases). Therefore, Raman spectroscopy can be a potential tool for *in situ* monitoring of such processes. Fig. 6 depicts an example of an explosives formulation comprising of CL-20 (nitramine class) and TNT (nitroaromatic class) at different weight ratios. Raman spectra were acquired for different compositions at higher wavenumber regions as these molecules have unique spectral features which can be attributed to C–H stretches.<sup>139–141</sup> Fibre optics based Raman spectroscopy can be employed for online monitoring of such processes which least human intervention. Although a less explored area, Raman spectroscopy can be a useful tool in the area of explosives research.

#### (b) Deep Raman spectroscopy for detection of sub-surfaces.

Historically, Raman spectroscopic studies were carried out

using backscattering ( $180^\circ$ ),  $90^\circ$ ,  $135^\circ$  and transmission ( $0^\circ$ ) geometries respectively. The choice of illumination to collection geometry is decided by the application and the type of the sample to be studied. For better spatial resolution, confocal backscattering geometry is a preferred choice which forms the basis of Raman microspectroscopy. On the contrary, for transparent and liquid samples,  $90^\circ$  and  $135^\circ$  are the preferred geometries. These geometries work fine for pure materials. Furthermore, Raman microspectroscopy provides spatial information of 2D heterogeneous materials up to a diffraction limited spot size ( $\lambda/2$ ). However, when the material of interest is layered or when it is concealed by other layers, backscattering geometry may not yield complete structural information due to the low scattering volume of backscattering geometry (depending upon the depth of field). Such situations demand for exploration of new methods for studying layered materials. In 2005, a new method was introduced for obtaining sub-surface information of layered materials and was termed as spatially offset Raman spectroscopy (SORS).<sup>142</sup> The principal of this method is to introduce an offset with respect to the illumination followed by collection of the Raman signals. The authors noted that at no offset, there is a strong propensity of Raman signal of the outer material. As the collection offset is increased, Raman signal from the inner layer becomes prominent. The intensity ratio of the inner vs. the outer layer improves significantly with offset. This technique offers discrimination of Raman signals of layered materials.<sup>143–145</sup> Since this technique depends upon the point of illumination, the depth of penetration may be restricted upto a few centimetres. However, one can also use transmission geometry to obtain bulk signature of materials.<sup>146</sup> This method has been successfully employed in pharmaceutical sector for quality control.<sup>147–151</sup> The next challenge is in the case of materials which are hidden beyond a few tens of millimetres. For instance, in the case of strongly scattering media, such as biological tissues, retrieval of Raman signal from a tissue buried under overlying layers non-invasively can be a daunting experimental task. If our target material is hidden deeper inside a scattering sample, the Raman signal will be dominated by that of the surface layer. Therefore, new Raman-based methods need to be developed for depth-specific measurements of such materials. Our laboratory has focused its attention towards exploration of novel methods to address the challenge. Understanding of photon migration studies shows that the excitation and Raman photons inside turbid media undergo multiple scattering events.<sup>152–156</sup> This leads to randomization of photons. As a consequence, Raman photons from any angle or geometry can be collected to identify substances lying deep within such media. This technique has been termed as Universal Multiple Angle Raman Spectroscopy (UMARS). Therefore, the collection geometry is independent of the illumination geometry. Additionally, Raman signals are collected from all sides and angles around the material under study. Fig. 7 shows the schematic of different geometries used in Raman spectroscopy. The subsequent sections will describe examples of SORS, backscattering, transmission and UMARS methods.



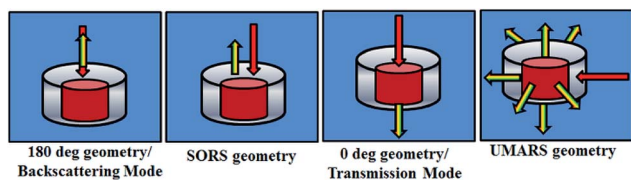


Fig. 7 Schematic of different experimental geometries employed for obtaining depth specific Raman spectroscopic signatures. Left panel to right depict backscattering, spatially offset Raman spectroscopy (SORS), transmission and Universal Multiple Angle Raman Spectroscopy (UMARS) geometries respectively.

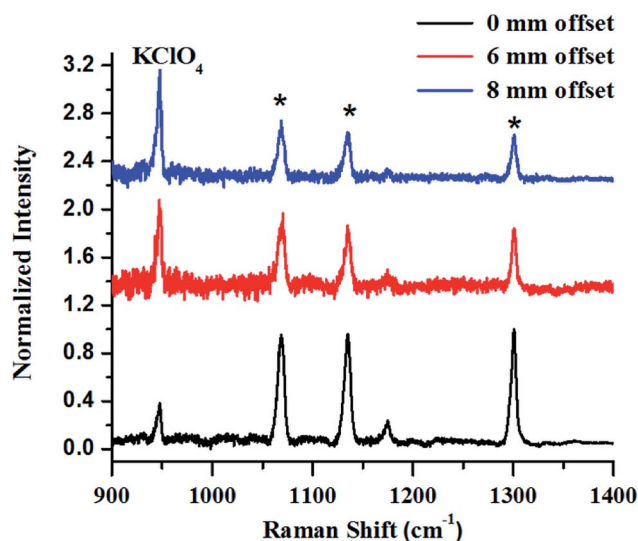


Fig. 8 (a) Raw SORS signals from KClO<sub>4</sub> powder filled in 25 mm square HDPE container with a wall thickness of 3 mm with 0 mm (a), 6 mm (b) and 8 mm (c) offsets respectively. Laser beam was focused on to the container. As the offset is increased to 8 mm, the signal of the sub-surface material improves.

The first example describes SORS for detection of KClO<sub>4</sub>, an oxidizer used towards making explosive formulations (Fig. 8). In this study, KClO<sub>4</sub> was taken inside a 3 mm thick high-density polyethylene (HDPE) container. Experiments revealed that a certain offset from the excitation point was necessary to discriminate the Raman signals of the content from that of the container. It may be noted that at no offset, the signal from the container was pronounced (peaks with asterisk). As the offset was increased, the Raman signal intensity of KClO<sub>4</sub> improved with respect to the container, while that of the container did not show any appreciable increase.

In the next set of studies, experiments were performed using transmission and backscattering geometry on the same material by using a focused beam of incident light (Fig. 9). We noted, however, that the transmission geometry yielded better content signal quality for the same container wall thickness (3 mm), as shown in the figure, and the signal from the container was remarkably suppressed. In this case the signal was collected at the back of the 25 mm thick square container, opposite to the point of excitation. This is possible in transmission geometry

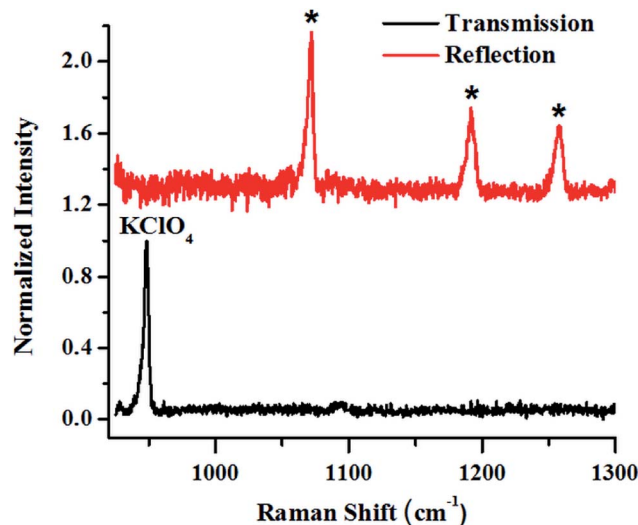


Fig. 9 Raman signals of layers of KClO<sub>4</sub> explosive in a 3 mm thick HDPE container obtained in reflection (backscattering) and right transmission mode.

because of the long migration paths of Raman photons in non-absorbing powder media.<sup>147</sup>

From the above-mentioned geometry-specific Raman experiments it is clear that backscattering mode yielded appreciable Raman signals of the top layer. This was overcome by using spatial offsets. However, transmission geometry provided Raman signals with even less interference from the top layer signals. It should be noted that transmission geometry is limited by the path length of the system due to the fact that after a certain depth, the photons would attenuate and low Raman signals can be acquired beyond that. In the cases where the material thickness exceeds a few centimetres, retrieving Raman signatures from material located at a depth will be a challenge by utilizing the above-mentioned geometries. It is worth mentioning that direct evidence of the extra pathlength for Raman photons from deeper layer can be obtained *via* time resolved Raman spectroscopic techniques.<sup>142</sup> For instance, Peterson *et al.*, have reported detection of explosives concealed behind opaque, diffusely scattering materials using picoseconds time resolved Raman spectroscopy.<sup>157</sup>

Our laboratory has demonstrated UMARS for the first time where we have obtained Raman signals in strongly scattering samples beyond a few centimetres.<sup>125,158–162</sup> As a demonstration, imaging experiments using the UMARS technique were conducted for bilayered solid compounds; in this case, ammonium nitrate (AN) was taken as a top layer and *t*-stilbene (TS) as a bottom layer. The excitation fibre was placed at the interface of the two layers, collection fibres 1–5 were placed around the bottom layer, TS, while fibres 6–10 were placed around the top layer, *i.e.*, AN. The experimental arrangement is shown schematically in the inset of Fig. 10. Before recording the image, Raman spectra were recorded for the separate components. Peaks centred at 992 cm<sup>-1</sup> and 1044 cm<sup>-1</sup> correspond to TS & AN, respectively. TS, being a very strong Raman scatterer, showed a higher peak intensity as compared to AN. When the



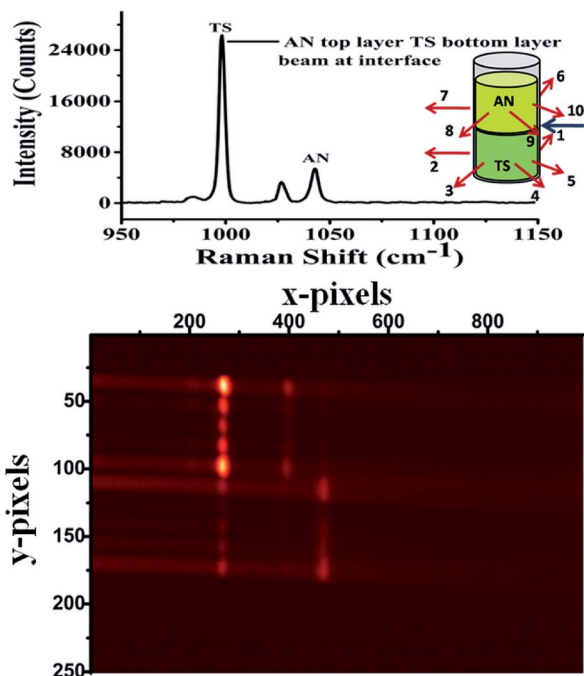


Fig. 10 UMARS imaging of bilayer solid powders ammonium nitrate (AN) & *trans*-stilbene (TS) using 10 optical fibres (830 nm diode laser, laser power: 300 mW, acquisition time: 100 s). The imaging experiments have been carried out by keeping the excitation beam at the interface of the two layers while fibre numbers 1 through 5 were placed around the bottom layer while fibres 6 through 10 were placed around the top layer. TS being a strong Raman scatterer showed intense spots (fibres 1–5, top) on the CCD as compared to AN. Identification of the spatial position is possible with UMARS imaging. Y axis: projection of 10 separate collection fibres on different pixel rows. Spectrum at the top of the figure corresponds to the sum of all y pixels (full vertical binning).

CCD was used in imaging mode, different collection fibres produced separate Raman spectra at different heights of the CCD. Y axis of the image shows the projection of 10 separate collection fibres on different pixel rows. Spectrum at the top of the figure corresponds to the sum of all y pixels, i.e., a full vertical binning of the signal. Since fibres 1–5 were placed around TS, those spots appeared on the top section of the CCD, while the bottom section contained mostly spots of AN. It was observed that in addition to the AN signal, fibres 6–10 also collected some signals from TS. This indicates that TS photons migrated to the top layer and contributed to the Raman signal. These results show that the spatial locations of the species present in the layers could be identified based on the position (image) of the fibres. In this setup, spectra of the individual fibres are collected simultaneously and can be evaluated separately. This is an important attribute of UMARS, as spatial information will be very important in the case of medical diagnostics. As expected, Rayleigh scattering was strongest in fibres 1 and 5 and fibres 6 and 9; closest to the point of excitation. The Raman signal of the bilayer clearly shows the TS and AN peaks.

In order to ascertain the location of the inner layer, a special experiment was designed. An HDPE cuboid container

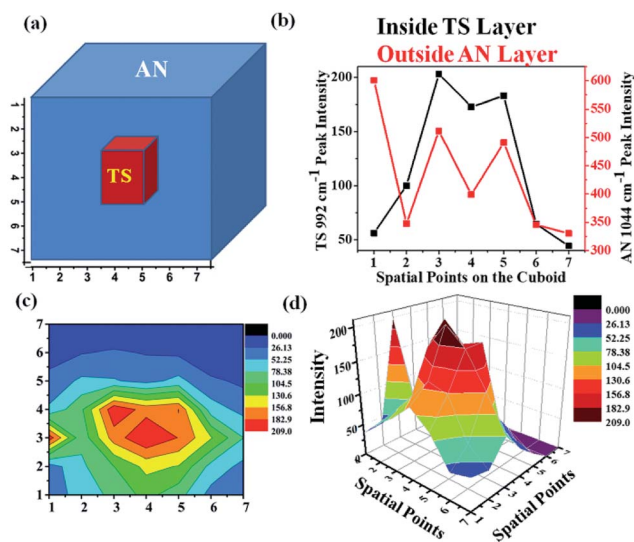
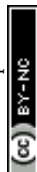


Fig. 11 Depth sensitive detection of bilayer solid powders ammonium nitrate (outer layer) and *t*-stilbene (inner layer) using UMARS where, (a) depicts the bilayer cuboid sample schematically, (b) is the Raman intensity variation of AN & TS as the signals were collected by keeping the y-axis constant ( $y = 4$ ) i.e., at the center of the cuboid and by varying the x at that y, i.e., from  $x = 1$  to  $x = 7$ ; (c) shows the contour of the bilayer mapped with respect to TS Raman intensity where red colours corresponds to maximum TS intensity, blue corresponds to a minimum value; and (d) shows the 3D surface plot of the variation of TS Raman intensity.

was taken and it was filled with AN as an outer layer and TS as an inner layer as shown in Fig. 11(a). The cuboid was divided into a  $7 \times 7$  grid. The excitation fibre was placed at each point in the grid, while the collection fibres were placed around the periphery maintaining the same plane as the input fibre. UMARS signals were collected from all the 49 points and the obtained spectra were analysed to retrieve depth specific molecular information. At first, the Raman spectral intensity variation of TS & AN were recorded by keeping the y-axis constant ( $y = 4$ ) i.e., at the centre of the cuboid and by varying the x at the y level, i.e., from  $x = 1$  to  $x = 7$  (Fig. 11b). It was observed that the Raman intensity of AN was strongest at  $x = 1$  while in the case of TS, the signal was weak at  $x = 1$ , followed by an increase in Raman intensity to reach a plateau and finally decreased. It was also noted from the spatial points 3 to 5, the intensity of TS remained more or less same. This implied that TS was present in higher amounts close to those spatial points as compared to other points. To get a clear picture, a contour and a 3D surface map was plotted for the TS Raman intensity ( $992 \text{ cm}^{-1}$  peak) for the  $7 \times 7$  matrix (49 spectral points) (Fig. 11c and d). This confirmed the presence of TS at the center of the cuboid. This example provides sufficient evidence of the capability of UMARS as a non-invasive, depth sensitive technique. Further experiments on phantoms and biological samples are being conducted in our laboratory to understand the potential of UMARS towards 3D imaging. We anticipate that UMARS would pave way towards Raman tomography for disease diagnosis in the future.



## Experimental

### Biomolecules

Bovine Serum Albumin (BSA), calf thymus DNA, D-(+)-glucose and cholesterol were purchased from Sigma Aldrich and used without further purification. Spectra were recorded from all the samples in powdered form using 785 nm laser and 50× L objective (N. A = 0.5) using Renishaw InVia Raman Microscope.

### Thermal denaturation of lysozyme

Hen egg white lysozyme (lysozyme) was purchased as lyophilized powder from Sigma and was dissolved in sterile MilliQ water to obtain a working concentration of 30 mg mL<sup>-1</sup>. For thermal denaturation studies, the protein solution was heated to a temperature of 85 °C for 8 hours. Spectra were recorded from the native and denatured protein solution using 785 nm laser and 50× L objective using Renishaw InVia Raman Microscope at room temperature.

### Liver and serum studies

6–8 weeks old healthy BALB/c mice were sacrificed and liver and blood were harvested. For collection of serum, blood was allowed to clot at 37 °C for 4 hours, followed by centrifugation at 4000 rpm for 5 min. For recording Raman spectra of liver, ~50 µm thin cryo-sections were used. All Raman experiments were conducted using Renishaw in Via system using a 785 laser and 50× objective (N. A = 0.75).

### Bacterial heat treatment

*E. coli* MG1655 K-12 strain was cultured in a shaking condition at 37 °C and 160 rpm for 8 hours. 1 mL of the culture was subjected to heat shock at a temperature of 85 °C for 15 minutes. The control and the heated samples were then washed with sterile water three times to remove any residual media components; ~3 µL of the bacterial sample was drop cast onto a magnesium fluoride coverslip before collecting Raman spectra. All Raman experiments were performed at room temperature using Renishaw in Via system using a 785 laser and 100× objective (N. A = 0.85).

### Data analysis

All spectra were processed using Wire 4.2, Origin 8.5 and MATLAB softwares. Statistical analysis (Mann–Whitney test) was performed using GraphPad Prism 6.

### Raman spectroscopic studies on materials

CL-20 and TNT were provided by DRDO laboratory for conducting the Raman experiments. The CL-20-TNT melt cast formulations were prepared in milligram quantities in the laboratory. Required quantity of TNT was first melted followed by addition of appropriate amount of CL-20 and homogenized using a wooden pestle. All chemicals used in SORS, reflection, transmission and UMARS experiments except ammonium nitrate (Nice Chemicals Pvt. Ltd.), were procured from Sigma Aldrich. The solid powder samples used in the experiments

were either filled in commercial high-density polyethylene (HDPE) containers, quartz cuvettes or glass vials of assorted sizes.

The Raman experiments on explosives formulation was conducted using Renishaw in Via system using a 514.5 nm Ar<sup>+</sup> laser. The power at the sample was 2.5 mW and the spectra were acquired for 10 seconds.

The SORS, reflection and transmission Raman experiments were carried out using the 830 nm laser diode. The spectrometer was a high resolution imaging spectrometer (Horiba Jobin Yvon, iHR320) fitted with a LN<sub>2</sub>-cooled CCD camera. Grating with a groove density of 1200 lines per mm was used for the experiments. In the case of UMARS experiments, the laser was delivered to the sample through an optical fibre of core diameter 400 µm and a numerical aperture of 0.22 and the Raman signals were collected using 10 fibres of the same core diameter and NA placed on the sample container sides at various angles. The ten collection fibres terminated in a vertical linear array, placed in front of a telescopic arrangement fixed to the entrance slit of the spectrometer. The power of the laser was maintained between 250–450 mW. The acquisition time for the experiments varied from 10 s to a few minutes.

The acquired data were processed using Wire 4.2 and Lab-spec softwares and were plotted using Origin 8.5 and Origin 9.1.

## Conclusion

Raman spectroscopy has proved to be a versatile tool in studying different diseases occurring because of the altered biochemistry in the body. Various advancements have been made in the field of Raman spectroscopy with respect to instrumentation and data handling methods. In this review, we have attempted to provide a glimpse of some of the research activities which are being pursued in our laboratory in the area of biophotonics, explosives detection and development of novel Raman techniques. We started by showing Raman spectra of pure molecules which comprise of the building blocks of cells and tissues. Examples of Raman spectroscopy of tissues are presented to understand the unique spectral features along with the complexity of the Raman signals. The second part of the review deals with Raman spectroscopic applications towards materials by taking explosives as a representative example. The last part of the review discusses the variants of Raman spectroscopy like SORS, transmission and UMARS for depth sensitive detection of layered materials. Finally, it is shown that UMARS imaging is possible and that chemical as well as spatial information of the materials lying underneath other materials can be clearly obtained using UMARS when the fibres are separately imaged on the CCD. The day is not far when Raman spectroscopy-based diagnostic methods will be routinely prescribed in clinics and public health care systems. One of the most anticipated uses of Raman spectroscopy is in clinical diagnostics. Most of the research on Raman is focused on biological research and imaging. It remains to be seen whether Raman spectroscopy would take off as portable equipment for point of care application. In his concluding remarks of his review paper, Prof Kiefer wrote, "Sometimes I am



asking myself what cannot be done with Raman spectroscopy – a wonderful analytical tool for science and technology which was discovered by Sir Chandrashekhar Venkata Raman and Krishnan already eight decades ago at the Indian Association for the Cultivation of Science in Calcutta, India. I am looking forward for many more years of significant advancements in this research field.<sup>163</sup>

## Conflicts of interest

There are no conflicts to declare.

## Acknowledgements

The authors gratefully acknowledge Department of Science & Technology, Defence Research & Development Organization, Council of Scientific & Industrial Research, Indian Institute of Science, Bangalore for their financial support. S. U. acknowledges the J. C. Bose fellowship from DST.

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