Label-free super-resolution stimulated Raman scattering imaging of biomedical specimens

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Abstract: Far-field super-resolution microscopy has unravelled the molecular machinery of 9 biological systems that tolerate fluorescence labelling. Conversely, stimulated Raman scattering 10 (SRS) microscopy provides chemically selective high-speed imaging in a label-free manner by 11 exploiting the intrinsic vibrational properties of specimens. Even though there were various 12 proposals for enabling far-field super-resolution Raman microscopy, the demonstration of a 13 technique compatible with imaging opaque biological specimens has been so far elusive. Here, 14 we demonstrate a single-pixel-based, combined with robust structured illumination, that enables 15 super-resolution in SRS microscopy. The methodology is straightforward to implement and 16 provides label-free super-resolution imaging of thick specimens, therefore paving the way for 17 probing complex biological systems when exogenous labelling is challenging. 18

19 1. Introduction

Far-field super-resolution imaging has emerged as a powerful tool in biology to unravel the 20 details of the complex molecular machinery at play at the nanoscale. However, the great majority 21 of super-resolution techniques are based on exogenous markers (fluorophores) that demand 22 chemical preparation protocols and further studies to determine cell viability and specificity to a 23 targeted molecule. Most importantly, fluorescence-based tools only report on the fluorophore 24 information – dynamical or structural – leaving open many fundamental questions on the other 25 outnumbering unlabelled molecular species: e.g., lipids and cholesterol molecular conformation 26 and local composition [1-3] within lipid domains have remained undetected in real cells, or 27 the local composition of the species forming membrane-less organelles which are currently 28 unknown [4, 5]. Therefore, Raman microscopies have emerged as ideal tools for probing 29 heterogeneous biological specimens [6], since they provide chemically resolved images using 30 the intrinsic vibrational properties of molecules, that is, a label-free method. Yet, reaching 31 fast super-resolution capabilities for opaque tissue imaging with Raman contrasts has remained 32 challenging [7]. 33

In the last decade, many attempts have been made to enable vibrational far-field super-resolution 34 in SRS. Computational super-resolution methods, exploiting structured illumination microscopy 35 (SIM), have been demonstrated for the spontaneous Raman contrast [8]. However, the usage of 36 an imaging spectrometer is not compatible with thick tissues, as the resolution enhancement is 37 only provided in one dimension, and the acquisition speeds are too slow for dynamic specimens. 38 Very recently, a solution to this issue has been put forward, but the imaging methodology is still 39 based on a wide-field geometry, potentially challenging to be applied in opaque specimens [9]. 40 Alternatively, coherent Raman microscopies (CRM) could provide fast acquisition speeds: 41 with the two most known contrast mechanisms being Coherent anti-Stokes Raman scattering 42 (CARS) [10] and stimulated Raman scattering (SRS) [11–13]. However, there are various 43 drawbacks that preclude biological specimens super-resolution imaging, in particular methods 44 exploiting coherent control of vibrational dynamics [14-25]. Furthermore, in CARS, interference 45 artifacts complicate chemical quantification analysis [26, 27]. In the case of the background-free 46

47 SRS process, the current mainstream is to exploit methods to control the dynamics of vibrational

energy levels, however using unconventional power levels that may be phototoxic for biological
specimens [20, 21, 23, 24, 28–31].

While combining computational super-resolution methods with SRS technology could over-50 come the above-mentioned issues, it cannot provide super-resolution capabilities in thick opaque 51 tissues. Generally, the mathematical framework of computational methods is based on wide-field 52 illumination which itself requires multi-pixel cameras. Unfortunately, wide-field cameras for 53 SRS are technologically challenging because of SRS' unconventional detection scheme: SRS 54 requires high-sensitivity radio-frequency lock-in amplifier (RF-LIA), which currently is only 55 reliable in a single-pixel scheme. Despite recent developments of multi-pixel RF-LIA [32, 33], 56 the pixel counts do not scale favorably for 1000's of pixels architecture needed in a camera. 57 Furthermore, wide-field illumination is not suitable for thick tissue imaging due to the lack of 58 sectioning capabilities: even if a camera did exist for using computational methods with SRS, it 59 would be challenging to use due to aberrations and out-of-focus light generated by the solvent or 60 the sample itself. 61

To address these challenges, we present a camera-less (aka single-pixel) chemically selective 62 super-resolution imaging methodology compatible with opaque thick biological specimens. 63 We are interested in demonstrating that this methodology indeed breaks the diffraction-limit 64 resolution barrier of SRS microscopy. We start by describing the mathematical model used in 65 the acquisition step, followed by a proof-of-principle aimed at demonstrating resolution beyond 66 the theoretical diffraction limit. We then finalize demonstrating that its sensitivity is compatible 67 with imaging biological specimens. Remarkably, the framework presented here has a simple 68 alignment procedure: it is simpler than conventional SRS microscopy, which demands overlap of 69 two tightly focused beams. 70

71 2. Concept

Specifically, we developed a single-pixel scheme (Fig. 1) compatible with computational super-72 resolution methods, therefore allowing for fast imaging capabilities exploiting SRS processes in 73 the form of stimulated Raman gain (SRG) (Fig. 1a). In our arrangement, a structured stationary 74 pump beam is shaped using a spatial light modulator (SLM) and is spatially and temporally 75 overlapped with a focused Stokes beam that scans over the sample (using a set of galvanometric 76 mirrors, Fig. 1b). After acquiring a series of SRS images with multiple structured illuminations 77 (Fig. 1c) the data is treated with algorithms based on standard SIM mathematical framework to 78 recover a super-resolved image [34, 35], as described below. We coin the method Single-pixel 79 blind-SIM SRS (or blind-S³ for short). 80

⁸¹ We describe the forward model of the acquisition procedure. In SRS, the signal detected ⁸² (ΔI_S), a modulation transfer between the Pump and Stokes beams, at one pixel location (of the ⁸³ un-processed image) is given by [12]:

$$\Delta I_S \propto \Im\{\chi^{(3)}\} I_P I_S \tag{1}$$

where $\Im{\chi^{(3)}}$ is the imaginary part of the complex-valued nonlinear susceptibility of the sample (related to the Raman cross-section), I_P and I_S the intensity of the Pump and Stokes beam, respectively. In the case of the proposed acquisition method in blind-S³, a static speckle pattern generated by the Pump beam spreads at the sample image plane where the Stokes beam is focused and scanned. To derive an image formation model, we assume a scalar approximation for the local intensity in one blind-S³ image:



Fig. 1. Principle of blind-S³. Schematic of the setup to achieve super resolution using the SRS process stimulated Raman Gain (SRG) (**a**) based on a single-pixel SIM scheme. Transverse (**b**1) and longitudinal (**b**2) planes of the scanning Stokes beam trajectory (red dash) over the stationary Raman-active specimen (blue) and structured Pump (green), in this case a speckle pattern. For every speckle realization, an SRS image is acquired forming an image stack that is passed to a SIM algorithm to reconstruct a super-resolved image (**b**3). (**c**) Conventional SRS, consisting in raster scanning co-propagating Pump and Stokes beams, is used as a control to demonstrate the increase in resolution when compared to standard imaging. Transverse (**c**1) and longitudinal (**c**2) planes of the Stokes and focused Pump beams (green and red dash) scanning trajectory over the stationary Raman-active specimen (blue).

$$\Delta I_S(x, y) \propto \iint \mathfrak{I}\{\chi^{(3)}(x', y')\} I_P(x', y') I_S(x - x', y - y') \, dx' \, dy'$$

$$\propto \left(\mathfrak{I}\{\chi^{(3)}\} I_P\right) \circledast I_S,$$
(2)

An image acquired with blind-S³ scheme obeys a forward model of the type $M_i = (O \times I)^2$ 90 I_{SIM}) \otimes I_{Stokes} , where M_i is an SRS image from a single speckle realization, O is the optical 91 response of the excited object (more precisely, $\mathfrak{I}{\chi^{(3)}}$), I_{SIM} is the spatial distribution of the 92 structured intensity at the Pump wavelength, I_{Stokes} is the effective PSF of the image formation 93 system, and [®] denotes a convolution operation. This means that each single image acquired 94 follow the standard forward models in computational super-resolution frameworks of incoherent 95 processes. We chose to work with non-sinusoidal SIM patterns in order to be compatible with 96 thick tissues: we use speckle patterns, since they are resilient in scattering specimens. While we 97 tested the methodology with two SIM algorithms using no prior knowledge on the structured 98 patterns I_{SIM} [34, 35], for the results presented we used the one described in Ref. [35]. In this 99 approximation, we disregard coherent effects as SRS processes are inherently phase-matched. 100

101 3. Experimental

102 3.1. Microscope design and details

Briefly, the output power of a femtosecond laser source (Coherent, Chameleon Ultra Vision, 103 800 nm, 80 MHz repetition rate, 150 fs pulse length) pumps an optical parametric oscillator (APE, 104 MIRA-OPO) that generates the Stokes beam, centered either at 1058 nm (Fig. 2, 3054 cm^{-1} 105 Raman-shift) or 1042 nm (Fig. 4, 2903 cm⁻¹ Raman-shift), and a small power fraction is used 106 as the Pump beam. The Stokes beam is spectrally narrowed using a combination of grating 107 (LightSmyth, T-1000-1040) and adjustable slit width for the purpose of increasing chemical 108 selectivity. The Pump beam is also spectrally narrowed in a pulse-shaper setup using two gratings 109 (LightSmyth Technologies, T-1400-800) and a digital micromirror device (DMD) placed in 110 the Fourier plane (a description of the methods using DMDs for SRS spectroscopy can be 111 found in Ref. [36]). The pump beam is amplitude-modulated at 1 MHz by an acousto-optic 112 modulator (AA Opto-electronic, MT80-B30A1,5 VIS). The specimen is z-displaced using a 113 piezo stage (Thorlabs, DRV517), and the signal generated by the sample is then collected by a 114 1.4 NA oil-immersion condenser, directed to a large-area detector (Thorlabs, DET100A2) and 115 demodulated by a lock-in amplifier (Zurich Instruments, MFLI). 116

We used two configurations for SRS microscopy. Regardless of the configuration used, both 117 beams are spatially and temporally combined at dichroic mirrors, whose location depends on the 118 modality of SRS in use, and focused by an objective (Nikon, Plan APO IR, 60x, NA=1.27). To 119 achieve the best compromise in terms of resolution enhancement (by having the Pump beam 120 as a structured illumination) and sensitivity (by having the Stokes beam as the demodulated 121 beam), we have designed a layout that allows us to quickly swap the direction of the Pump beam 122 between the conventional SRS or blind- S^3 configurations using a combination of a half-wave 123 waveplate and a polarizing beam splitter cube. For the blind- S^3 configuration, the Pump beam is 124 sent onto a SLM (Meadowlark Optics, HSP512L-1300) to modulate the wavefront with a random 125 phase. The SLM throughput is higher than 80% but it could be further enhanced by replacing the 126 SLM by engineered diffusers since they have no absorption. Galvanometric mirror scanners are 127 used to move either Pump and Stokes beams together (conventional) or Stokes only (blind-S³). 128 Typical average power measured before the objectives were 13 mW (conventional) and 41 mW 129 (blind- S^3) for the Pump, and 25 mW for the Stokes beams. However, we note that the energy 130 density levels used for blind- S^3 are inherently lower than the conventional SRS configuration: 131 we have estimated a 5 times lower effective energy densities (i.e. product of the energy densities 132 of the Pump and Stokes energy densities), taking into consideration the speckle envelope and 133 the longer integration time in the blind-S³ procedure, when compared to conventional method. 134 Finally, after imaging in conventional SRS and blind- S^3 , we image a large FOV to detect any 135 sign of phototoxicity such as shown in the wide FOV images of the biological specimens. 136

137 3.2. Sample preparation

Samples presented in Fig. 2 were prepared by drop-casting the polystyrene beads on a coverslip 138 and embedded in deuterated water to decrease the spectral congestion with the water vibrational 139 response background. The various diameters (and standard deviation) used were: 239 nm (6 nm, 140 PS Research Particles), 372 nm (10 nm, Polysciences, Inc.), 520 nm (16 nm, Thermo Scientific), 141 740 nm (22 nm, Thermo Scientific) and 990 nm (30 nm, Polysciences, Inc.). Mice brain slices 142 were kindly provided by Laurent Bourdieu and experimental procedures were conducted in 143 accordance with the institutional guidelines and in compliance with French and European laws 144 and policies. All procedures were approved by the 'Charles Darwin' Ethics Committee (project 145 number 26667). More precisely, 6-months old C57BL6 male mice were sacrificed, the extracted 146 brain was then stored overnight in a solution of 4% paraformaldehyde and finally rinsed in 147 phosphate buffer solution (PBS). Coronal slices of thickness 100 µm were then cut and stored in 148

¹⁴⁹ PBS. Prior to experiments, the slices were placed between two cover slips with a 120- μ m-thick ¹⁵⁰ spacer. HeLa cells (ATCC) were incubated with 400 μ M oleic acid, washed, fixed with 4% ¹⁵¹ paraformaldehyde, and stored at 4°C before imaging.

152 3.3. Resolution estimation

We assume that the theoretical transverse resolution results from the product of two focused Gaussian beams with two different wavelengths λ_p and λ_s for the Pump and Stokes wavelength respectively. Here, we use the Raman resonance 3054 cm⁻¹ and the Rayleigh criteria to asses the resolution limit of each beam: $\Delta r_P = \frac{1.22\lambda_P}{2NA} = 384$ nm and $\Delta r_S = \frac{1.22\lambda_P}{2NA} = 508$ nm for the Pump and Stokes beam respectively where NA = 1.27, $\lambda_P = 800$ nm and $\lambda_S = 1058$ nm. Therefore, the theoretical resolution limit is $\Delta r_{SRS}^{conv} = \frac{1}{\sqrt{\frac{1}{\Delta r_P^2} + \frac{1}{\Delta r_S^2}}} = 307$ nm for conventional SRS while it is

¹⁵⁹ $\Delta r_{SRS}^{blindS^3} = \frac{\Delta r_{SRS}^{conv}}{\sqrt{2}} = 217 \text{ nm for blind-S}^3.$

160 4. Results and discussion

4.1. Proof-of-concept of super-resolution capabilities beyond the diffraction limit

We first demonstrate the improvement in the transverse resolution, surpassing the diffraction-limit 162 of usual SRS microscopy. In order to evaluate the gain in resolution, we compare blind-S³ to 163 the conventional scanning methods. For conventional SRS, the theoretical transverse resolution 164 is $\Delta r^{Conv} = 307$ nm. This theoretical value is technically challenging to achieve with high 165 numerical aperture (NA) objectives in the near-IR because the wavelengths of the two beams differ 166 by hundreds of nanometers (spectral span necessary for fast quantification of lipids, proteins, 167 and nucleic acids in SRS microscopy), in opposition to the visible range where diffraction 168 limited performance has been reported [37]. Conversely, blind-S³ transverse spatial resolution 169 results from the doubling in resolution dictated by SIM and the speckle grain size limited by 170 diffraction, leading to $\Delta r^{blind S^3} = 217$ nm. To show the superior transverse resolution in 171 blind-S³, we imaged 239 nm-diameter polystyrene beads with the two modalities (Fig. 2a-b). 172 Clearly, conventional SRS (Fig. 2a) cannot resolve the beads transversely as the beads size is 173 smaller than the theoretical resolution limit. After multiple speckle pattern illuminations, we feed 174 the resulting images to a blind SIM algorithm to reconstruct a super-resolved image. Notably, 175 blind-S³ methodology (Fig. 2b) resolves several beads in the in-focus layer. The line profiles 176 reveal the distance between the centers of the beads (242 nm) which matches well to the distance 177 of close contact between two beads. We note that the effective region-of-interest (ROI) of blind- S^3 178 is modulated by the speckle envelope, hence decreasing the similarity of the two images in the 179 edges of the beads cluster, yet not affecting the resolution gain (see below). The present findings 180 show that the blind-S³ methodology goes beyond the fundamental far-field diffraction-limit 181 resolution of SRS microscopy, by improving the resolution $\geq \sqrt{2}$, without addition of exogenous 182 signal enhancers and using excitation energy densities lower than conventional SRS microscopy, 183 therefore decreasing probability of nonlinear phototoxic effects. 184

Remarkably, super-resolution in blind-S³ comes with intrinsic high z-sectioning capabilities. 185 In each illumination during the blind- S^3 procedure, an image is formed based on a wide-field 186 geometry model, that is, an object is convoluted with a linear point-spread function (PSF). In a hypothetical conventional wide-field SRS microscope using multi-pixel cameras, the excitation 188 beams would overlap a large volume. This would in turn reduce the sensitivity due out-of-focus 189 shot-noise, therefore deteriorating image quality and resolution due to the background noise 190 (shot-noise). Conversely, the blind-S³ methodology improves sectioning, without resorting to 191 cameras, as the nonlinear optical response is local in the longitudinal direction: because SRS 192 signals are only generated within the overlap region of the two beams, each SRS image does 193 not contain appreciable out-of-focus shot-noise. To demonstrate the sectioning capabilities of 194



Fig. 2. Proof-of-concept of blind-S³ capabilities to image beyond the diffraction-limit. Conventional (**a**) and blind-S³ (**b**) images of 239 nm-diameter polystyrene beads, and line profiles (**c**) showing the increase in transverse resolution of blind-S³ (line) compared to conventional methods (dash). (**d**) Conventional SRS (dash) and blind-S³ (line) sectioning capabilities characterization. All scale bars: 500 nm. Pixel dwell times are 73 μ s and 300 μ s for conventional and blind-S³ methods, respectively.

¹⁹⁵ blind-S³, we probed a thin film of oil (of few μ m) by scanning it in the longitudinal direction. ¹⁹⁶ Note that we collect the signal generated for each z-position on a ≈ 10 mm-wide detector, hence, ¹⁹⁷ not in a confocal geometry. Clearly, conventional SRS and blind-S³ give a peaked response which ¹⁹⁸ means that those two techniques have inherent longitudinal sectioning (Fig. 2d). Indeed, the ¹⁹⁹ conventional SRS microscope is able to show such z-sectioning capability, due to its nonlinear ²⁰⁰ longitudinal PSF.

Contrary to conventional methods in super-resolution microscopy, in blind-S³ it is not 201 straightforward to compare the reconstructed images with a "ground truth" object. This arises 202 from the fact that the FOV in blind- S^3 is modulated by the speckle envelope, which is smaller 203 than conventional SRS microscopy. Therefore, we devised another methodology to insure if 204 the reconstruction was indeed reaching super-resolution capabilities. We imaged commercially 205 available calibrated polystyrene beads of various sizes, which are well-known to aggregate and 206 form close-packed structures. Therefore, we can use the bead close contact distance as a proxy 207 for the bead diameter. We measured the close contact distances of several beads for several 208 sizes ranging from smaller than to several times the resolution limit, and also for two different 209 objectives with different NAs. Although the method is somewhat subjective, we were careful 210 to chose "spot centers" that had the smallest distances possible. Following this procedure, we 211 noticed that maximum spot-to-spot center were indeed limited by the bead size, that is, in the 212 360 nm bead diameter we did not see 240 nm spatial fluctuations. The outcome of this procedure 213 is shown Fig. 3 and the agreement between the nominal bead diameter and the retrieved diameter 214 therefore confirms that the features observed in the blind-S³ reconstructions indeed correspond 215 to physical features beyond the diffraction limit. 216

217 4.2. Biological compatibility

To demonstrate blind-S³ compatibility with biological specimens, we image standard cell lines and mouse brain tissues. Conventional SRS reveals several μ m-large droplets within the cell in the FOV (Fig. 4a). Close-up images show different cluster morphology (Fig. 4b), and increased resolution gain with blind-S³ from the line profiles of selected ROI (Fig. 4c). To demonstrate



Fig. 3. Transverse resolution analysis for blind- S^3 . Outcome analysis of the images of various close-contact beads pairs: We use close-contact distances as a proxy for the bead diameter. The inset shows representative images used for analysis, with the dashed lines representing some of the beads chosen for evaluation.

capabilities for aberrant and opaque tissues, we have further imaged highly scattering brain slices at 8 μ m-deep in the sample (Fig. 4d-e) with line profiles demonstrating increased resolution power of the myelin structures (Fig. 4f). The close-up images with super-resolution capabilities (Fig. 4e) reveal that the structure of the myelin in the tissue is actually not as symmetrically perfect as inferred from the low-resolution images. These results show that the method is compatible with thick tissue imaging, despite being completely opaque, a situation where using a hypothetical SRS widefield camera approach may fail due to background shot-noise.

We consider the effects of the idle speckle upon illumination of the sample. One potential 229 hazardous effect is due heating. While the Stokes is point scanning, there are other regions 230 of the speckle pattern that constantly illuminate the sample therefore potentially presenting a 231 phototoxic effect. Here, we assume the peak power of the speckle is too weak to induce nonlinear 232 photodamage (at least not observed in our experiments and shown by the large FOV images after 233 the acquisition procedure), and we consider what is the temperature rise due to the small, but 234 non-negligible, absorption of water at the pump wavelength. This is a safe assumption as the 235 speckle power used in the experiments is spread over a region that is over 10×10^{10} larger that the spot 236 size of the conventional SRS imaging system: this would represent a $100 \times$ weaker peak power. To 237 analyze the heat effects, we use a well-established heat propagation framework used in the context 238 of optogenetics [38] which have shown to be accurate in previous experiments. The simulation 239 was performed with a speckle illumination power of 1 W ($20 \times$ higher than the experiments) and 240 envelope size of 10 μ m. Since heat-diffusion is faster than the image acquisition, the speckle 241 grains are washed out and only the speckle envelope is important to be considered. Hence, we 242 plot the maximum of this thermal envelope in Fig. 5 for a illumination that stopped at 10 s and a 243 continuous one. One can see that the steady-state value saturates at $0.023 \text{ K W}^{-1}\text{MHz}^{-1}$, which 244 corresponds to temperature rise < 0.1 K in our experiments. Note that the recently reported 245 photothermal SRS signal [39] could be present, and therefore have a higher temperature rise 246 than due to the Pump speckle itself. However, in the simulations above, we consider the Pump 247 speckle envelope, which does not go through the thermal SRS effect as a whole. Furthermore, 248 this thermal SRS heat mechanism would be highly local and quickly dissipated as the overlap of 249



Fig. 4. Bio-compatibility capabilities of blind-S³ at reduced excitation energy densities. (a) Large FOV imaging of lipid droplets within HeLa cells (conventional SRS). Two zoomed-in ROIs (b) are depicted by dashed boxes with conventional SRS (left panels) and blind-S³ (right panels) methods, with various line profiles shown (c, **i**, **ii** and **iii**) for conventional SRS (dash) and blind-S³ (line). (d) Large FOV image of opaque 100 µm thick mouse cerebellum (conventional SRS). A zoomed-in ROI (e) is depicted by dashed boxes with conventional SRS (left panel) and blind-S³ (right panel) methods, with a line profile chosen (f) for conventional (dash) and blind-S³ (line) methods. All scale bars: 500 nm. Pixel dwell times are top row (bottom row), 90 μ s and 180 μ s (100 μ s and 270 μ s) for conventional and blind-S³ methods, respectively, in panel **b**, and 100 μ s and 300 μ s for conventional and blind-S³ methods, respectively, in panel **e**.

the Pump and Stokes beams is small and as we use water (fast thermal dissipation). Finally, to

reduce the temperature rise due to the idle speckle illuminating the sample, one could alleviate

this by conjugating the excitation speckle plane with a scanning mirror (for instance, using a

253 digital micromirror device).



Fig. 5. Maximum temperature rise of the speckle evelope.

254 4.3. Discussion

Here, we have demonstrated above that the proposed method reaches resolution beyond the diffraction limit. We further demonstrated that the low excitation power used ensures low phototoxicity. Nevertheless, we discuss below potential issues in the methodology and how to overcome them with further engineering.

Further technical improvements could greatly overcome current limitations in these proof-of-259 principle experiments. In regards to speed and/or increasing the FOV, the current implementation 260 used classical frequency-domain spectral narrowing methods which have lower throughput 261 therefore limiting the largest FOV available (i.e. larger FOV requires more laser power to keep 262 the energy density constant). To address this issue, we envision an improvement using spectral 263 focusing methods, therefore increasing the FOV of the measurement as it can use all laser 264 power available. Despite these proposed follow-up improvements, the blind-S³ technique is 265 able to super-resolve in strongly opaque biological tissues: indeed, in biomedical applications, 266 aberrations deteriorate the PSF of the microscope, perhaps explaining the lower resolution 267 attained in the biological specimens observed here. Here, while the penetration depth of brain 268 tissues was limited to 8μ m, with recent advances of deep SRS imaging [40], this penetration 269 depth could in principle be extended. Finally, we speculate that blind- S^3 could be a route to 270 reach far-field nanoscopy (sub-100-nm resolution). A straightforward route for nanoscopy is 271 to decrease the excitation wavelengths in the UV [37]. However, UV radiation are known to 272 enhance phototoxicity (for instance, by generating DNA photoradicals [41]). With blind- S^3 , one 273 could still use excitation lasers in the visible range, and reach resolutions below the 100-nm 274 barrier because of the structured illumination approach. 275

276 5. Conclusions

In conclusion, we have designed and demonstrated a single-pixel super-resolution technique that is straightforward to implement, i.e. simpler than a conventional SRS microscope. We demonstrated that this technique can image beyond the diffraction limit ($\sqrt{2}$), and that it did $_{\tt 280}$ not show phototoxic effects in biological specimens imaged. blind-S 3 is an universal approach

in the sense that it depends neither on the specific vibrational mode ultrafast dynamics [29],

nor it requires vibrational signal enhancers [42,43], or *a priori* knowledge about the specimen,

for instance, as gained in the training of neural networks methods [44, 45]. Therefore, these achievements have overcome a decade-long challenge in SRS imaging, paving the way for

²⁸⁵ investigating matter in its most natural environments.

After the initial submission of the current work [46], an alternative computational approach has been experimentally demonstrated for SRS super-resolution of biological specimens [47] using deconvolution methods.

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