

Label-free metabolic imaging by mid-infrared optoacoustic microscopy in living cells

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We develop mid-infrared optoacoustic microscopy (MiROM) for label-free bond-selective live-cell metabolic imaging, enabling spatiotemporal monitoring of carbohydrates, lipids, and proteins in cells and tissues. Using acoustic detection of optical absorption, MiROM converts mid-infrared sensing into a positive-contrast imaging modality with negligible photodamage and high sensitivity. We use MiROM to observe changes in intrinsic carbohydrate distribution from a diffusive spatial pattern to tight co-localization with lipid droplets during adipogenesis.

Label-free dynamic imaging of biomolecules in living cells remains challenging for optical microscopy; at wavelengths below 700 nm, it is fundamentally hampered due to lack of chemical specificity or by cellular phototoxicity. Contrarily, chemically specific vibrational imaging modalities such as spontaneous or coherent Raman scattering and mid-infrared (**mid-IR**) microscopy have enormously extended possibilities for endogenous biomolecular imaging¹⁻³. Stimulated Raman scattering (**SRS**) microscopy, for instance, can rapidly image tissues at high spatial resolution^{4,5}. However, Raman imaging may photodamage cells⁶ and its sensitivity is above 1 mM¹, which misses biomolecules in the physiologically relevant micro- to

31 nanomolar range. Raman imaging of carbohydrates has been demonstrated mainly in fixed
32 cells^{7,8}, in living cells after using labelling (*i.e.* isotopic labeling) or glucose analogs^{5,9}, and in
33 densely-packed starch granules in microalgae¹⁰. Direct vibrational excitation by mid-IR
34 absorption complements Raman imaging³ (**Supplementary Table 1**). Based on photon
35 absorption, rather than scattering, mid-IR spectroscopy/imaging offers cross-sections up to
36 eight orders of magnitude larger than Raman imaging for certain molecules^{3,11,12}, sensitively
37 detecting the CH bond and the fingerprint spectral region³. Nevertheless, this modality is poorly
38 suited to metabolic studies of living cells because water strongly attenuates mid-IR light and
39 because it applies negative-contrast detection (*i.e.* the stronger the attenuation, the weaker
40 the signal detected). Typically, samples are placed within thin cuvettes (~10-25 μm thick) and
41 irradiated with high-powered mid-IR sources^{13,14}. Such confinement perturbs normal cellular
42 behavior and proliferation.

43 Here we introduce mid-IR optoacoustic microscopy (**MiROM**), a bond-selective imaging
44 modality based on biomolecule-specific vibrational transitions and radiation-less deexcitation
45 for highly efficient optoacoustic generation/detection (**Fig. 1**). We hypothesized that MiROM
46 could offer label-free detection for all types of biomolecules with high signal-to-noise ratio
47 (**SNR**) and contrast-to-noise ratio (**CNR**). MiROM relies on detection of ultrasound waves,
48 which are attenuated much less than mid-IR photons; ultrasonic detection also renders MiROM
49 as a positive-contrast method; *i.e.* the stronger the attenuation, the stronger the signal
50 detected. The unique combination of positive-contrast sensing and low-attenuation detection-
51 path balances signal loss, allowing deeper imaging than standard mid-IR microscopy. Here
52 MiROM was implemented in trans-illumination, which allowed diffraction-limited optical
53 excitation with confocal ultrasound detection (**Fig. 1a**). A mid-IR absorption-contrast
54 micrograph was obtained by raster-scanning the sample within the focal plane, simultaneously
55 acquiring optoacoustic signal from specific biomolecular vibrations. With the system presented
56 here, using an average laser power of only 330 μW , we obtained a limit of detection (**LOD**) of
57 2.5 mM for dimethyl sulfoxide (**DMSO**) (**Supplementary Fig. 1a,b**). For comparison, coherent
58 anti-Stokes Raman scattering (**CARS**) has an LOD of 70 mM for DMSO at an average laser

59 power of ~ 100 mW^{12,15}, while SRS detects up to 21 mM of DMSO at similar irradiation
60 power^{12,16}. Moreover, we obtained an LOD of 1.5 μ M for protein (albumin) in D₂O using only
61 500 μ W laser power (**Supplementary Fig. 1c-g**), which promises live-cell chemical
62 microscopy at significantly lower risk of phototoxicity⁶ (**Supplementary Fig. 2**).

63 First, we demonstrated the full biomolecular-contrast coverage of MiROM *in vitro* in
64 comparison with standard ATR-FTIR spectroscopy (**Fig. 1b,c; Supplementary Fig. 3;**
65 **Supplementary Tables 2-5**). Next, we demonstrated its bond-selective live-cell imaging
66 capabilities by mapping the lipid and protein content in HeLa cells, undifferentiated 3T3-
67 L1/PreBAT preadipocytes, and differentiated 3T3-L1 adipocytes. The symmetric CH₂ vibration
68 of lipids (around 2850 cm⁻¹) and the amide II band of proteins (around 1550 cm⁻¹), mainly NH
69 bending and CN stretching, were excited. Imaging wavenumbers were fine-tuned by live-cell
70 optoacoustic spectroscopy (**Fig. 1h,i**). The spatial (lateral) resolution of the system (~ 5.3 μ m
71 at 2850 cm⁻¹, **Supplementary Fig. 4a,b**) resolved single cells and individual lipid droplets
72 (**LDs**) in differentiated 3T3-L1 adipocytes (**Fig. 1d-h**). In HeLa cells and undifferentiated
73 preadipocytes, micrographs at the CH₂ vibration (lipid-map) show the phospholipid membrane
74 (**Supplementary Fig. 5**) at CNRs up to 22:1 (SNRs around 30:1, **Methods**). In differentiated
75 adipocytes, the dominant contrast comes from LDs because they contain more triglycerides
76 than the cell membrane and culture medium (**CM**). CNRs up to 220:1 and SNRs up to 223:1
77 were observed.

78 Positive contrast in amide II micrographs (protein-map) originated mainly from the overall
79 protein content, with a weak contribution from water. CNRs up to 41:1 and SNRs around 80:1
80 (**Supplementary Fig. 6a**). LDs showed negative contrast because they are hydrophobic and
81 contain less protein than the cytoplasm. The absolute contrast of LDs in the protein-map was
82 nevertheless high enough to be detected at SNRs up to 46:1, perhaps because of LD-
83 associated proteins (**Supplementary Fig. 6a**).

84 Besides lipids and proteins, carbohydrates were also detected in living 3T3-L1 adipocytes after
85 excitation between 1085 and 1000 cm⁻¹ (C–O stretching and C–O–H deformation). Localized

86 contrast was found in the cell body around the growing LDs with CNRs up to 24:1 and SNRs
87 around 40:1 (FOV: 50 μm x 50 μm ; imaging time: < 1 min/frame, **Fig. 2a-e**). One possible
88 explanation for the presence of carbohydrates in the cell body might be the capture and
89 accumulation of glucose for biosynthesis of triglycerides to be packed into LDs during
90 lipogenesis. This observation is supported by measurements in differentiating 3T3-L1 cells
91 where the molecular contrast (lipid, protein, and carbohydrate) was monitored in large FOVs
92 (1.5 mm x 1.5 mm) during several days (**Fig. 2f-n**). Glucose uptake was manipulated by
93 changing the concentrations of glucose and insulin in the medium (**Methods**). Carbohydrate
94 contrast increased in cells when incubated in 25 mM glucose and 1 $\mu\text{g}/\text{mL}$ insulin (**Fig. 2g-i,n**;
95 days 2, 4, and 6), and it decreased when incubated in 5 mM glucose only (**Fig. 2j,k,n**; days 8
96 and 10). These changes in MiROM carbohydrate contrast agreed with a standard colorimetric
97 assay of total carbohydrate (**Supplementary Fig. 7, Methods**). The carbohydrate contrast
98 was initially diffuse in the cell body and later co-localized with LDs during incubation
99 (**Supplementary Fig. 8**). This may reflect the appearance of glycoproteins or glycolipids in
100 LDs or their membranes.

101 We used MiROM to monitor lipid and protein dynamics during isoproterenol-induced lipolysis
102 in white adipocytes (differentiated 3T3-L1) and brown adipocytes (differentiated PreBAT)
103 during 4 hours. Lipid-maps (at 2857 cm^{-1}) and protein-maps (at 1550 cm^{-1}) were taken every
104 5 min before and after addition of isoproterenol to the medium (**Methods**). In both adipocyte
105 types, overall lipid content slowly increased before lipolysis, reflecting ongoing lipogenesis.
106 After induction, lipid contrast decreased continuously and nearly linearly (**Fig. 1m,n**;
107 **Supplementary Fig. 9a-d**). Different white adipocytes exhibited different lipolysis rates, with
108 some adipocytes unaffected by isoproterenol. Absorption of smaller LDs by larger ones (*i.e.*
109 LD remodeling) was continuously observed in some adipocytes (**Fig. 1m, ROI 1**). Other
110 adipocytes dimmed noticeably after lipolysis induction (**Fig. 1m, ROI 2**). The same
111 heterogeneous response to isoproterenol was observed in brown adipocytes (**Supplementary**
112 **Fig. 9a,c,d**). As expected, lipolysis was faster and more extensive in brown than in white
113 adipocytes: by 2 hours, lipid contrast had changed up to 30% in brown adipocytes, compared

114 to 18% in white adipocytes (**Videos 1a,b** and **2a,b**). Similar heterogeneous changes in protein
115 contrast were observed in cells during lipolysis (**Videos 3a,b**; **Supplementary Fig. 9e,f** and
116 **Supplementary Fig. 10**). This may reflect adipokine secretion, protein degradation and/or
117 protein translation. Changes in protein contrast may also reflect conformational changes
118 because amide II is relatively conformation-dependent¹⁷.

119 To demonstrate the potential of MiROM to image deeper in thicker samples than with standard
120 mid-IR imaging, we applied MiROM in a 4-mm-thick slice of freshly excised pancreatic (mouse)
121 tissue (**Fig. 1j-l**). Lipid micrographs show the pancreatic acinar glands in positive contrast
122 (CNR up to 58:1), while protein micrographs show, in negative contrast, the compartments
123 where acinar glands are embedded (CNR up to 21:1) (**Supplementary Fig. 6b**). Similar or
124 higher CNRs were observed in other similarly thick tissues (**Supplementary Fig. 11** and
125 **Supplementary Fig. 12**). A maximum imaging depth of 90 μm was obtained for the acinar
126 glands (**Supplementary Fig. 13**), compared to 575 μm in fat/polyamide-suture phantoms
127 (**Supplementary Fig. 14**). Regarding imaging speed, MiROM enables imaging FOVs of 5 mm
128 x 5 mm in steps of 10 μm (pixel size) in \sim 16 min.

129 MiROM offers unprecedented high contrast, image quality, sensitivity, and specificity for
130 endogenous biomolecular microscopy of living cells and thick unprocessed freshly-excised
131 tissues with negligible photodamage. For the first time, we visualize how carbohydrates initially
132 spread throughout young adipocytes, then co-localize with LDs upon adipocyte maturation.
133 MiROM can monitor intrinsic lipid/protein changes of $<1\%$ during lipolysis. MiROM is based on
134 vibrational excitation by mid-IR absorption and positive-contrast detection, so it offers great
135 sensitivity in the fingerprint spectral region, with LODs of 2.5 mM for DMSO and 1.5 μM for
136 albumin at laser powers in the 100's of μW . Most LDs visualized by confocal microscopy were
137 also resolved by MiROM, though the two types of images differed slightly, especially for
138 structures $< 5 \mu\text{m}$ (**Supplementary Fig. 15**). Resolution of MiROM may be improved by pump-
139 and-probe optoacoustic/optothermal signal read-out^{11,18,19} or ultra-wide bandwidth ultrasound
140 detectors²⁰.

141 Beyond carbohydrates, lipids, and proteins, MiROM can be used to analyze nucleic acids and
142 water in practically any other cell culture or tissue (shown *in vitro* in **Supplementary Fig. 3**).
143 In this way, MiROM supports live-cell metabolic research and analytical histology, while filling
144 an important gap in label-free biomolecular imaging and considerably extending the contrast
145 range of optoacoustic microscopy.

146

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154

155

156 **Author Contributions**

157 **M.A.P.** created the imaging concept, designed, built, and characterized the imaging system.
158 **M.A.P.**, **A.A.**, and **J.R.** designed and performed the experiments on adipocytes. **M.A.P.** and
159 **J.R.** designed and performed the experiments on excised tissues. **A.C.** synchronized and
160 automated the imaging system. **M.R.S.** performed the image processing and prepared the art
161 work. **M.A.P.** and **A.S.** designed and performed the molecular contrast validation experiments
162 and viability test on cells. **M.A.P.**, **F.G.**, and **B.S.** designed and performed the spectral
163 validation of the system. **M.A.P.** processed the results, prepared the images, and wrote the
164 manuscript. **M.A.P.**, **A.A.**, **J.R.**, and **M.S.** analyzed the results on lipolysis. **M.S.** and **S.H.**
165 supervised the study on lipolysis. **V.N.** supervised the whole study. All authors edited the
166 Manuscript.

167

168 **Ethics declarations**

169 *Competing Interests*

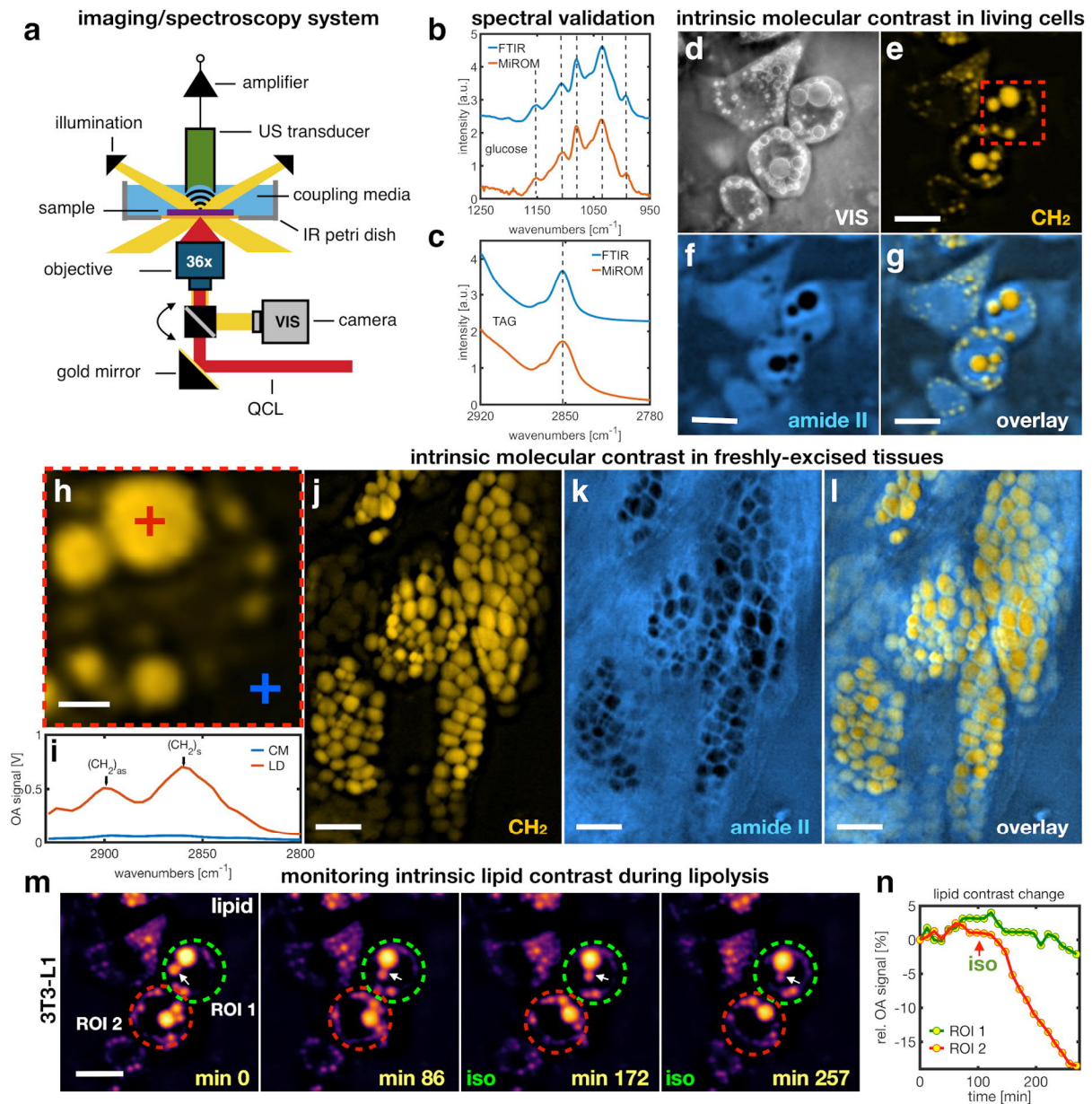
170 **V.N.** have financial interests in iThera Medical GmbH, equity owner and consultant; SurgVision
171 BV / Bracco Sp.A, member of the Scientific Advisory Board; Spear UG, owner. These
172 companies, however, did not contribute to this work. **V.N.** and **M.A.P.** are inventors on a
173 provisional patent application related to mid-infrared optoacoustic microscopy.

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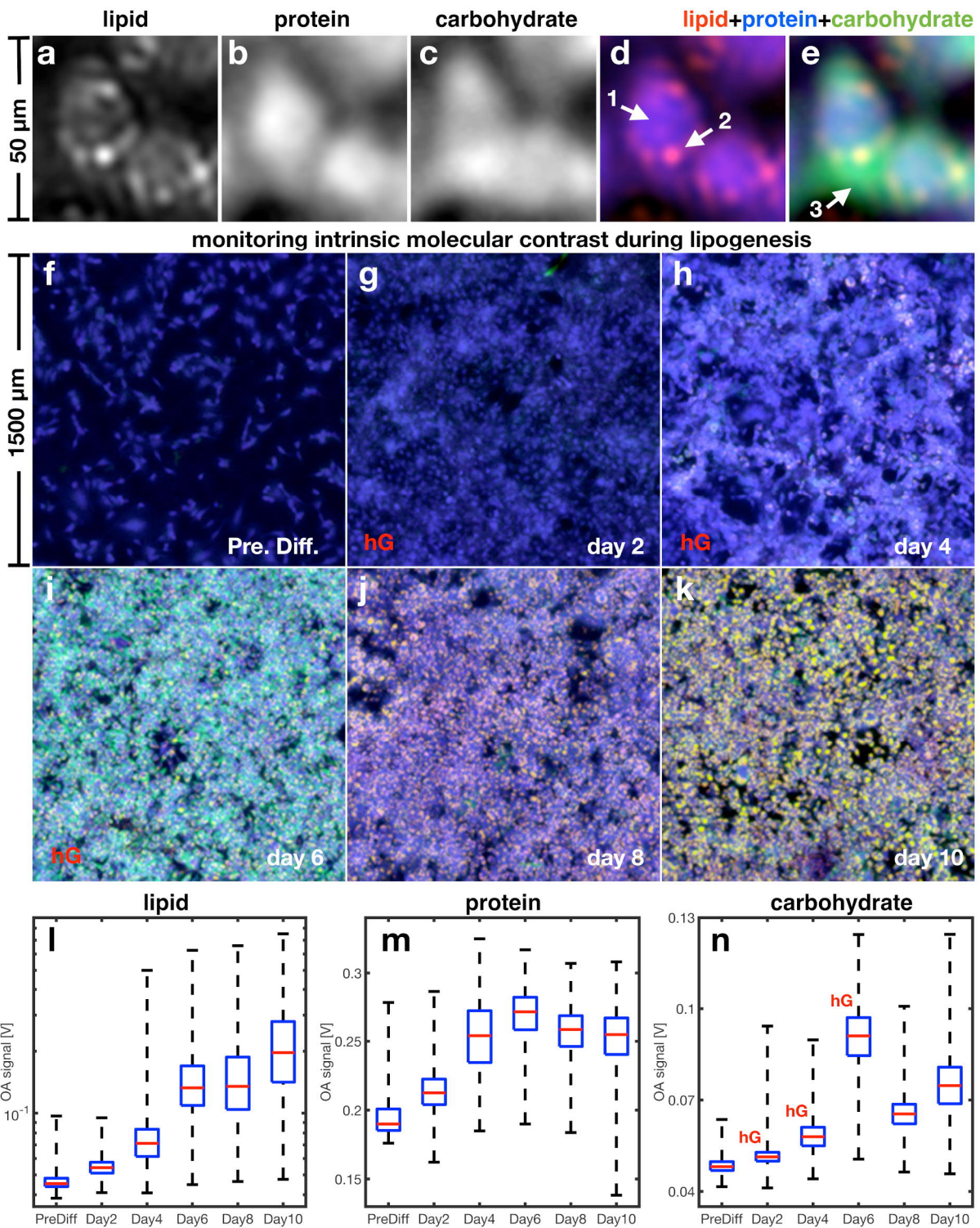
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221

222 **Figure 1: Mid-IR optoacoustic microscopy (MiROM).** a, Excitation-sample-sensor
 223 configuration for MiROM. Focused ultrasound (US) transducer and objective are confocally
 224 aligned to the sample plane. b,c, Comparison of MiROM and ATR-FTIR spectrum of glucose
 225 and 1,2-dioleoyl-3-palmitoyl-rac-glycerol (TAG) *in vitro*. The vertical dashed lines indicate good
 226 spectral matching between both methods. d, brightfield image of differentiated 3T3-L1 cells.
 227 e-g, MiROM micrographs of the cells in d with endogenous lipid contrast (e) at 2857 cm^{-1} (CH_2
 228 vibration) and protein contrast (f) at 1550 cm^{-1} (amide II). g, Overlay of lipid and protein maps.
 229 h, Zoom-in on a single adipocyte; dashed red square in e, scale bar: 10 μm . Two spots for
 230 spectral analysis and fine tuning of the imaging wavelength have been marked; on a lipid

231 droplet (**LD**) (red cross) and on the cell media (**CM**) (blue cross). **i**, Optoacoustic spectra in the
232 CH vibrational region for the selected spots in **h**. **j-l**, MiROM micrographs of freshly excised
233 pancreatic mouse tissue of 4 mm thickness, scale bar: 100 μm . **j**, Lipid map at 2850 cm^{-1} . **k**,
234 Protein map at 1550 cm^{-1} . **l**, Overlay of lipid and protein maps. Here, clusters of pancreatic
235 acinar glands embedded in protein are observed. **m,n**, Monitoring induced lipolysis in
236 differentiated 3T3-L1 adipocytes. **m**, Lipid monitoring sequence at 2857 cm^{-1} . Two region-of-
237 interest (**ROI**) enclosing individual adipocytes are marked, green dashed circle for ROI 1 and
238 red dashed circle for ROI 2. The white arrow follows the process of LD remodeling in a single
239 adipocyte enclosed in ROI 1. Time and presence of ISO is indicated at the bottom corners of
240 each frame (see also **Videos 2a,b**). **n**, Relative lipid contrast change for ROI 1 and 2 in **m**; the
241 red arrow indicates when ISO was added. Scale bar in **e-g**, **m**: 40 μm . Data in **b-n** are
242 representative of five independent experiments.



243

244 **Figure 2: Monitoring of carbohydrate contrast on 3T3-L1 adipocytes during LD**

245 **formation. a-e**, MiROM micrographs of 3T3-L1 cells at differentiation day 6; imaging speed: <

246 1 min per channel. **a**, Lipid map at 2853 cm^{-1} . **b**, Protein map, sum of amide I and II. **c**,

247 Carbohydrate map, sum of 1081 cm^{-1} and 1084 cm^{-1} . **d**, Overlay of lipid (red) and protein (blue)

248 maps. **e**, Same picture in **d** adding the carbohydrate (green) map. Arrow 1 indicates proteins

249 in the cell body, arrow 2 indicates an LD, and arrow 3 indicates an area of carbohydrate
250 accumulation around the growing LDs. **f-k**, Merged lipid (2857 cm^{-1}), protein (1541 cm^{-1}), and
251 carbohydrate (1022 cm^{-1}) maps of 3T3-L1 cells at different incubations days towards LD
252 formation; imaging speed: 16 min per channel. The overall carbohydrate contrast increases
253 after differentiation is started and it is broadly distributed in the cells at day 6 (CNR of 24:1 and
254 SNR of 40:1) around the areas of LD formation as in **e**. At day 8 and 10, however, carbohydrate
255 contrast is found only sparsely but highly co-localized with the LDs, see also **Supplementary**
256 **Fig. 8. i-n**, Box-plots ($n=90,000$) of OA contrast for each channel in micrographs **f-k**; box plots
257 indicate the upper and lower quartiles (box limits), median (center line), minimum and
258 maximum values (whiskers). During differentiation (day 2, 4, and 6), the cells show a rapid
259 increase of molecular contrast. Once the cell media is changed to normal incubation media
260 (low glucose concentration, no insulin) after day 6, lipid accumulation is observed at a lower
261 rate while proteins reach a rather stable level and the mean carbohydrate contrast drops,
262 correlating with extra cellular glucose and insulin modulation. The label '**hG**' (red font) indicates
263 when the cells were cultured with differentiation media (high glucose concentration + insulin).
264 Images **a-k** are representative of five independent experiments.

265 **Online Methods**

266 *System description*

267 A broadly tunable pulsed Quantum Cascade Laser (**QCL**) (MIRcat, Daylight Solutions, CA,
268 USA) is used for optoacoustic generation and biomolecular specificity; the spectral range of
269 the QCL is $3.4\text{--}11.0\ \mu\text{m}$ ($2941\text{--}909\text{ cm}^{-1}$) with a spectral linewidth $\leq 1\text{ cm}^{-1}$ (FWHM). The pulse
270 duration of the QCL is set to 20 ns at a repetition rate of 100 kHz then focused into the sample
271 by a 0.5 NA reflective objective (36x, Newport Corporation, CA, USA). The mid-IR absorption-
272 map of the sample is obtained by scanning the sample along the focal plane by motorized
273 stages (Prior Scientific/Physik Instrumente (PI), Cambridge/Karlsruhe, UK/DE) simultaneously
274 detecting the optoacoustic signal by a 20 or a 25 MHz central frequency focused ultrasound
275 transducer (Imasonic/Sonaxis, Voray sur l'Ognon/Besancon, France). The focused ultrasound
276 transducer and the reflective objective are coaxially aligned to share the same focal plane

277 where the sample is placed on a custom-made mid-IR transparent stainless-steel or acrylic-
278 glass petri-dish using a ZnSe window (Edmund Optics, Mainz, DE) or a ZnS window (Crystal
279 GmbH, Berlin, DE) as bottom substrate. For the live-cell studies, the cell media served as
280 acoustic coupling between the US transducer and the cells. Deionized water was used as
281 coupling media otherwise.

282 To remove interference from atmospheric CO₂ and water vapor, the mid-IR beam-path is
283 purged with a constant flow of dry N₂. A Mercury-Cadmium-Tellurium (**MCT**) detector (Daylight
284 Solutions, CA, USA) is used for optical reference and a VIS laser pointer, co-aligned with the
285 QCL beam, serves as aiming beam for easy optical adjustment. For validation, co-registration,
286 and easy ROI selection, oblique VIS illumination (Edmund Optics, Mainz, DE) is used to obtain
287 standard brightfield micrographs with a general purpose monochromatic camera (Edmund
288 Optics, Mainz, DE) (see **Fig. 1a** and **Supplementary Fig. 4a**).

289 The laser power at selected and relevant wavenumbers for this work was measured by a mid-
290 IR specific power meter (PE9-ES-C, Ophir-Spiricon, Darmstadt, DE) at 500 ns laser pulse
291 duration and 10 kHz repetition rate just before entering the reflective objective. The average
292 laser power at the sample was calculated scaling the measured values to the pulse duration
293 of 20 ns and repetition rate of 100 kHz taking also into account the obscuration (17%) of the
294 reflective objective. These laser excitation powers were used for all measurements reported
295 here; imaging, spectroscopy, and viability test. This is summarized as follows:

| Wavenumber (cm ⁻¹) | Measured average power at 500 ns and 10 kHz (mW) | Equivalent laser power at 20 ns, 100 kHz, and 17% obscuration (mW) |
|--------------------------------|--|--|
| 2850 | 0.8 | 0.27 |
| 1650 | 1.6 | 0.53 |
| 1550 | 2 | 0.66 |
| 1085 | 1.9 | 0.33 |

296

297 *Signal recovery, contrast, and noise.*

298 The raw optoacoustic signals were amplified by 63 dB (MITEQ, NY, USA) filtered with a 50
299 MHz low pass filter (Mini-Circuits, NY, USA) and then recorded at a sampling rate of 250 MS/s
300 on a 12 bit DAQ card, or at 200 MS/s on a 16 DAQ card, (Gage Applied, Lockport, USA). The
301 intensity of each pixel composing the micrographs shown in this work is the peak-to-peak
302 amplitude value resulting from the average of 50 or 100 optoacoustic transients (A-lines);
303 corresponding to a pixel dwell time of 1 ms or 500 μ s, respectively, at the pulse repetition rate
304 of 100 kHz.

305 The signal-to-noise ratio (*SNR*) is defined here as the ratio between the peak-to-peak amplitude
306 value of the optoacoustic signal (OAS_{pkpk}) over the peak-to-peak amplitude value of the noise
307 level ($Noise_{pkpk}$) before the arrival of the optoacoustic signal. For instance, the maximum SNR
308 of the lipid-map for the white adipocytes discussed above is 223:1, corresponding to a relative
309 error of 0.45 %. In terms of absolute values this corresponds to an optoacoustic peak-to-peak
310 amplitude of 702.2 mV and a peak-to-peak amplitude of the noise of 3.1 mV. In the protein-
311 map the maximum SNR is close to 80:1, or 1.3 % relative error. This is calculated from an
312 absolute optoacoustic peak-to-peak amplitude of 247 mV and a noise level of around 3 mV.

$$313 \quad SNR = OAS_{pkpk}/Noise_{pkpk}$$

314 The contrast-to-noise ratio (*CNR*) is defined here as the intensity difference between a point in
315 the sample (OA_S) and a point in the background (OA_B) divided over the peak-to-peak amplitude
316 value of the noise level.

$$317 \quad CNR = |OA_S - OA_B|/Noise_{pkpk}$$

318 The mean contrast-to-background ratio (mean CBR) described in **Supplementary Fig. 7** is the
319 mean contrast obtained from the image histogram divided over the background intensity (OA_B).
320 As explained below, prior to measurements on cells and tissues, the contrast, resolution,
321 spectral accuracy, sensitivity, and imaging depth of the system was tested in synthetic
322 samples: polyamide sutures and polyethylene microspheres embedded in agar, NIST spectral
323 gold standard, and dimethyl sulfoxide (**DMSO**). (**Supplementary Fig. 1, Supplementary Fig.**
324 **4b-d, Supplementary Fig. 14, and Supplementary Table 6**).

325

326 *Spectral validation*

327 First, the absorption spectrum of a NIST traceable Polystyrene calibration film for IR
328 spectroscopy (Sigma-Aldrich, St. Louis, USA) was measured in transmission mode by MiROM.
329 The position of the absorption peaks obtained by MiROM was compared with the certified
330 values of the NIST gold standard. We obtained a minimum deviation of 0.1 cm^{-1} around 1583
331 cm^{-1} and a maximum deviation of 2.6 cm^{-1} around 2850 cm^{-1} of the spectrum measured by
332 MiROM regarding the gold standard (see **Supplementary Table 6**).

333 Second, the ability to identify specifically and precisely different biomolecules was
334 demonstrated comparing the optoacoustic spectra obtained by MiROM with the spectra of the
335 same molecules (the same samples) obtained with a standard ATR-FTIR spectrometer (Bruker
336 Corporation, MA, USA).

337 Four samples representing each of the fundamental classes of biomolecules (namely: glucose,
338 DNA, triglycerides, and albumin) were prepared and measured with MiROM and ATR-FTIR as
339 follows:

340 A 50g/L water solution of glucose was prepared dissolving 2.5 g of D-(+)-glucose (Sigma-
341 Aldrich, St. Louis, USA) in 50 mL of distilled water.

342 A 5% water solution of DNA (Deoxyribonucleic acid sodium salt from *Escherichia coli* strain B,
343 Sigma-Aldrich, St. Louis, USA) was prepared dissolving 1 mg of DNA in 20 μL of distilled water.

344 A 5- μL drop of this solution was placed in a carbon tape ring located on a dish and dried under
345 vacuum. The sample was covered with a plastic film to avoid water dilution.

346 A 10 mg/mL solution of 1,2-dioleoyl-3-palmitoyl-rac-glycerol (TAG) (Sigma-Aldrich, St. Louis,
347 USA) was prepared dissolving 1 mg in 100 μL of a chloroform/methanol solution (2:1). A 10-
348 μL drop of this solution was placed on the window of a dish and dried under vacuum.

349 An 80g/L D_2O solution of albumin was prepared dissolving 2.0 g of albumin (Carl Roth) in 25
350 mL of heavy water. As commonly done in standard FTIR spectroscopy, for albumin, D_2O was
351 used instead of H_2O in order to avoid the strong absorption peak of water around 1650 cm^{-1}
352 (**Supplementary Fig. 3d**), overlapping the amide I band of proteins.

353 All the measurements were performed on the custom-made mid-IR petri-dish with a ZnSe
354 window using carbon tape (SPI Suppliers, PA, USA) as spectral reference. The optoacoustic
355 spectra were measured with a resolution of 2 cm^{-1} and an averaging time of 100 ms per
356 wavenumber. For comparison, a 5- μL drop of each solution was measured with the ATR-FTIR
357 spectrometer equipped with a diamond ATR crystal. The same resolution of 2 cm^{-1} , as with the
358 MiROM system, was used to record the spectra with the ATR-FTIR spectrometer. For water
359 solutions, water was measured and subtracted from both, the optoacoustic and FTIR spectra.
360 As observed in **Fig. 1b,c**; **Supplementary Fig. 3** and **Supplementary Tables 2-5**, MiROM
361 was able to accurately detect the absorption bands of biomolecules.

362 *Determination of the limit of detection (LOD)*

363 We determined the LOD of MiROM for two reference molecules: DMSO in H_2O and albumin
364 in D_2O . For DMSO, the optoacoustic spectrum at different concentrations (from 664 mM to
365 0.02 mM) was measured between 1250 and 909 cm^{-1} , averaging time ~ 100 ms per
366 wavenumber (**Supplementary Fig. 1a**). The area under the absorption band of the S=O
367 vibration at 1010 cm^{-1} was selected for determination of the lowest detectable concentration of
368 DMSO, the optoacoustic signal at 1066 cm^{-1} and 984 cm^{-1} were used for spectral baseline
369 correction and normalization, respectively. The area under the normalized optoacoustic
370 intensity of the S=O vibration (**Supplementary Fig. 1b**) as well as its optoacoustic spectra,
371 show that DMSO can be detected, above the spectral baseline, at a concentration of 2.5 mM
372 with a SNR of 6:1. Here, the baseline, or noise level, is defined by the difference of two
373 measured optoacoustic spectra of water. The maximum average laser power in the spectral
374 range measured is $330\text{ }\mu\text{W}$ at around 1085 cm^{-1} , measured as described above.

375 For albumin, the optoacoustic spectrum at different concentrations (from $750\text{ }\mu\text{M}$ to $1.5\text{ }\mu\text{M}$)
376 was measured between 1700 and 1600 cm^{-1} (the amide I band), averaging time ~ 100 ms per
377 wavenumber (**Supplementary Fig. 1d**). The area under the amide I band was selected for
378 determination of the lowest detectable concentration of albumin, the optoacoustic signal at
379 1700 cm^{-1} and 1600 cm^{-1} were used for spectral baseline correction and normalization,
380 respectively. The area under the normalized optoacoustic intensity of the amide I band

381 (**Supplementary Fig. 1e**) as well as its optoacoustic spectra show that albumin can be
382 detected, above the spectral baseline, at a concentration of 1.5 μM with an SNR of 4.5:1. Here,
383 the baseline, or noise level, is defined by the difference of two measured optoacoustic spectra
384 of heavy water. The maximum average laser power in the spectral range measured is 530 μW
385 at around 1650 cm^{-1} , measured as described above.

386 The SNR for the LOD determination (SNR_{LOD}) is defined here as the area of the normalized
387 optoacoustic absorption band of the analyte ($Area_{NOAS}$; the signal), after buffer subtraction,
388 divided by the area of the baseline ($Area_{BL}$; the noise) (absolute value).

$$389 \quad SNR_{LOD} = Area_{NOAS} / Area_{BL}$$

390 *Image processing*

391 In order to enhance visibility and compensate for spatial resolution of mid-IR microscopy in the
392 range of subcellular compartments of interest ($\sim 5 \mu\text{m}$), the images were bicubic interpolated
393 to a pixel size of 250 nm and deconvolved with the experimental determined point-spread
394 function (see **Supplementary Fig. 4b**) by a 3- or 5-step iterative Wiener deconvolution.
395 Furthermore, the images were post processed by a 2-pixel Gaussian filtering, outlier removal
396 if necessary, a contrast enhancement to a 0.3% saturation, and a histogram normalization. For
397 resolution analysis and SNR determination, the images were kept unprocessed.

398

399 *Preparation and measurement of white and brown adipocytes*

400 3T3-L1 mouse white preadipocyte cells were plated in the custom plates and cultured till
401 confluency in growth media consisting of: Dulbecco's Modified Eagle's Medium (DMEM) low
402 glucose (1 g/L) (Life Technologies, Paisley, GBR) supplemented with 10% fetal bovine serum
403 (FBS, Merck, Darmstadt, DE) and 1% pen/strep (Life Technologies, Bleiswijk, NLD). Cells
404 were kept in incubator at 37°C, 5% of CO_2 . The process of differentiation lasted for 6 days. On
405 day 0 and day 2, differentiation media consisting in DMEM high glucose (4.5 g/L), 10% FBS,
406 1% pen/strep, supplemented with 1 $\mu\text{g}/\text{mL}$ insulin (Sigma, Steinheim, DE), 0.25 μM
407 dexamethaxone (Sigma, Steinheim, DE), 0.5 mM 3-isobutyl-1-methylxanthine (Sigma,
408 Steinheim, DE) and 1/1000 volume ABP (50 mg/mL L-ascorbate, 1 mM biotin, 17 mM

409 pantothenate; Sigma, Steinheim, DE) was added to the cells. On day 4, differentiation media
410 supplemented only with insulin and ABP was used. From day 6 and on (after MiROM
411 measurements) the cells were kept in growth media (low glucose, no insulin).

412 The PreBAT cell line was created and provided by Hoppmann, Perwitz et al. (2010) by
413 immortalizing preadipocytes from the intrascapular BAT of newborn mice using the SV40
414 Large T antigen. The differentiation process also lasted for 6 days starting with induction on
415 day 0 with DMEM growth media (4.5 g/L glucose; Life Technologies, Paisley, Scotland)
416 containing 20 % FBS (Merck, Darmstadt, DE) and 1 % Pen/Strep (Life Technologies, Bleiswijk,
417 Netherlands) with the addition of 20 nM insulin (Sigma, Steinheim, DE), 1 μ M triiodothyronine
418 (T_3) (Sigma, Steinheim, DE), 0.125 mM indomethacin (Sigma, Steinheim, DE), 2 μ g/ml
419 dexamethasone (Sigma, Steinheim, DE) and 0.5 mM 3-isobutyl-1-methylxanthine (Sigma,
420 Steinheim, DE). The cells were kept in the incubator at 37°C, 5% of CO₂. On day 2 and day 4,
421 growth media containing only insulin and T_3 was added and on day 6 the cells were given only
422 growth media. At the end of differentiation, both cell lines showed abundant amounts of lipid
423 droplets.

424 All MiROM measurements on cells were performed using growth media; changing from
425 differentiation media to growth media right before imaging whenever needed.

426

427 *Quantification of total carbohydrates content*

428 The total amount of carbohydrates in the differentiated 3T3-L1 cell lysates were measured
429 using the Cell Biolabs' Total Carbohydrate Assay Kit (STA-682, Cell Biolabs, San Diego, CA,
430 USA) following the manufacturer's protocol. After each MiROM measurement (at day 0, 2, 4,
431 6, 8 and 10), the cells were removed from the customized plates upon trypsin digestion and
432 centrifuged at 2000 rpm, for 3 min. The supernatant was discarded and the pellet was washed
433 once with PBS. After centrifugation, the cells were collected and stored at -80 °C. The cells
434 were resuspended at 1-2 x 10⁶ cell/mL in 1X Assay Buffer. Each pellet was vigorously
435 homogenized and centrifuged to remove the debris. Cell lysates were assayed undiluted in a
436 96-well plate, 30 μ L of each solution was mixed with 150 μ L of concentrated sulfuric acid and

437 incubated for 15 min at 90°C. After addition of a developing solution, containing 5% of phenol
438 in 1X diluent buffer, a chromogen is formed and it can be detected at 490 nm. The samples
439 were analyzed with the microplate reader CLARIOstar (BMG LABTECH) at optical density
440 (OD) 490 nm. For each point, the background OD (acquired before the addition of the
441 developing solution) was subtracted. Experiments were carried out in duplicate.

442

443 *Cell viability study.*

444 To assess the negligibility of photodamage induced by MiROM, a standard viability test on
445 undifferentiated 3T3-L1 preadipocytes was performed. A 10 µL-drop of cells was plated in a
446 custom-made MiROM dish, with a ZnSe window of 12.7 mm diameter. After overnight
447 incubation in growth media, at 37°C, 5% of CO₂, a FOV of 2 mm x 2 mm was irradiated with a
448 pulse laser of 100 kHz during a total irradiation time of 4 h 40 min: 20 scanning loops (frames)
449 for each excitation wavenumber (1557 and 2850 cm⁻¹), corresponding to 7 min per frame.
450 During the measurement, the cells were kept at 37 °C. Cell viability was assessed using
451 erythrosine B exclusion assay. Cell survival was expressed as the percentage ratio of viable
452 irradiated cells in comparison with the corresponding viable not-irradiated controls. For
453 statistical elaboration OriginPro9.1 Software was used. Reported data corresponded to the
454 mean values ± standard deviation (SD) obtained from three different experiments.

455

456 *Preparation and measurement of mouse tissues*

457 Male C57BL/6J mice (8-10 weeks old; Charles River Laboratories Inc, Charleston, USA) were
458 kept at 24±1°C and fed with standard rodent diet (Altromin 1314, Altromin Spezialfutter GmbH
459 & Co, Germany) with free access to water, with constant humidity and on a 12-h light–dark
460 cycle. After the mice were sacrificed the organs were harvested and directly placed on the
461 sample holder where they were covered with low temperature melting agar (2%) and de-
462 ionized water as coupling media.

463

464 **Data Availability Statement**

465 The data that support the findings of this study are available from the corresponding authors
466 upon reasonable request.