

Imaging isotopically labeled bacteria at the single cell level using high-resolution optical infrared photothermal spectroscopy

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Figure S1 (A) PCA scores plot of preprocessed O-PTIR spectra acquired from a substrate containing a mixture of *E. coli* cells grown on ¹²C/¹³C-glucose and/or ¹⁴N/¹⁵N-ammonium chloride; each point is a spectrum from a single bacterium; values on the axes in parentheses are the percent total explained variance (TEV) by each PC. (B) PCA loadings plot of the O-PTIR spectral data with PC-1 (dashed line) and PC-2 (full line) showing the major peaks affected by the labelling process. (C) Comparison of loadings related to PC-1 and PC-2 obtained for spectra recorded in two configurations. Dashed lines represent spectral data acquired from each condition spotted into four different CaF₂ substrates; full lines relate to data collected from labelled and unlabelled cells on the same substrate. The plots are offset for clarity.

Figure S2 Single-point O-PTIR spectra acquired from single bacterial cells numbered 1-6 on Figure 4A, in which from inspection of the amide I and II positions we identified one unlabeled bacterium (cell 2, blue line); one ¹³C-labeled bacterium (cell 5, red line), three cells grown on ¹⁵N-ammonium chloride (cells 3, 4, 6, yellow lines), and one cell cultivated on both compounds (cell 1, green line). The plots are offset for clarity.

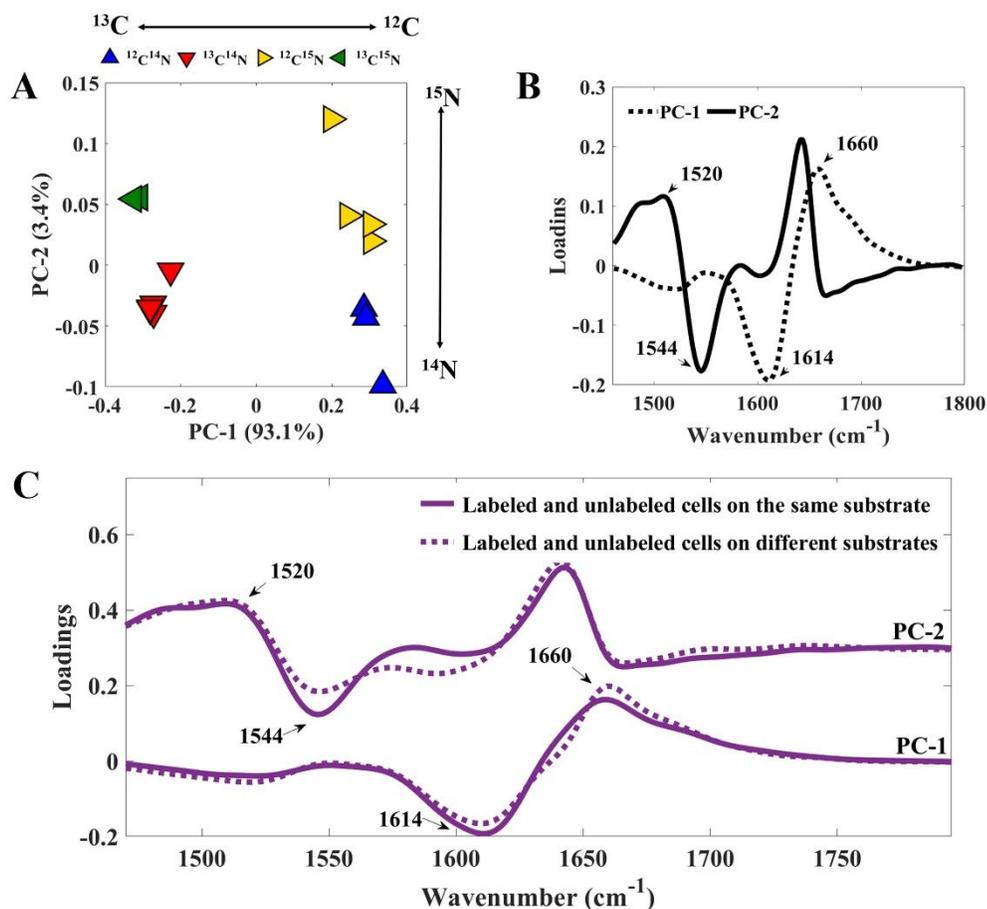


Figure S1 (A) PCA scores plot of preprocessed O-PTIR spectra acquired from a substrate containing a mixture of *E. coli* cells grown on $^{12}\text{C}/^{13}\text{C}$ -glucose and/or $^{14}\text{N}/^{15}\text{N}$ -ammonium chloride; each point is a spectrum from a single bacterium; values on the axes in parentheses are the percent total explained variance (TEV) by each PC. (B) PCA loadings plot of the O-PTIR spectral data with PC-1 (dashed line) and PC-2 (full line) showing the major peaks affected by the labelling process. (C) Comparison of loadings related to PC-1 and PC-2 obtained for spectra recorded in two configurations. Dashed lines represent spectral data acquired from each condition spotted into four different CaF_2 substrates; full lines relate to data collected from labelled and unlabelled cells on the same substrate. The plots are offset for clarity.

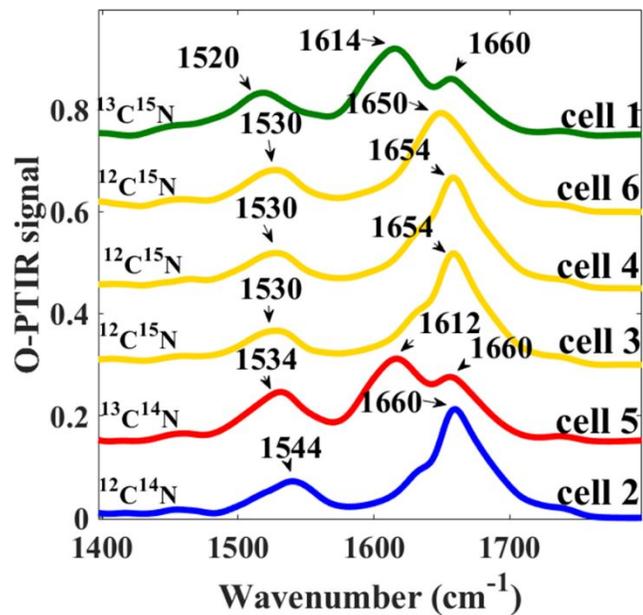


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