## **Support Information**

## Metabolic Labeling of Peptidoglycan Enabled Optical Analysis of Probiotic Vitality

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**Figure S1.** (A) Representative flow cytometry results of metabolic labeling on *L. paracasei* with the addition of different structures of amino acids. (B) Representative confocal images of *L. paracasei* via metabolic labeling with different conditions. Red, DBCO-Sulfo-Cy5. Scale bar, 5 µm.



**Figure S2.** (A) Representative standard plate count results of *L. paracasei* after NaN<sub>3</sub> treatment or not. (B) Representative confocal images of *L. paracasei* with live and dead staining after NaN<sub>3</sub> treatment or not. Green, Syto-9; Red, Propidium iodide. Scale bar, 2

 $\mu$ m. (C) Representative confocal images of *L. paracasei* with CFDA staining after NaN<sub>3</sub> treatment or not. Green, CF (carboxyfluorescein). Scale bar, 5  $\mu$ m. (D) Representative flow cytometry results showing bacterial live and dead staining of *L.paracasei* after NaN<sub>3</sub> treatment or not. (E) Representative flow cytometry results showing CFDA staining of *L.paracasei* with NaN<sub>3</sub> treatment or not. Negative refer to control without the addition of CFDA.



**Figure S3.** (A) Representative standard plate count results of *L. paracasei* after treatment at different temperatures. (B) Representative confocal images of *L. paracasei* with live and dead staining after treatment at different temperatures. Green, Syto-9; Red, Propidium iodide. Scale bar, 5 µm. (C) Representative flow cytometry results showing bacterial live and dead staining of *L.paracasei* after treatment at different temperatures.



**Figure S4.** (A) Fluorescence intensity of bacterial solutions via metabolic labeling after treatment of different temperatures (n=3). <sup>ns</sup>p>0.05, \*p<0.05, \*\*p<0.01, \*\*\*p<0.001, \*\*\*\*p<0.001. (B) Representative confocal images of *L. paracasei* via metabolic labeling after treatment at 25 °C and 90 °C. Red, DBCO-Sulfo-Cy5.Scale bar, 5  $\mu$ m. (C) Fluorescence intensity of bacterial solutions via metabolic labeling after the operation of lyophilization or not (n=3). <sup>ns</sup>p>0.05, \*p<0.05, \*\*p<0.01, \*\*\*\*p<0.001, \*\*\*\*p<0.0001.



**Figure S5.** (A) Fluorescence intensity of bacterial solutions through metabolic labeling after treatment of different pH values (n=3). <sup>ns</sup>p>0.05, \*p<0.05, \*\*p<0.01, \*\*\*p<0.001,

\*\*\*\*p<0.0001. (B) Representative confocal images of *L. paracasei* via metabolic labeling after treatment at different pH values and culture medium. Red, DBCO-Sulfo-Cy5.Scale bar, 5  $\mu$ m. (C) Fluorescence intensity of bacterial solutions via metabolic labeling after the treatment of SGF or SIF (n=3). <sup>ns</sup>p>0.05, \*p<0.05, \*\*p<0.01, \*\*\*p<0.001, \*\*\*\*p<0.0001.



**Figure S6.** (A) Representative standard plate count results of *L. paracasei* after treatment at different pH values. (B) Representative confocal images of *L. paracasei* with live and dead staining after the treatment of pH=1 and 2. Green, Syto-9; Red, Propidium iodide. Scale bar, 5 µm. (C) Representative flow cytometry results showing bacterial live and dead staining of *L.paracasei* after the treatment of pH=1 and 2.



**Figure S7.** (A) Representative standard plate count results of *L. paracasei* after treatment in normal saline for 2h (NS-2h), SGF for 2h, normal saline for 4h (NS-4h), and SIF for 4 h. (B) Representative confocal images of *L. paracasei* with live and dead staining after being cultured in normal saline for 2h (NS-2h), SGF for 2h, normal saline for 4h (NS-4h), and SIF for 4 h. Green, Syto-9; Red, Propidium iodide. Scale bar, 5  $\mu$ m. (C) Representative flow cytometry results showing bacterial live and dead staining of *L.paracasei* after the treatment in normal saline for 2h (NS-2h), SGF for 2h, normal saline for 4h (NS-4h), and SIF for 4 h.



**Figure S8.** (A) Representative confocal images of *L. rhamnosus* GG, *L. casei* and *B. subtilis* via metabolic labeling under normal culturing conditions. Red, DBCO-Sulfo-Cy5.

Scale bar, 5  $\mu$ m. (B) Fluorescence intensity of the bacterial solutions via metabolic labeling with different storage time (n=3). <sup>ns</sup>p>0.05, \*p<0.05, \*\*p<0.01, \*\*\*\*p<0.001, \*\*\*\*p<0.0001.