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Evaluation of sample preservation and storage methods for metaproteomics analysis of intestinal microbiomes

Angie Mordant and Manuel Kleiner

Corresponding Author(s): Angie Mordant, North Carolina State University

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North Carolina State University
Plant & Microbial Biology
4510 Thomas Hall
112 Derieux Place
Raleigh, NC 27695

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1752 N St., NW
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E-mail: spectrum@asmusa.org

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