Journal Pre-proof

Evaluation of N-terminal labeling mass spectrometry for characterization of partially hydrolyzed gluten proteins

Wanying Cao, Joseph L. Baumert, Melanie L. Downs

PII: S1874-3919(19)30310-0

DOI: https://doi.org/10.1016/j.jprot.2019.103538

Reference: JPROT 103538

To appear in: Journal of Proteomics

Received date: 9 May 2019

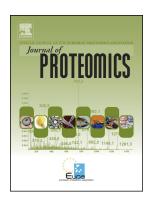
Revised date: 17 September 2019

Accepted date: 27 September 2019

Please cite this article as: W. Cao, J.L. Baumert and M.L. Downs, Evaluation of N-terminal labeling mass spectrometry for characterization of partially hydrolyzed gluten proteins, *Journal of Proteomics* (2018), https://doi.org/10.1016/j.jprot.2019.103538

This is a PDF file of an article that has undergone enhancements after acceptance, such as the addition of a cover page and metadata, and formatting for readability, but it is not yet the definitive version of record. This version will undergo additional copyediting, typesetting and review before it is published in its final form, but we are providing this version to give early visibility of the article. Please note that, during the production process, errors may be discovered which could affect the content, and all legal disclaimers that apply to the journal pertain.

© 2018 Published by Elsevier.



Journal Pre-proof

Evaluation of N-terminal Labeling Mass Spectrometry for Characterization of Partially Hydrolyzed Gluten Proteins

Wanying Cao, Joseph L. Baumert, Melanie L. Downs*

Food Allergy Research and Resource Program, Department of Food Science and Technology, Food Innovation Center, 1901 North 21st Street, University of Nebraska-Lincoln, Lincoln, NE 68588

Corresponding Author

*Dr. Melanie L. Downs, Food Allergy Research and Resource Program, Department of Food Science and Technology, Food Innovation Center, 1901 North 21st Street, University of Nebraska-Lincoln, Lincoln, NE 68588-6205. Phone: 402-472-5423. Email address: mdowns2@unl.eduABSTRACT

Gluten, a group of proteins found in wheat, barley, and rye, is the trigger of celiac disease, an immune disorder that affects about 1% of people worldwide. The toxicity of partially hydrolyzed gluten (PHG) in fermented products is less well understood due to the significant analytical challenges in PHG characterization. In this project, an N-terminal labeling mass spectrometry method, terminal amine isotopic labeling of substrates (TAILS), was optimized for the in-depth analysis of PHG and validated using a test protease (trypsin) with known cleavage specificity. Gluten N-termini in test and control groups were labeled with heavy and light formaldehyde, respectively. Trypsin-generated neo N-termini were identified by exhibiting an MS1 Log2 H:L

Download English Version:

https://daneshyari.com/en/article/13472915

Download Persian Version:

https://daneshyari.com/article/13472915

<u>Daneshyari.com</u>