

## Accepted Manuscript

Manipulating trypsin digestion conditions to accelerate proteolysis and simplify digestion workflows in development of protein mass spectrometric assays for the clinical laboratory

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PII: S2376-9998(17)30001-6  
DOI: <https://doi.org/10.1016/j.clinms.2017.10.001>  
Reference: CLINMS 25

To appear in: *Clinical Mass Spectrometry*

Received Date: 8 January 2017  
Revised Date: 9 October 2017  
Accepted Date: 17 October 2017

Please cite this article as: Y.Z. Zheng, M.L. DeMarco, Manipulating trypsin digestion conditions to accelerate proteolysis and simplify digestion workflows in development of protein mass spectrometric assays for the clinical laboratory, *Clinical Mass Spectrometry* (2017), doi: <https://doi.org/10.1016/j.clinms.2017.10.001>

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August 25, 2017

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**Abbreviations:**

MS: mass spectrometry; LC: liquid chromatography; MRM: multiple reaction monitoring; DP: declustering potential; CE: collision energy; apoA1: apolipoprotein A-1; RBP4: retinol-binding protein 4; TTR: transthyretin/prealbumin; CO9: complement component 9; CRP: C-reactive proteins; HEPES: 4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid; AA: ammonium acetate; IAA: iodoacetamide; DTT: dithiothreitol; AMBIC: ammonium bicarbonate; SDC: sodium deoxycholate; TPCK: tosyl phenylalanyl chloromethyl ketone; ACN: acetonitrile; Q1 ion: precursor ion; Q3 ion: product ion; PTMs: post-translational modifications

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