

## **MEETING ABSTRACTS**

## MASS SPECTRAL DETECTION OF DIETHOXYPHOSPHO-TYROSINE ADDUCTS ON PROTEINS FROM HEK293 CELLS USING MONOCLONALANTIBODY DEPY FOR ENRICHMENT

Seda Onder 1,2 \*, Lawrence M. Schopfer 2, Ozden Tacal 1, Thomas A. Blake 3, Rudolph C. Johnson 3, Oksana Lockridge 2\*

- \* Presenting author, Seda Onder
- <sup>1</sup> Department of Biochemistry, School of Pharmacy, Hacettepe University, Ankara, 06100, Turkey sedaonder@hacettepe.edu.tr, tacal@hacettepe.edu.tr
- <sup>2</sup> Eppley Institute, University of Nebraska Medical Center, Omaha, NE, 68198, USA lmschopf@unmc.edu, olockrid@unmc.edu
- <sup>3</sup> Division of Laboratory Sciences, National Center for Environmental Health, Centers for Disease Control and Prevention, 4770 Buford Highway NE, Atlanta, GA, 30341, USA rmj6@cdc.gov, fsi3@cdc.gov

Chronic illness from exposure to organophosphorus toxicants is hypothesized to involve modification of unknown proteins. Tyrosine readily reacts with organophosphorus toxicants in proteins that have no active site serine. We developed a monoclonal antibody, depY, that specifically recognizes diethoxyphospho-tyrosine in proteins and peptides, independent of the surrounding amino acid sequence <sup>1</sup>. Our goal was to identify diethoxyphosphorylated proteins in human HEK293 cell lysate treated with chlorpyrifos oxon. Cell lysates treated with chlorpyrifos oxon were examined by ELISA and capillary electrophoresis Western blot. Tryptic peptides were analyzed by liquid chromatography-tandem mass spectrometry. The depY antibody recognized diethoxyphosphotyrosine containing proteins by ELISA and Western blotting. Mass spectrometry identified 40 diethoxyphosphotyrosine peptides from 24 proteins in immunopurified samples, but found only 9 diethoxyphosphotyrosine peptides from 6 proteins when the same sample was not immunopurified on depY. The most abundant proteins in the cell lysate, Histone H4, Heat shock 70 kDa protein 1A/1B, Heat shock protein HSP 90 beta, and Alpha-enolase, were represented by several diethoxyphosphotyrosine peptides. It was concluded that use of immobilized depY improved the number of diethoxyphosphotyrosine peptides identified in a complex mixture. The mass spectrometry results confirmed the specificity of depY for diethoxyphosphotyrosine peptides independent of the context of the modified tyrosine, which means depY could be used to analyze modified proteins in any species.

Keywords: chlorpyrifos oxon; diethoxyphospho-tyrosine antibody; mass spectrometry; ELISA; Western blot

## References

1. Onder, S., Dafferner, A. J., Schopfer, L. M., Xiao, G., Yerramalla, U., Tacal, O., Blake, T. A., Johnson, R. C., and Lockridge, O. (2017) Monoclonal Antibody That Recognizes Diethoxyphosphotyrosine-Modified Proteins and Peptides Independent of Surrounding Amino Acids. *Chem Res Toxicol* 30, 2218-2228