

# DERIVATIZATION REAGENTS FOR NEGATIVE MODE LC-MS

Ernesto Zapata<sup>1</sup>, Koit Herodes<sup>1</sup>, Ivo Leito<sup>1</sup>

<sup>1</sup>*Institute of Chemistry, University of Tartu, Ravila 14a, 50411 Tartu, Estonia*  
e-mail: edjzapata@outlook.com

Liquid chromatography coupled to mass spectrometry (LC-MS) is a very flexible analytical technique that allows to analyse different analytes in a big variety of matrices [1-3]. One of its biggest disadvantages is the susceptibility to matrix effects. There are different approaches that can be used to overcome this problem. Derivatization of analytes is a good alternative that improves not only chromatographic retention and selectivity as well as LoD/LoQ, which altogether pave the road to reduce matrix effects.

Previous studies have shown that LC-MS/MS analysis using negative ionization mode, produces cleaner chromatograms [4-5]. In this study, a derivatization method was developed using novel derivatization reagents based on azo-benzene. Amino acids derivatives were analysed in positive and negative ionization modes and MS/MS fragmentation patterns were studied.

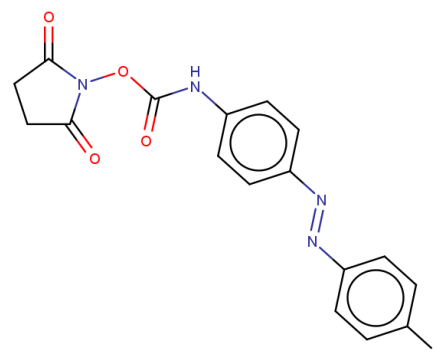


Figure 1. Generic structure of azo-benzene derivatization reagents.

## References

1. Zhu, K., Li, P., Feng, M. et al. *Environ Monit Assess* (2015) 187: 539. <https://doi.org/10.1007/s10661-015-4773-2>
2. Xin, X., Fan, R., Gong, Y. et al. *Eur Food Res Technol* (2014) 238: 837. <https://doi.org/10.1007/s00217-014-2153-8>
3. Gremmel, C., Frömel, T. & Knepper, T.P. *Anal Bioanal Chem* (2017) 409: 1643. <https://doi.org/10.1007/s00216-016-0110-z>
4. Ziegler, J. & Abel, S., *Amino Acids* (2014) 46:2799. <https://doi.org/10.1007/s00726-014-1837-5>
5. Oldekop, M.-L., Rebane, R., & Herodes, K. (2017). *European Journal of Mass Spectrometry*, 23(5), 245–253. <https://doi.org/10.1177/1469066717711026>