protocols.io

🌐 Native Protein Analysis on TIMSTOF Pro mass spectrometer V.2

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dx.doi.org/10.17504/protocols.io.bzv2p68e

LCMSMethods.org Version 2

lcmsmethods

Method described in David Robert et al.,
https://pubs.acs.org/doi/10.1021/jacs.1c02713

DOI

dx.doi.org/10.17504/protocols.io.bzv2p68e

> David Roberts 2021. Native Protein Analysis on TIMSTOF Pro mass
> spectrometer. **protocols.io**
> https://dx.doi.org/10.17504/protocols.io.bzv2p68e
> lcmsmethods

The second version has a zipped .m file for the TIMSTOF Pro method

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1   Method for native protein analysis on a TIMSTOF Pro mass
   📎 **Native_200-5000mz_LowMass1800mz_tuned0pt95to1pt60.m.zip**
   spectrometOriginal study by David Robert et al.,