

Intelligent workflows for automated analysis of mass spectrometry-based proteomics data Güler. A.T.

Citation

Güler, A. T. (2022, April 7). *Intelligent workflows for automated analysis of mass spectrometry-based proteomics data*. Retrieved from https://hdl.handle.net/1887/3281870

Version: Publisher's Version

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Note: To cite this publication please use the final published version (if applicable).

Intelligent Workflows for Automated Analysis of Mass Spectrometry-based Proteomics Data

Arzu Tuğçe Güler

ISBN: 978-94-6423-731-3

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Cover design: Arzu Tuğçe Güler Printing: ProefschriftMaken.nl

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Proefschrift

ter verkrijging van de graad van doctor aan de Universiteit Leiden, op gezag van rector magnificus prof.dr.ir. H. Bijl, volgens besluit van het college voor promoties te verdedigen op donderdag 7 april 2022 klokke 11.15 uur

door

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