IMPERIAL

Protein Digestion, Peptide Mass and Peptide Fragmentation with MZCal: A User-Friendly Phone-Compatible Application

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MZCal

Teaching resource

Supports iOS and Android

Mobile-compatible web application

Optimised for lab settings

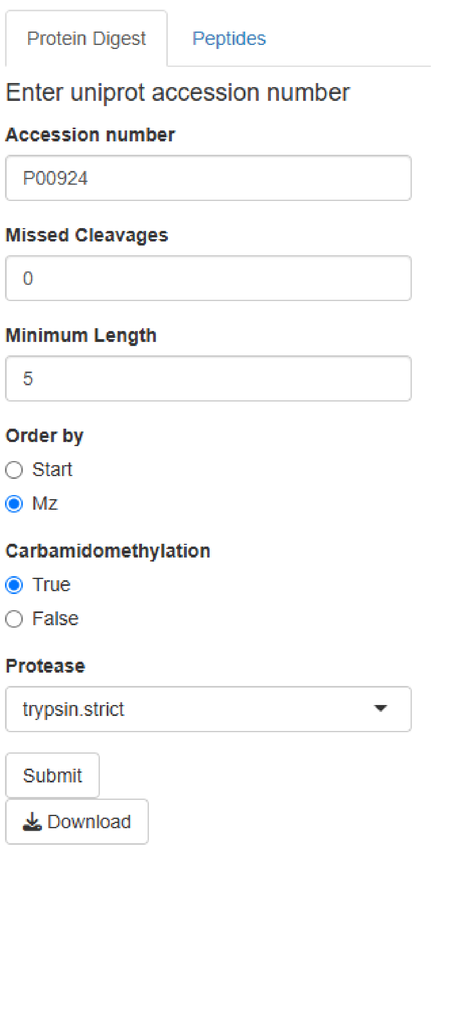
In Silico Digestion

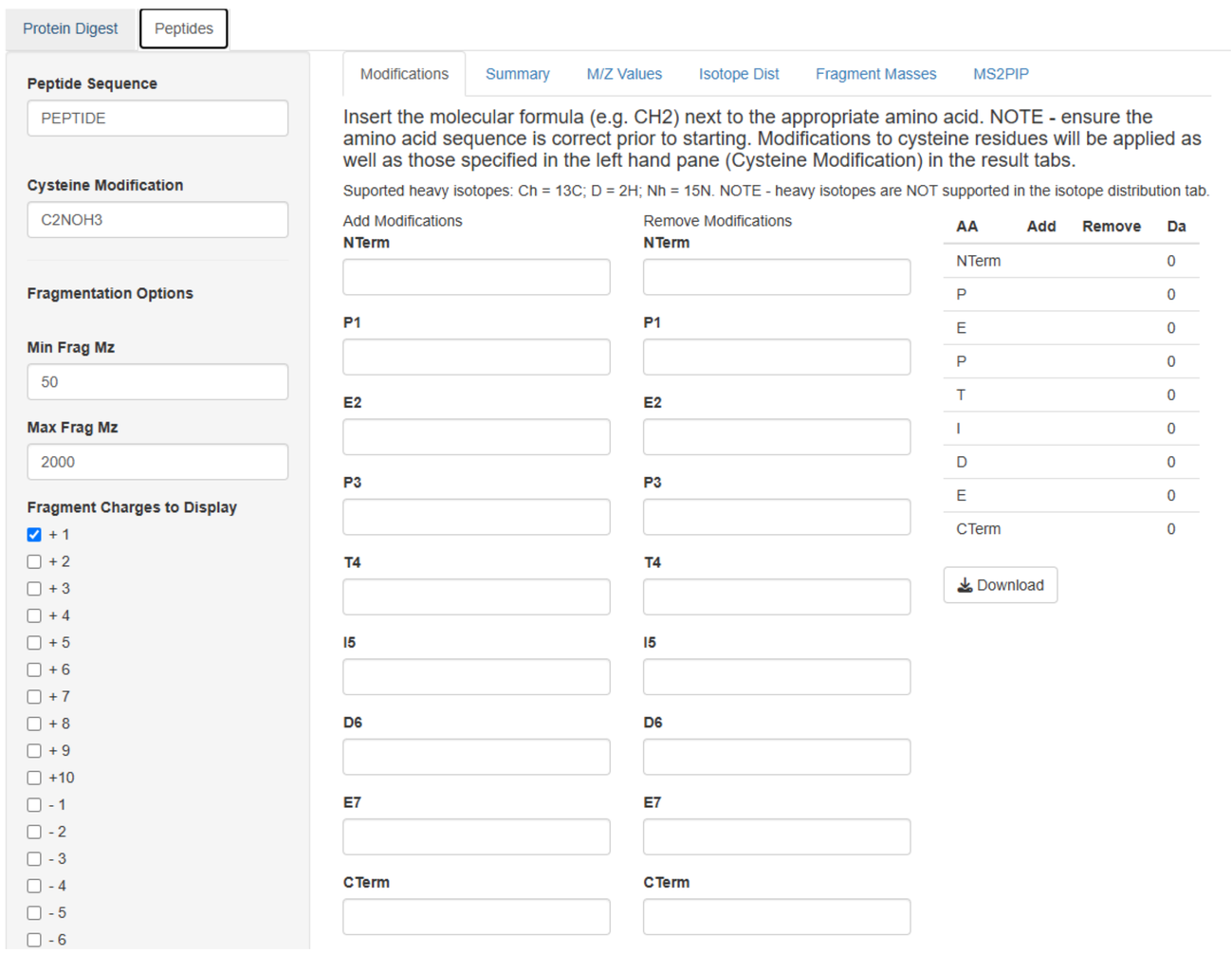
Peptide Mass

Dynamically Assign Modifications

Physiochemical Characteristics

Spectral Prediction MS2PIP *Degroeve (2012)*





TRY ME!

https://harrywhitwell.shinyapps.io/MZCal/

Publication

https://doi.org/10.1016/j.jprot.2025.105456

Sven Degroeve, Lennart Martens, MS^{2}PIP: a tool for MS/MS peak intensity prediction, *Bioinformatics*, Volume 29, Issue 24, December 2013, Pages 3199–3203, https://doi.org/10.1093/bioinformatics/btt544

Waters

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