

Relevant Experiences

- Research Project *uPEPperoni: A 5' UTR Short Open Reading Frame searching tool*
- Supervisor Associate Professor Joe Rothnagel
- Description A tool for searching of sORF in user's input and similar sORF in other protein sequences within the program database and visualization of their positions using heatmap.
- Contribution Revamping code base and bug fixing.
Shifting part of the database from file based to RDBMS.
Integrating and improving functionality.
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- Research Project *SWATHLib, an automated workflow for theoretical ion library generation for glycopeptides*
- Supervisor Associate Professor Benjamin L. Schulz
- Description DDA based spectral library together with complexity from post-translational modification are a suboptimal representation of ions within SWATH data. A workaround for this problem is a tedious process of creating a manually curated ion library. SWATHLib is created to address these problems.
- Contribution Writing of the JavaScript front-end using Angular 7 for automated user input collection.
Implementing a Python back-end for processing of user requests.
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- Research Project *Evolution and constraints of N-glycosylation Sequons*
- Supervisor Associate Professor Benjamin L. Schulz
- Description Analysis of N-glycosylation sequons in Homo sapiens glycoprotein homologs, their alignment and phylogenetic relation.
- Contribution Programming of automation workflow, logic and analyses