



# Knowledge-based CI and ML in Biomedicine

Data Mining Laboratory Classes



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# Data Mining Assignments



- ✓ Download any dataset of biomedical data (vectorized or sequential) where you can search for frequent patterns or association rules.
- ✓ Make yourself familiar with [the data mining library SPMF](#) containing more than two hundred data mining and knowledge exploration methods:
  - association rule mining,
  - itemset mining,
  - sequential pattern
  - sequential rule mining,
  - sequence prediction,
  - periodic pattern mining,
  - episode mining,
  - high-utility pattern mining,
  - time-series mining,
  - clustering and classification.



# Data Mining Assignments

- ✓ Implement one of the methods (Apriori, Eclat, FP-tree) presented during the lecture or reuse (using [wrappers](#) for Python) a selected method from [the data mining library](#).  
You can also use any other Python library or methods built in Python after your choice!
- ✓ Use the implemented or adapted method(s) from the library to find:
  - all frequent patterns in the dataset meeting the defined criterium of the minimum support;
  - all closed and maximum patterns in the dataset meeting the defined criterium of the minimum support;
  - all associative rules meeting the defined criteria of the minimum support and the minimum confidence;
- ✓ Present your final results (found patterns and associative rules) and check whether the computed results are correct.
- ✓ Send the code and the results as an answer to the assignment in MS Teams.

# Literature and Bibliography



1. Robert Layton, [Learning Data Mining with Python](#), Second Edition, O'Reilly.
2. Robert Layton, [Learning Data Mining with Python \(on-line\)](#), Second Edition, O'Reilly.
3. Useful [Python libraries for CI and DM](#).
4. Python libraries and modules for data mining:
  1. [Numpy](#)
  2. [Scipy](#)
  3. [Pandas](#) – Python Data Analysis Library
  4. [Scikit-learn](#)

