

SESSION SIXTEEN

RUN DATA

- Cushaw3
- In Cushaw3.0.3:
 - Make bwt file
 - `./cushaw3 index -p <bwt_file_base> <read data>.fasta`

JOB SUBMISSION SCRIPT

- Use CHY and ATRI data

BIO PYTHON

- Set of tools stored in package for computational biology

DYNAMIC PROGRAMMING

- Method of problem solving by solving subproblems
 - Breaking down problems
- Problems get progressively larger as components are broken down
- Why may this be the best method for Longest Common Subsequence finding?

STARTING OUR SCRIPT