Active learning strategies for bioinformatics teaching

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CEES is instituted in the control of the colored and Evolutionary Synthesis



Background

- After implementing active learning techniques, I find my teaching has become more interesting, more active, more
 interactive, and thus hopefully better
- This poster summarizes and demonstrates the techniques I have implemented my bioinformatics curriculum for analysis of high-throughput sequencing data (part of the INF-BIO 5121 and 9121 courses)

The assembly exercise Pen and paper

- Students receive printed sheets with short 14 character snippets ('reads') randomly drawn from a real English text
- In groups of 4 to 5 students, students try to reconstruct the source text.
- Students are very active and really make a collaborative effort, and seem to be enjoying it
- Enables a discussion of many important aspects of the challenges of genome assembly





ivory.idyll.org/blog/the-assembly-exercise.html

A class of active learners
Source: https://twitter.com/steeljawpanda/status/636233683444350976

Building small graphs Interactive notebook

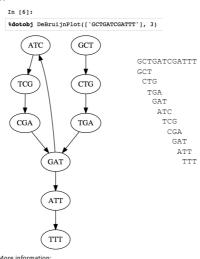
- Students explore small De Bruin graphs in an interactive electronic notebook (Jupyter) with python code
- The code visualizes the nodes and the effect of choosing different kmer sizes on the graph's structure

Exercise 1: simple De Bruijn graph

Do the exercise by execute the code cell below

Let's first generate a really simple De Bruijn graph of a short read 'GCTGATCGATTT' with kmer size 4 $\,$

Study the graph and see if it indeed encodes the read now change the kmer size to 3 and again trace the read: what happened?



More information:

fixlexblog.wordpress.com/2014/11/04/on-the-benefits-of-open-for-teaching

Quiz!

Are you using active learning in your teaching or are you planning to?



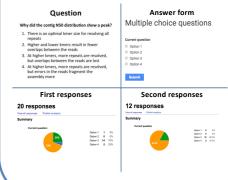
Finding the optimal settings Google spreadsheet

- Students collaboratively explore parameter space by each picking a number (kmer size), and running a velvet assembly using that number (kmer)
- Students log the resulting assembly metrics on an editable google spreadsheet, which has a graph showing the assembly results for the different settings
- The graph is shown on the presenter's screen in the classroom and updates in real time
- The goal is to teach the effect of choosing different kmer sizes for De Bruin Graph assembly



Peer instruction Google form

- I pose a multiple choice question
- Students discuss the question in pairs
- Each pair gets to vote through an online system, I use a simple google form for this
- I show the tally of the votes, using the 'summary' function of the google form
- Student pairs discuss the tally and consider whether they want to change their opinion
- Students vote again and the tally is shown again
- The answers are discussed by me



Credited to Eric Mazur: 'ConcepTests'

Further reading

https://flxlexblog.wordpress.com/2015/08/31/activelearning-strategies-for-bioinformatics-teaching-2/

Active learning' techniques benefit learning. Freeman et al (2014) PNAS 10.1073/pnas.1319030111

Get a copy of this poster through figshare: http://dx.doi.org/10.6084/m9.figshare.1541056 or scan the QR code



