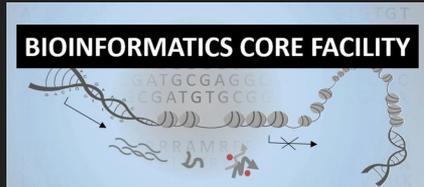


Welcome to the ABC

abc.au.dk



13.June 2024



Health
Data Science
Sandbox

The ABC concept

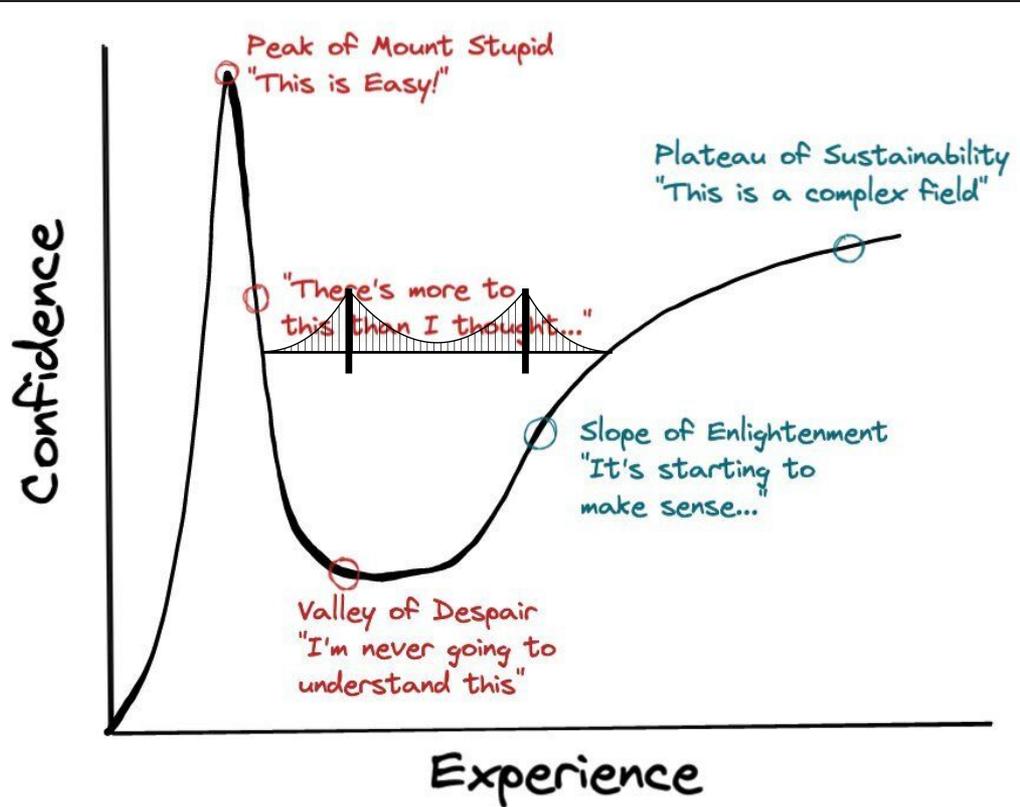
- Bi-weekly Informal venue
 - Independent sessions
 - **All documented at abc.au.dk**
- Open to any coding/bioinf experience level
- Integrated into the supporting infrastructures

The ABC format

- 15 min intro
 - What's new
 - Accessible delving into a topic
- Anyone showing something cool or useful?
- Coding support and code-together
 - python, R, bash
 - HPC, technicals
 - various omics (scRNA, bulkRNA, genomics, ...)
- Periodical proposal of workshops, tutorials, in-depth sessions

The ABC aim

- Reducing or avoiding the frustration phase
- Learn new things without the stress of formal meetings
- Stay up-to-date
- Know what other people do to code and analyze data



Who we are & What we do

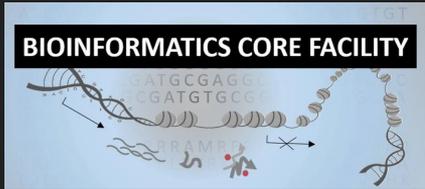
```
print("Hello world!")

n <- readLines("stdin")
print(paste("Entered value: ",n))

#Slice
slices
lbls <-
"France
pie(s1
Country:

#if stat
x = 0.1
if( x < 0.
{
  x <-
}

#for stat
x <- c(1,2,4,8,16)
for (loop in x)
{
  print(loop)
}
```

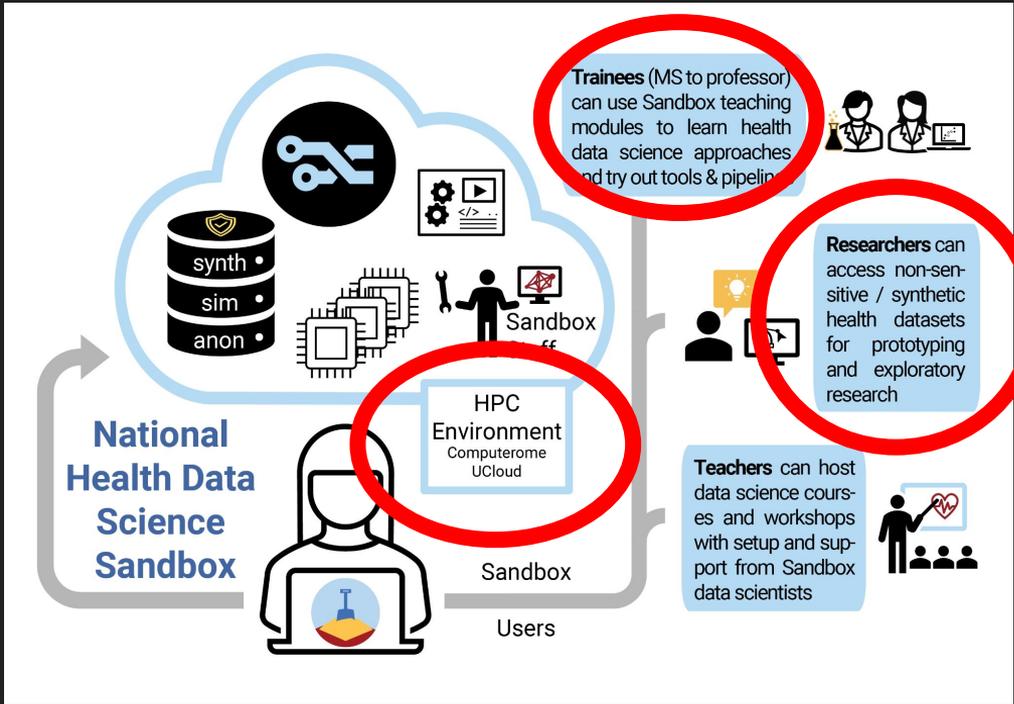
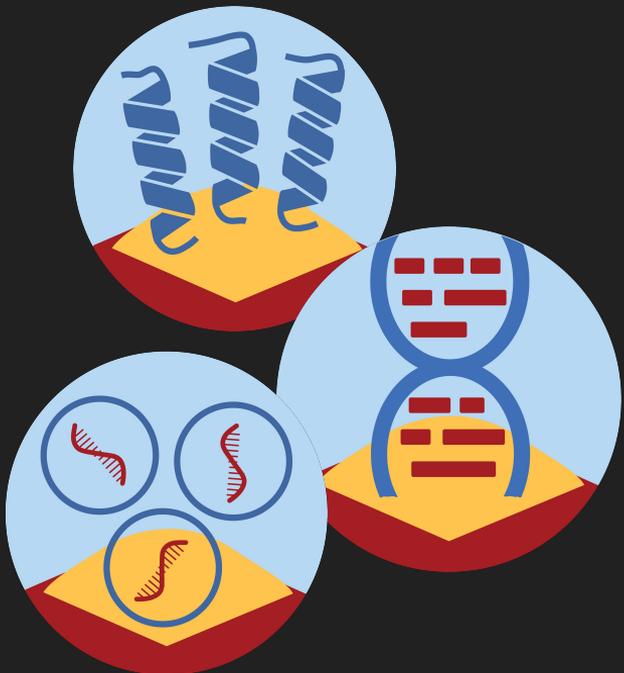


Health
Data Science
Sandbox



Samuele

Health Data Science Sandbox @ BiRC





Samuele

What are my tools / What I can help with at the cafe

Languages

R
Python
Bash
Docker
Some C++
Matlab (!?)

Data

Genomics
Bulk RNA
Single cell RNA
- Spatial
- ATACseq
Some proteomics

Tools

HPC
Github
Conda
JupyterLab
Latex
Quarto / Markdown
Some RStudio

BIOINFORMATICS CORE FACILITY

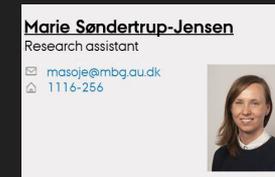
Established autumn 2022

Aim: to provide basic bioinformatics training, services as well as full-scale scientific collaborations for all scientists at Aarhus University as well as external academic and industrial partners.

- One priority is **to provide biologist-friendly solutions** to implement the use of OMICs and other large-scale data at the Department of Biomedicine.

Core activities:

- Data analysis
- Consultation
- Training
- Research collaboration
- Knowledge dissemination
- Fundraising



Now: poll

- At [menti.com](https://www.menti.com)
- Use code: 1412 3273

- The next ABC (and last before the summer holidays) is Thursday the 27th June 2024

Exercises

- A: download and install R
- B: basic R commands
- C: work on your own data