#### Bioinformatics and Bioinformaticians

Johan Nylander

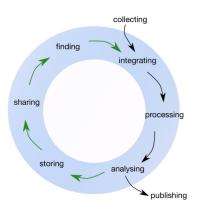
NRM.se / NBIS.se / SciLifeLab.se





#### Introduction

In a time where we see transitions towards biology as a data science, and towards a 'life cycle' view of research data



Griffin et al. 2017, F1000Research

#### Introduction



**EDUCATION** 

Unmet needs for analyzing biological big data: A survey of 704 NSF principal investigators

Lindsay Barone \*\*, Jason Williams\*, David Micklos\*

#### Introduction

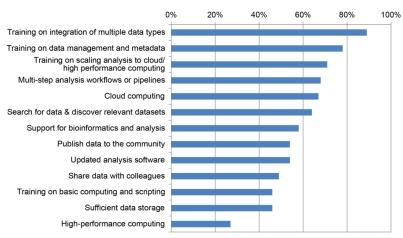


Fig 3. Unmet data analysis needs of National Science Foundation (NSF) Biological Sciences Directorate (BIO) principal investigators (PIs) (percent responding negatively, 318  $\leq n \leq$  510).

https://doi.org/10.1371/journal.pcbi.1005755.g003

## Bioinformatics - What's in a Word?



"Scientist studying DNA sequences"

 "The application of information technology to the field of molecular biology" (wikipedia.org)

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- Collection, archiving, organization, and interpretation of biological data
- Use of computers to address problems that are specific to biologists
- ► Informational technology + computer science + mathematics and statistics

# A Bioinformatician is, then, Someone who...

- Applies information technology to the field of molecular biology
- ► Collects, archives, organizes, and interprets biological data
- Uses computers to address problems that are specific to biologists
- Uses informational technology + computer science + mathematics and statistics

Hi, my name is Johan, and I'm a Bioinformatician

Anyone else who wants to share?

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Individual with a superior understanding of biology,
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- Have strong communication skills and ability to tell a story using numbers and data

(The following are quotes from job ads)

- Individual with a superior understanding of biology,
   computer science, and the latest trends in analyzing big data
- Have strong communication skills and ability to tell a story using numbers and data
- Ability to apply advanced statistical methods, machine learning, data science to molecular biology

- Strong background in programming and algorithms
- ► Fluent in one programming language (Python, C, C++ or Java) and familiarity with scripting languages (bash, Perl, R)

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- ► Fluent in one programming language (Python, C, C++ or Java) and familiarity with scripting languages (bash, Perl, R)
- Expert knowledge of Unix operating system, and High-performance computing (HPC) systems

- Have hands-on experience with bioinformatics methods appropriate for NGS applications (targeted sequencing, RNASeq, CHiPSeq, and de novo genome assembly)
- Have hands-on experience with genome alignment, mapping, variant calling and annotation (e.g. BWA, Bowtie, STAR, GATK, samtools, bcftools)

 Provide advanced bioinformatics analyses within research projects

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- Development of tools and workflows for such analyses

 Develop custom databases and web portals for managing raw and processed experimental data

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- Design and implement reusable bioinformatics analysis pipelines for processing next-generation sequencing, microarray, genomics, proteomics and chemogenomics data

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- Design and implement reusable bioinformatics analysis pipelines for processing next-generation sequencing, microarray, genomics, proteomics and chemogenomics data
- Develop novel algorithms and integrated data visualization applications when existing software packages are not available or are not adequate

Educating other scientists in bioinformatics through collaboration within supported projects, teaching at national courses, and through participating in various networks

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- Participating in the writing of scientific articles

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- Perform other duties as required

# One Real Example of Skills Requirements

#### The following profiles are expected:

- 1. Analysis of standard HTS data (variant calling, genome assembly, differential expression, ...),
- 2. Metagenomics
- 3. Metatranscriptomics
- 4. Single-cell analysis
- 5. Analysis of metabolomic data
- 6. Epigenomics and epitranscriptomics
- 7. Flux cytometry data analysis
- 8. Biological databases
- 9. Proteomics (Mass Spectrometry for Biology)
- 10. Omics data analysis and integration

- A Bachelor's degree in life sciences or computational sciences is required, although a MSc or PhD is preferred.
- Master's degree in Computer Science or Bioinformatics or a related computational discipline, or equivalent experience
- ▶ PhD or MSc in Bioinformatics, Computer Science, Computational Biology, Electrical Engineering/Signal processing or a related field with > 1 years' experience

## A Bioinformatician is Not a Unicorn

 A biologist may have broad knowledge of biology, but will have expertise in just part

## A Bioinformatician is Not a Unicorn

- A biologist may have broad knowledge of biology, but will have expertise in just part
- Similarly, a bioinformatician is not universal

# Common Expected Duties?

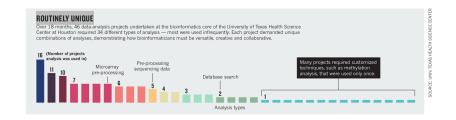
- Organize other people's data
- Analyze other people's data
- Educate other people in the field of bioinformatics
- ► **Support** other people in analysis
- Develop tools for other people to use

# Things that Bioinformaticians Do - and How Much of What?

- ► Organize ~ 20%
- **► Analyze** ~ 20%
- **► Educate** ~ 20%
- **► Support** ~ 30%
- **▶ Develop** ~ 10%

# Things that Bioinformaticians Do - and How Much of What?

#### 85% of the time a bioinformatician does "research"!



Chang 2015 Nature Communications

# Career Opportunities and Salaries for Bioinformaticians?

- Need to offer career opportunities for bioinformatitians - Can you?
- Academic suicide if not first or last author in a world based on counting publications
- Competetive salaries in academia?

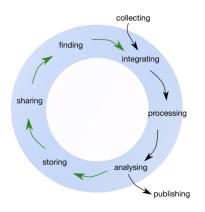
# Getting Bioinformatics Done - By Whom?

- ➤ You?
- ► Your students?
- ► Your postdocs?
- ► Your collaborators?
- **.**..
- ► A company?

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### What Should I Learn?



#### What Should I Learn?

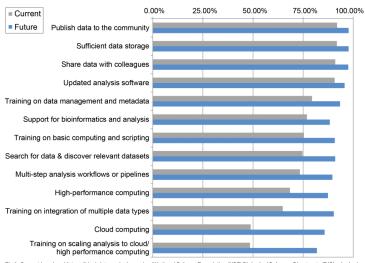
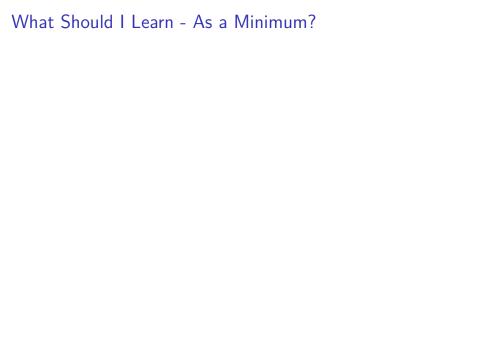


Fig 2. Current (grey) and future (blue) data analysis needs of National Science Foundation (NSF) Biological Sciences Directorate (BIO) principal investigators (PIs) (percent responding affirmatively,  $387 \le n \le 551$ ).



#### What Should I Learn - As a Minimum?

- Be able to look at your data?
  - Locate your data
  - Learn how to identify the "non-reproducible" parts of the data
  - ► Transfer data
    - **.**

#### What Should I Learn - As a Minimum?

```
$ sshfs user-delivery01234@grus.uppmax.uu.se: ~/grus
$ firefox ~/grus/P12345/00-Reports/*multiqc_report.html
$ backup2nrm ~/grus/P12345
```

## What Should I Learn - As a Minimum?

- ▶ Be able to look at your data?
- Other suggestions?

# Do I Need to Learn Programming?

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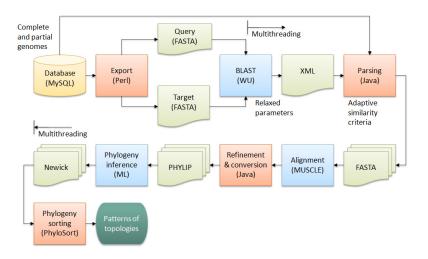
No, but Why not?

- Actually programming can be easy
- Doesn't require knowledge of computer science

► A script is like a protocol:

```
for infile in *.fas
do
  alifile="${infile%.fas}.ali"
  # 1. Do multiple sequence alignment
  mafft --auto "$infile" > "$alifile"
  # 2. Infer a phylogenetic tree
  igtree -s "$alifile" -nt AUTO -m TEST
done
```

▶ You start with an existing one and tweak it

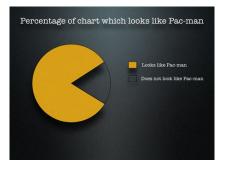


You have a big advantage over the bioinformaticians:

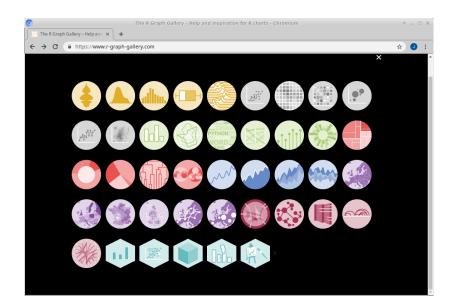
- You generated the data
- You understand the biology
- You understand how things go wrong

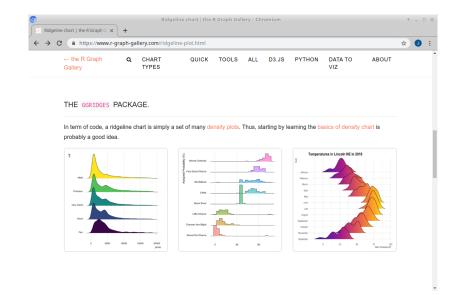
#### Learn How to Visualize Your Data

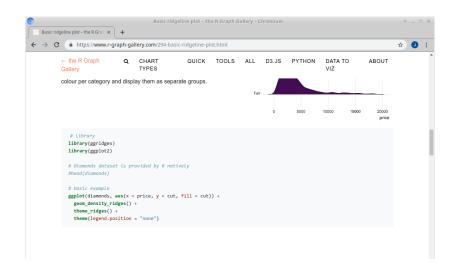
- Great for learning about the data
- Easy to pick out ouliers and trends



https://www.r-graph-gallery.com/







```
library(ggridges)
library(ggplot2)
ggplot(diamonds, aes(x = price, y = cut, fill = cut)) +
  geom_density_ridges() +
  theme_ridges() +
  theme(legend.position = "none")
```

# Do I Need to Learn Programming?

► Well, up to you (your loss...)

## Do I Need to Learn Programming?

- ► Well, up to you (your loss...)
- Have your student(s) learn programming!
  - Include a programming course as part of their PhD
  - Have your student(s) gather other students in "study circles"

#### Courses and Teach Yourself

https://tess.elixir-europe.org/

