



Open-source, transparent, and reproducible bioinformatics

PUBS2015

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National Microbiology Laboratory
Public Health Agency of Canada

Overview

- [Who are we?](#)
- [Data](#)
- [Bioinformatics](#)
- [Open-source software](#)
- [Reproducibility](#)
- [Galaxy](#)

Who Are We?

- Franklin Bristow

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 <https://github.com/fbristow>

- Eric Marinier

 eric.marinier@phac-aspc.gc.ca

 <https://github.com/emarinier>

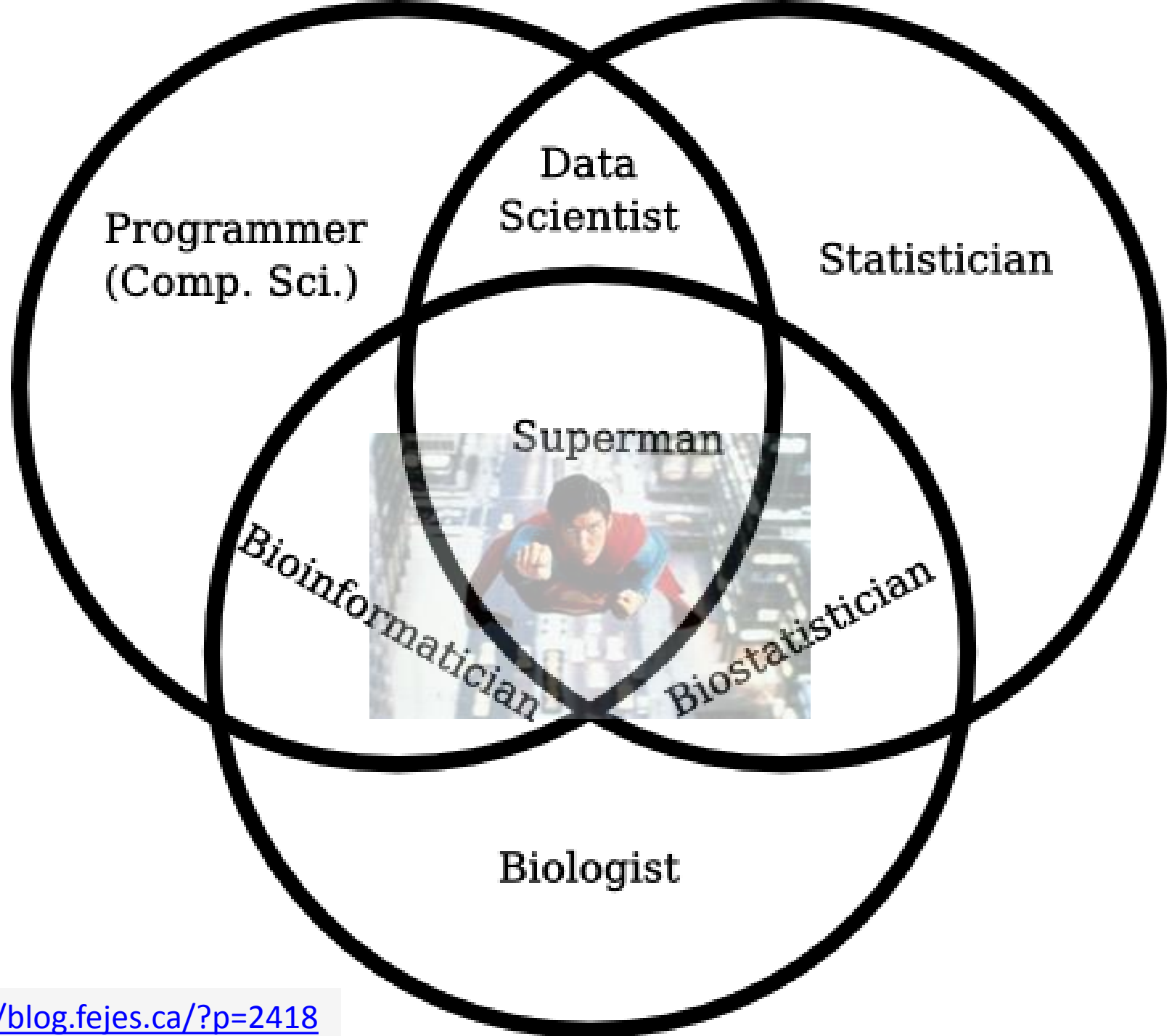
Data



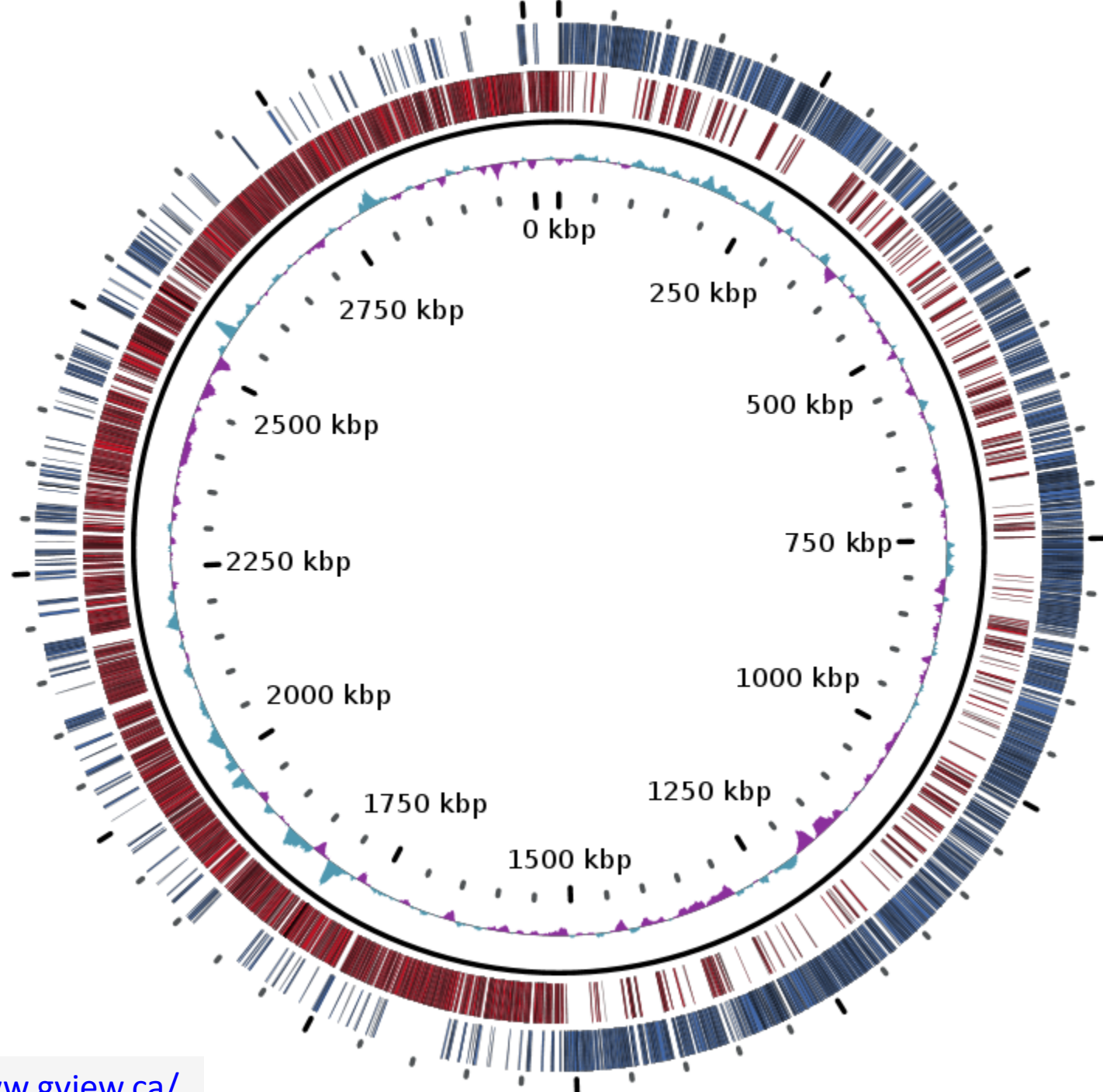
Data



http://prodstatics3cdn1.purewow.com/images/articles/2014_08/Drought400x290.jpg







Open- Source Software



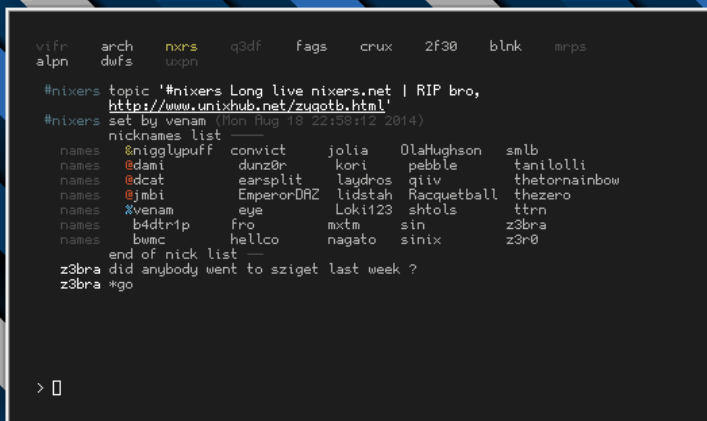
http://en.wikipedia.org/wiki/Richard_Stallman

Open-Source Software



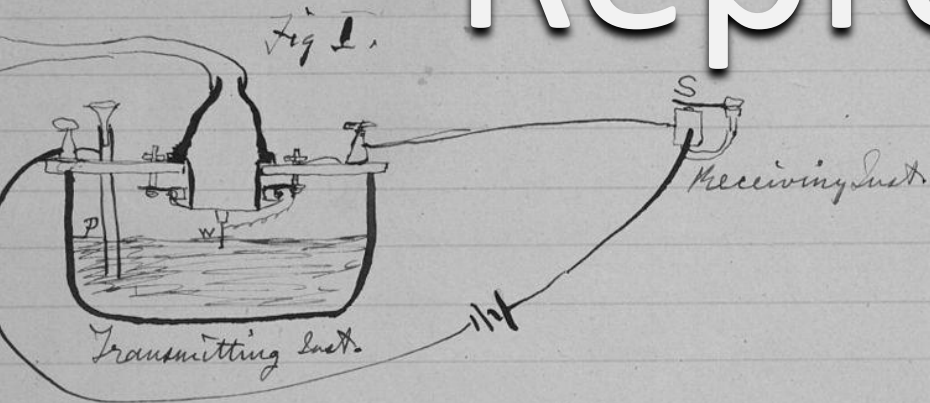
<http://en.wikipedia.org/wiki/Tux>
http://en.wikipedia.org/wiki/Linus_Torvalds

Command-line Software



March 10th 1876

Reproducibility



The improved instrument shown in Fig. 1 was constructed this morning and tried this evening. P is a brass pipe and W the platinum wire. M the mouth piece and S the armature of the Receiving Instrument.

Mr. Watson was stationed in one room with the Receiving Instrument. He pressed one closely against S and closed his other with his hand. The Transmitting Instrument placed in another room and the doors of rooms were closed.

I then shouted into M the following message: "Mr. Watson - Come here - I want to

I asked him to repeat the words. He answered "You said 'Mr. Watson - come I want to see you'." We then changed places and I listened at S while Mr. Watson read a few passages from a book into mouth piece M. It was certainly that articulate sounds proceeded from S effect was loud but indistinct and muffled.

If I had read beforehand the passages by Mr. Watson I should have recognized every word. As it was I could make out the sense - but an occasional word here and there was quite distinct. I made out "to" and "out" and "for" and finally the sentence "Mr. Bell do understand what I say? Do - you - der - stand - what - I - say" quite clearly and intelligibly. It was audible when the armature S was

moved -
http://en.wikipedia.org/wiki/Lab_notebook



Reproducibility

Galaxy

https://usegalaxy.org

Galaxy

Analyze DataWorkflowShared DataVisualizationCloudHelpUser

Using 0 bytes

Tools

search tools

Get Data

Lift-Over

Text Manipulation

Convert Formats

Filter and Sort

Join, Subtract and Group

Extract Features

Fetch Sequences

Fetch Alignments

Get Genomic Scores

Statistics

Graph/Display Data

Phenotype Association

snpEff

BEDTools

Genome Diversity

EMBOSS

Regional Variation

FASTA manipulation

Evolution

Multiple Alignments

Metagenomic analyses

Motif Tools

NGS TOOLBOX BETA

NGS: QC and manipulation

NGS: Mapping

NGS: SAM Tools

NGS: Peak Calling

NGS: Variant Analysis

NGS: RNA-seq

Operate on Genomic Intervals

NGS: GATK Tools (beta)

NGS: VCF Manipulation

NGS: Picard (beta)

Galaxy is an open source, web-based platform for data intensive biomedical research. If you are new to Galaxy [start here](#) or consult our [help resources](#).



Tweets

 Galaxy Project @galaxyproject 56m @Triconference? @jtx speaking on Accessible and Reproducible Large-Scale Analysis with Galaxy at 2:20 #usegalaxy #TRICON Expand

 Galaxy Project @galaxyproject 4h Galaxy code is now on @github github.com/galaxyproject/... #usegalaxy pic.twitter.com/O9dohBqgv Show Photo

 Galaxy Project @galaxyproject 17 Feb Visualize your tabular data in Galaxy using the Charts tool. Details &

Tweet to @galaxyproject

History

search datasets

Unnamed history

0 bytes

This history is empty. You can load your own data or get data from an external source

PENNSSTATE



The Galaxy Team is a part of the [Center for Comparative Genomics and Bioinformatics](#) at Penn State, and the [Department of Biology](#) and at [Johns Hopkins University](#).

JOHNS HOPKINS UNIVERSITY

This instance of Galaxy is utilizing infrastructure generously provided by the [iPlant Collaborative](#) at the [Texas Advanced Computing Center](#), with support from the [National Science Foundation](#).

TACC

iPlant Collaborative

The Galaxy Project is supported in part by [NSF](#), [NHGRI](#), [The Huck Institutes of the Life Sciences](#), [The Institute for CyberScience](#) at Penn State, and [Johns Hopkins University](#).

This is a free, public, internet accessible resource. Data transfer and data storage are not encrypted. If there are restrictions on

Galaxy

Galaxy

https://usegalaxy.org

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Analyze Data Workflow Shared Data Visualization Cloud Help User Using 0 bytes

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workflows

reproducible

biology

data

NGS

science

informatics

anal

Penn State

The Galaxy Team is a part of the Genomics and Bioinformatics at the Department of Biology and at the

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Unnamed history

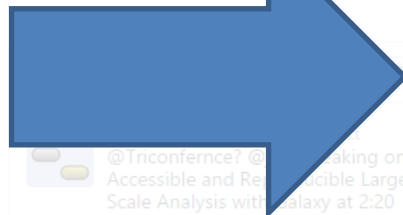
This history is empty. You can log in to own data or get data from an external source

Toolbox



http://cdn.toysrus.com.au/www/732/files/581070-home-depot-18-piece-tool-box.jpg

History



History

search datasets

Unnamed history

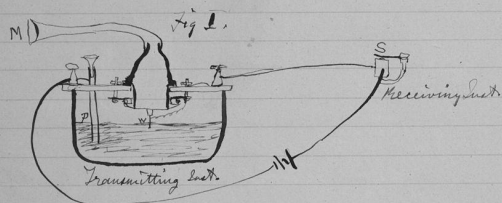
0 bytes

i This history is empty. You can [load your own data](#) or [get data from an external source](#)

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March 10th 1876



1. The improved instrument shown in Fig. 1. I was constructed this morning and tried this evening. P is a brass pipe and W the platinum wire M the mouth piece and S the armature of the Receiving Instrument.

Mr. Watson was stationed in one room with the Receiving Instrument. He pressed one ear closely against S and closed his other ear with his hand. The Transmitting Instrument was placed in another room and the doors of both rooms were closed.

I then shouted into M the following sentence: "Mr. Watson - Come here - I want to

see you." To my delight he came and declared that he had heard and understood what I said.

I asked him to repeat the words - ~~He said~~ He answered "You said 'Mr. Watson - come here - I want to see you'."

We then changed places and I listened at S while Mr. Watson read a few passages from a book into the mouth piece M. It was certainly the case that articulate sounds proceeded from S. The effect was loud but indistinct and muffled.

If I had read beforehand the passage given by Mr. Watson I should have recognized every word. As it was I could not make out the sense - but an occasional word here and there was quite distinct. I made out "to" and "out" and "further"; and finally the sentence "Mr. Bell do you understand what I say? Do-you-un-der-stand-what-I-say" came quite clearly and intelligibly. No sound was audible when the armature S was re-moved.

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Galaxy

https://usegalaxy.org/workflow/create#

Galaxy

Analyze DataWorkflowShared DataVisualizationCloudHelpUser

Using 0%

Your account has not been activated yet. Feel free to browse around and see what's available, but you won't be able to upload data or run jobs until you have verified your email address. [Resend verification.](#)

Tools

Workflow Canvas | Unnamed workflow

Details

NGS TOOLBOX BETA

NGS: QC and manipulation

Select high quality segments

Build base quality distribution

Draw quality score boxplot

Quality format converter (ASCII-Numeric)

Filter by quality

FASTQ to FASTA converter

ILLUMINA DATA

ROCHE-454 DATA

Remove sequencing artifacts

Barcode Splitter

Clip adapter sequences

Collapse sequences

Draw nucleotides distribution chart

Compute quality statistics

Rename sequences

Reverse-Complement

Trim sequences

FastQC:Read QC reports using FastQC

Combine FASTA and QUAL into FASTQ

Filter FASTQ reads by quality score and length

FASTQ Groomer convert between various FASTQ quality formats

Manipulate FASTQ reads on various attributes

FASTQ Masker by quality score

FASTQ joiner on paired end

Input Dataset

output

FastQC:Read QC

Short read data from your current history

Contaminant list

html_file (html)

Input dataset

Name:

Edit Step Attributes

Annotation / Notes:

Add an annotation or notes to this step; annotations are available when a workflow is viewed.

Workflow Editor

Links

- Lecture notes and exercises:
 - <https://github.com/emarinier/>
- Public Galaxy instances:
 - <https://usegalaxy.org>
 - <https://orione.crs4.it>