Materiais de apoio

Introdução ao uso de dados genômicos em Sistemática - BotInv2024

Em construção ...

Linguagens de programação para iniciantes

Bash

https://www.bioinformatics.babraham.ac.uk/training.html#sequenceqc

http://korflab.ucdavis.edu/bootcamp.html

https://lifehacker.com/a-command-line-primer-for-beginners-5633909

R

http://ecor.ib.usp.br/doku.php

https://www.datacamp.com/learn/r

Python

https://www.bioinformatics.babraham.ac.uk/training.html#sequenceqc

General (diversos cursos)

https://github.com/hbctraining/Training-modules

https://speciationgenomics.github.io/

Tutoriais processamento de dados NGS

https://bioinformatics.rbge.org.uk/ (RBGEs bioinformatics and data management)

https://github.com/sidonieB/Workflows (NGS Workflows, Sidonie Bellot)

https://github.com/LPDagallier/Phylogenomics_Workshop (Phylogenomics workshop, Léo-Paul Dagallier)

https://speciationgenomics.github.io/ (Speciation & Population Genomics, Mark Ravinet & Joana Meier)

https://github.com/carol-siniscalchi/IntroNGS (Introduction NGS App to Systematics, Carol Siniscalchi)

Referências