Introduction and housekeeping

Using Networks to Study Microbes

Alberto Santos & Yesid Cuesta-Astroz

Outline

- Who are you?

- Introduction to the course

- Course schedule

- Tools

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Who are you?

Alberto Santos - Associate Professor and Head of the Data Science Platform at the Danish Technical University (DTU) - Denmark

- Research Group Lead **Multi-omics Network Analytics** (MoNA)

  - **Multi-omics**

  Using multimodal data to have a comprehensive view on (micro) biology problems

  - **Network**

    Exploiting graphs to structure, represent, integrate and analyse data

  - **Analytics**

Applying Data Science and Machine Learning to answer complex biological questions

Who are you?

Yesid Cuesta-Astroz - Assistant profesor Instituto Colombiano de Medicina Tropical, Universidad CES

- Researcher ICMT - CES

  - Pathogen genomics and bioinformatics

    - Working on the study of the biological diversity of microorganisms from omics approaches.

    - Modelling the host-pathogen interactions using protein-protein interaction networks.

Who are you?

Introduction

Introduction to the Course
Networks to Study Microbes

- Technological advances are making **large-scale omics datasets** available to **study microbes and microbial communities**

- We need **computational methods** to **model and understand** these complex data

- This course will provide an overview of **omics data types** and **data resources** available and how to **integrate**, **analyse** and **interpret** these data with **networks**
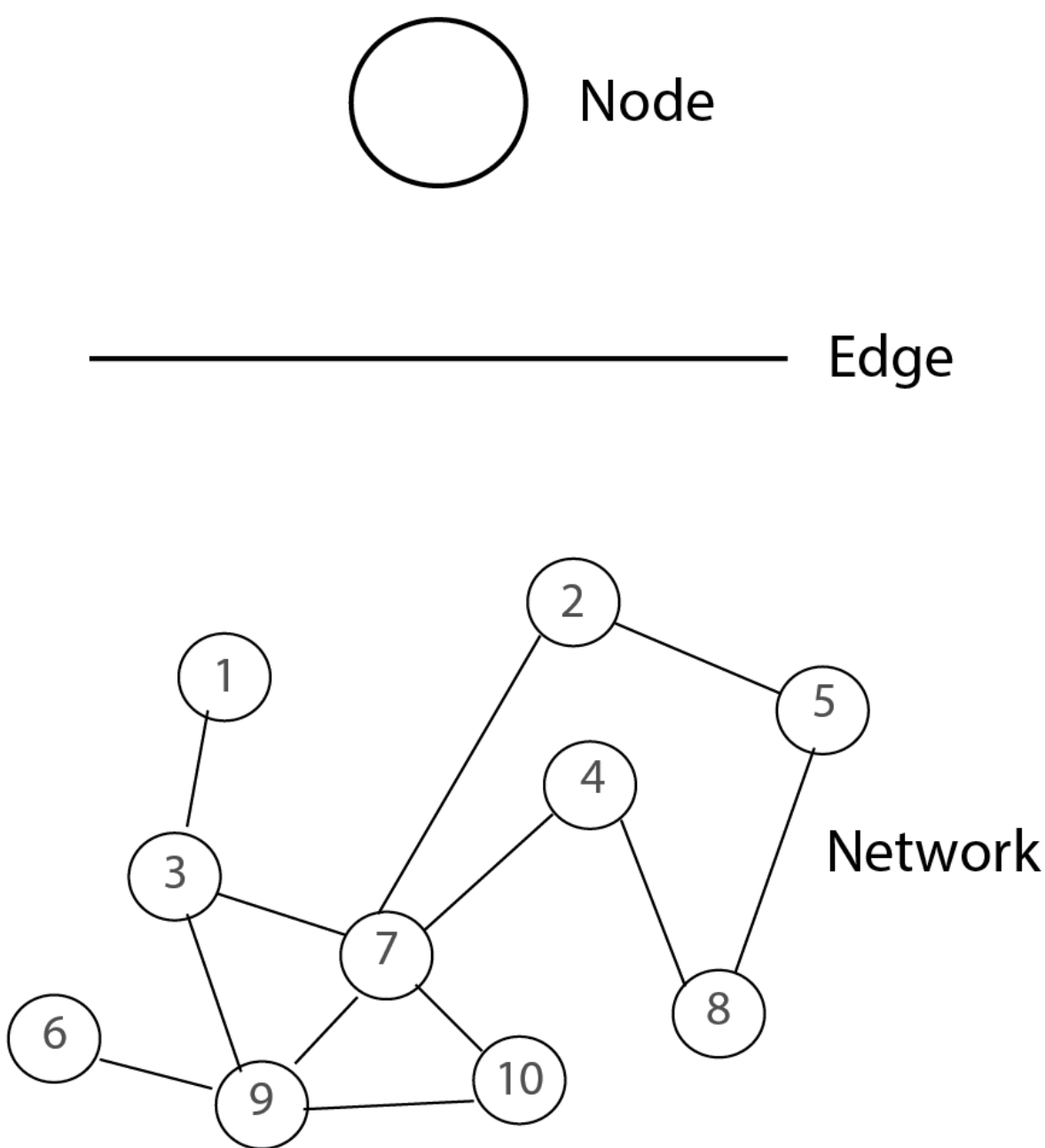
Introduction to the Course

What are networks?

- **Data structures** of components (**nodes**) connected by relationships (**edges**)

- These structures allow:

  - Quick **integration** of **heterogeneous data** based on relationships

  - **Graph theory** methods can be used to **analyse** and **interpret** data, e.g., topological properties can be used to explain:

    - The possible **role** of specific components

    - The **flow** of information

    - The **robustness** of the system
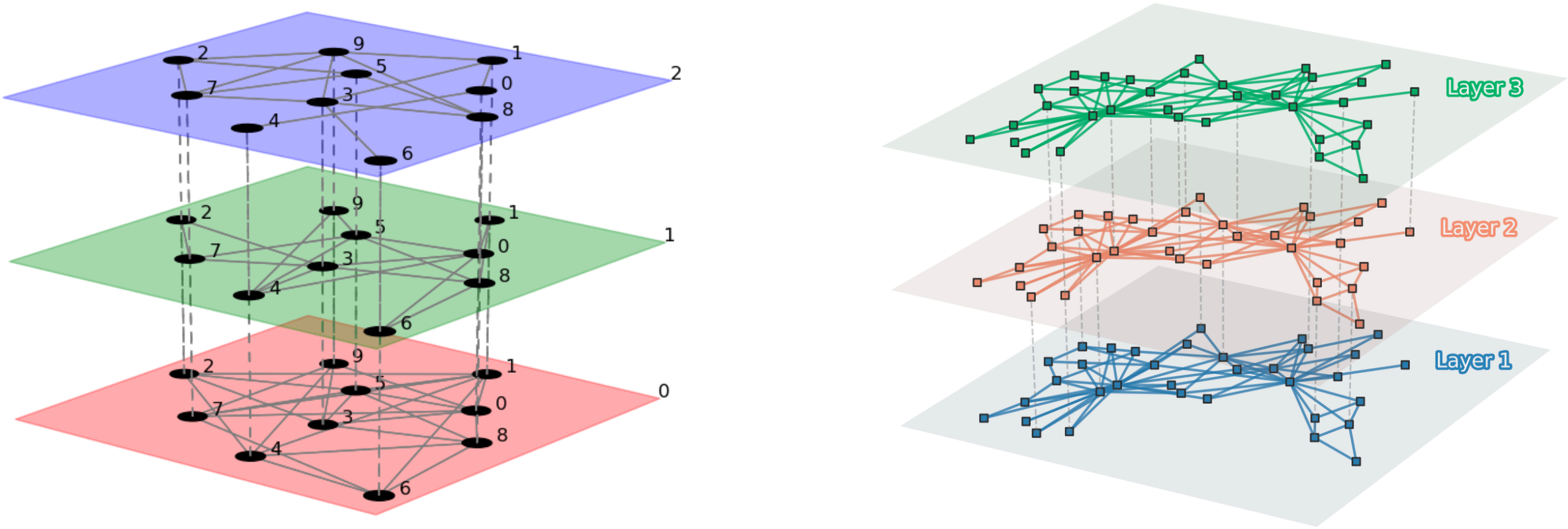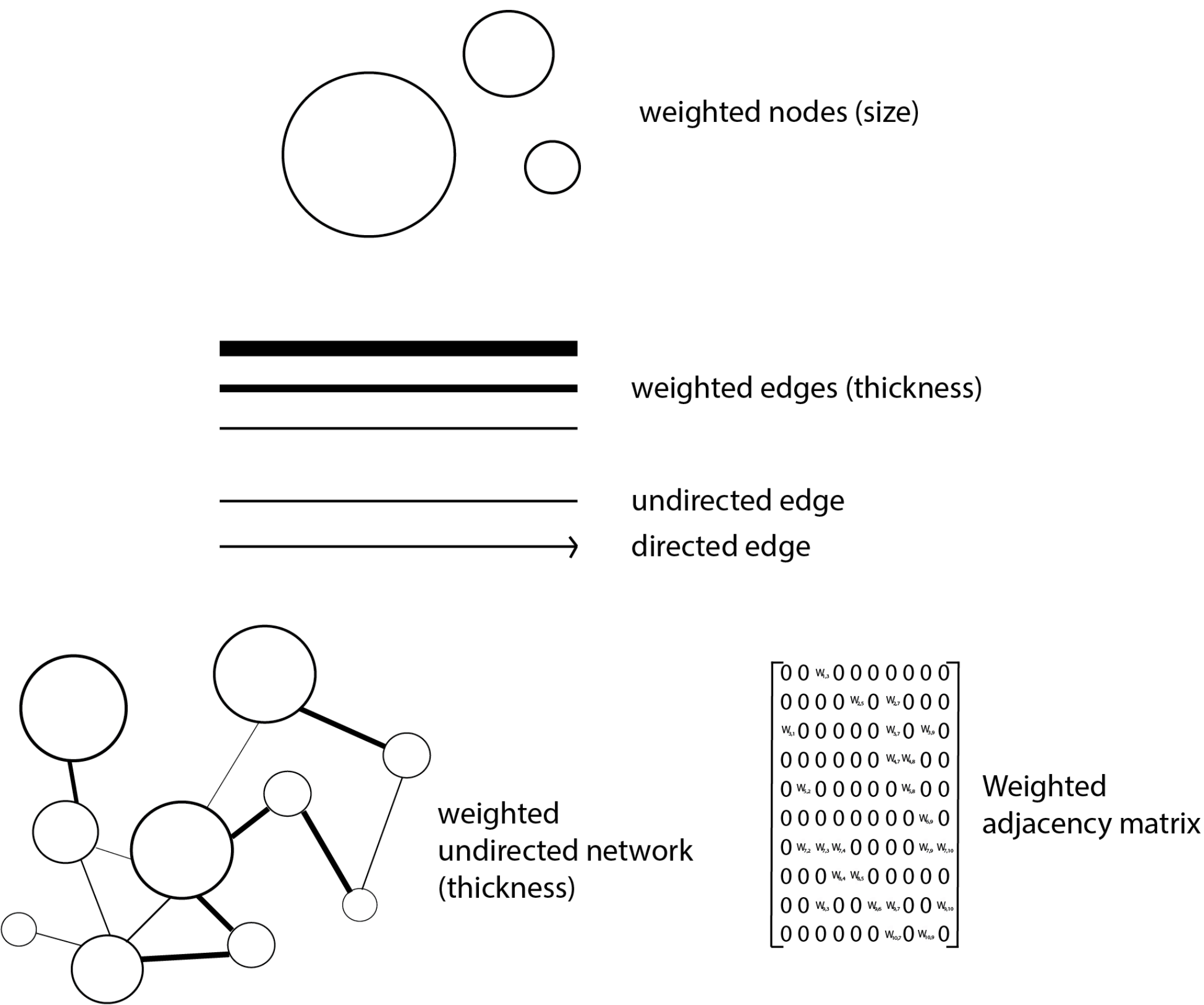
  - **Visualize** data

Networks

Node

Edge

weighted nodes (size)

weighted edges (thickness)

undirected edge

directed edge

Network

Adjacency matrix

$$\begin{bmatrix} 0&0&1&0&0&0&0&0&0&0 \\ 0&0&0&0&1&0&1&0&0&0 \\ 1&0&0&0&0&0&1&0&1&0 \\ 0&0&0&0&0&0&1&1&0&0 \\ 0&1&0&0&0&0&1&0&0 \\ 0&0&0&0&0&0&0&1&0 \\ 0&1&1&1&0&0&0&0&1&1 \\ 0&0&0&1&1&0&0&0&0 \\ 0&0&1&0&0&1&1&0&0&1 \\ 0&0&0&0&0&1&0&1&0 \end{bmatrix}$$

weighted
undirected network
(thickness)

Weighted
adjacency matrix

Schedule

Time	DAY 1	DAY 2	DAY 3
8:30-9:20	Introduction and Housekeeping	Working with Data in Python I	Analysing Networks I
9:20-10:10	An Omics View on Microbes I	Working with Data in Python II	
10:10-10:30	Coffee break	Coffee break	Coffee break
10:30-11:20	An Omics View on Microbes II	Visualizing Data in Python	Analysing Networks II
11:20-12:10	Data Resources	Visualising Networks I	
12:10-13:30	Lunch	Lunch	Lunch
13:30-14:20	Introduction to Python I	Visualising Networks II	Team Project
14:20-14:40	Coffee break	Coffee break	Coffee break
14:40-16:00	Introduction to Python II	Network Exercises	Team Project
16:00-16:50	Recap and Q & A	Recap and Q & A	Team Project Presentations and Q&A

Tools

Course Website
https://github.com/Multiomics-Analytics-Group/networks_to_study_microbes

- We use GitHub to maintain the contents of the course

- **GitHub** is a framework generally used to maintain and version control software development projects

- **Version control** is a system that records changes to a file or a set of files over time and who made them so that you can recall specific versions later
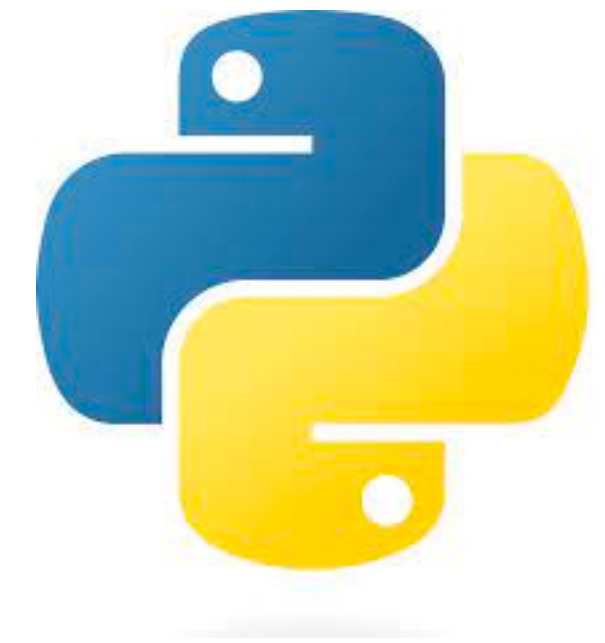
https://en.wikipedia.org/wiki/GitHub

https://git-scm.com/book/en/v2/Getting-Started-About-Version-Control

Basic Programming
Python

- Python is a **high-level, general-purpose programming language**

- It is a good language for both **beginners** and **advanced programmers**:

  - Its design philosophy **emphasises code readability**

  - Many **learning resources** available

  - **Large community** behind (i.e. stackoverflow for questions)

  - **Excellent Libraries** available, especially for data science

Jupyter Notebooks
https://jupyter.org/

- **Web-based development environment** for **creating**, **running** and **sharing Python** (and other languages) code

- A **notebook** is an interactive document that combines **live code**, **equations**, **text or markdown**, and **visualisations** (output of your code)

- Notebooks are divided into **cells** that **run sequentially**! (Need to pay attention)

- It **requires** having **Python installed** on your local machine

Colab Notebooks

https://research.google.com/colaboratory/faq.html

- Google Colab is based on **Jupyter Notebook** open source project **hosted on Google's servers**

- Advantages:

  - Requires **no setup** to use (no python installation)

  - Provides **free** access to **computing resources** on Google's servers including GPUs

  - **Notebooks** can be **shared** just as you would with Google Docs or Sheets.

  - You can **import** existing **Jupyter notebooks**

- **Own data and notebooks** need to be accessed through **Google Drive** — Need Google account

Cytoscape
https://cytoscape.org/

- An open source **software platform** for **visualising** and **analysing** complex **networks**

- Used for any kind of networks but **specialised** on **biological domains**:

  e.g, Molecular interaction networks and biological pathways and integrating these networks with annotations, gene expression profiles and other state data.

- **Additional features** are available as freely available **Apps** (https://apps.cytoscape.org/)

Relevant Links
Extra readings and resources

- **Network biology: understanding the cell's functional organisation.** Albert-László Barabási & Zoltán N. Oltvai. Nature Reviews Genetics 2004 (https://www.nature.com/articles/nrg1272)

- **Network analysis of protein interaction data** — Online tutorial — EMBL EBI (https://www.ebi.ac.uk/training/online/courses/network-analysis-of-protein-interaction-data-an-introduction/)

- **Network Biology: A short introduction to the core concepts** https://www.youtube.com/watch?v=H1bGk8PGvf8

- **Network Visualization: a short introduction to the core concepts of network layout and clustering** https://www.youtube.com/watch?v=OunX7ISRbgA

- **A Guide to Conquer the Biological Network Era Using Graph Theory.** Mikaela Koutrouli, Evangelos Karatzas, David Paez-Espino, Georgios A. Pavlopoulos. Frontiers in Bioengineering and Biotechnology 2020 (https://www.frontiersin.org/articles/10.3389/fbioe.2020.00034/full)

- **Python for Beginners** https://www.python.org/about/gettingstarted/

- **How to Use Jupyter Notebook: A Beginner's Tutorial** https://www.dataquest.io/blog/jupyter-notebook-tutorial/

- **Getting started with Python and Jupyter Notebooks** https://colab.research.google.com/github/jckantor/CBE30338/blob/master/docs/01.01-Getting-Started-with-Python-and-Jupyter-Notebooks.ipynb