

## Day 1 – Introduction & Setup

- Welcome, course goals, and expectations
  - **Bioinformatics primer:** Central Dogma (DNA → RNA → Protein) and applications in health tech
  - Installing **Python**, **Anaconda**, and understanding **Google Colab** (main platform)
  - Running your first Python script in Colab
  - Quick intro: What is a **Python library** and why do we use them?
- 

## Day 2 – Python Basics for Bioinformatics

- Python data types (strings, numbers, lists, dictionaries)
  - Loops and conditionals (if/else, for/while)
  - Functions: writing and reusing code
  - Importing libraries (Pandas, Biopython basics)
  - *Mini-task:* Write your first “**Hello Bioinformatics!**” program
  - *Mini-project:* **DNA Base Counter** – count nucleotides from a FASTA file
- 

## Day 3 – Working with Biological Data

- Understanding file formats: FASTA and CSV
  - Reading and parsing sequence files
  - Using **Biopython** for sequence operations
  - *Hands-on Project:* **GC Content Calculator** – extract DNA sequences & calculate GC %
  - Application in health tech: mutations, genetic markers, and diagnostics
- 

## Day 4 – Version Control with GitHub

- Why researchers & coders use GitHub (collaboration, version control, portfolio)
  - Creating a GitHub account
  - Uploading your first script (DNA Counter) to GitHub
  - *Mini-project:* Share your **GC Content Calculator** on GitHub
- 

## Day 5 – Introduction to Streamlit (App Development Basics)

- What is Streamlit & why it's perfect for bioinformatics tools
- Setting up your **first Streamlit app**
- Adding text, inputs, and file uploads

- *Mini-project:* Build a **DNA Base Counter App** (interactive version of Day 2 project)
- 

## Day 6 – Making Apps Interactive

- Adding buttons, sliders, and user inputs
  - Displaying results dynamically
  - Uploading FASTA files and showing results in tables
  - *Mini-project:* **Mutation Finder Tool** – search for specific sequences/mutations
- 

## Day 7 – Data Visualization in Bioinformatics Apps

- Why visualization matters in biology & medicine
  - Using **Matplotlib/Plotly** in Streamlit
  - Visualizing DNA/protein sequence lengths (histograms)
  - Color-coding mutations in sequence data
  - *Mini-project:* **GC Content Visualizer**
- 

## Day 8 – Bioinformatics Use Case Design

- Case studies: genomics, diagnostics, personalized medicine
  - Mapping **real-world problems** to **app features**
  - Workshop: Design your **own app idea**
  - *Activity:* Pitch your app concept to peers
- 

## Day 9 – Building Your Custom App

- Structuring your Streamlit app (best practices)
  - Importing and cleaning datasets
  - Adding interactivity & visualizations
  - Styling and branding your app (make it professional)
  - Debugging tips
  - *Mini-project:* Begin building **your custom app**
- 

## Day 10 – Demo Day & Next Steps

- Each participant demos their custom app

- Peer and mentor feedback
- Certificates of Participation & Innovation Awards
- What's next?
  - More advanced Python libraries (scikit-learn, NumPy)
  - Cloud deployment (Streamlit Cloud, Hugging Face Spaces)
  - Continuing your GitHub portfolio