

CS-1390/ PHY-1390-1  
Introduction to Machine Learning  
Monsoon 2024  
Assignment 2

## Clustering Task

**Data:** *Provided separately with the classroom post.*

### Data Description

| Column Name               | Description  |
|---------------------------|--|
| Variant                   | Indicates the specific variant or type for each sample.  |
| Sample_ID                 | Unique identifiers for each sample, often combining multiple reference codes.  |
| Sample_Code               | A unique code for each sample, typically used as the primary identifier for each record.                               |
| Entry_Date                | The date when the sample was recorded or added to the database (in standard date format).                              |
| Origin                    | Geographical origin of the sample, including country and region where applicable.                                      |
| Source_Category           | Describes the source of the sample, such as Natural_Source (environmental/other) or Medical_Source (clinical).         |
| Genetic_Group             | Genetic group identifier based on molecular analysis, useful for genetic comparisons.                                  |
| Intra_Group_Similarity    | A numerical value representing the genetic similarity within the same group.   |
| Inter_Group_Difference    | A numerical value representing the genetic difference between different groups.  |
| Reference_ID              | External reference identifier associated with the sample, used for cross-referencing with other databases.             |
| Genomic_Data              | Genomic data identifier for samples where molecular sequencing information is available.                               |
| Resistance_Profile        | Describes the resistance profiles detected for each sample, indicating genetic markers linked to treatment resistance. |
| Predicted_Characteristics | Characteristics predicted from molecular data or other methods, indicating the predicted properties of the sample.     |

**Any clustering algorithm can be used.**

### Submission Guidelines

1. Submit a python notebook (with cell outputs, we will not run your code).
2. Include an extensive report within the python notebook throughout your code. Use markdown.
3. Grading emphasis will be on coding and plotting/describing clusters.
4. Use of libraries is not allowed. Code everything from scratch.
5. Any code similarities with your peers and code from the internet, as well as any AI generated code will result in a 0, along with further severe consequences under discretion of the professor.