

# Fall 2025: CAP5610 – HW 1

In HW1, you will explore and compare **Feature Extraction** and **Feature Selection** using a high-dimensional biological dataset. The goal is to implement **Autoencoder** (feature extraction) and **Concrete Autoencoder** (feature selection), visualize and analyze results, and present your findings in a structured report.

#### Data:

• Gene Expression: <u>lncRNA 5 Cancers.csv</u>

• Cancer Types: KIRC, LUAD, LUSC, PRAD, THCA

• Features: 12,309 lncRNAs

# Tasks:

# Task 1: [25 points] Explore the Data

Load the provided dataset and see how many data points and features it has. Count how many samples belong to each type of cancer and make a simple bar chart.

# Task 2: [25 points] Classification Using All Features Employing SVM

Use those ~12K features to train an SVM with RBF kernel. Report (i) confusion matrix for one fold, (ii) the macro, micro, and weighted precision/recall/F1 along with the accuracy, (iii) plot AUC-ROC and AUC-PR curve. Use 5-flod cross-validation.

#### Task 3: [25 points] Autoencoder for Feature Extraction

Train an Autoencoder that reduces the features to 50 <u>latent</u> dimensions. Use those 50 features to train an SVM with RBF kernel. Report (i) confusion matrix for one fold, (ii) the macro, micro, and weighted precision/recall/F1 along with the accuracy, (iii) plot AUC-ROC and AUC-PR curve.

### Task 4: [25 points] Concrete Autoencoder for Feature Selection

Train a Concrete Autoencoder to select 50 <u>actual</u> features. Train an SVM with RBF kernel on those 50 selected features. Report (i) confusion matrix for one fold, (ii) the macro, micro, and weighted precision/recall/F1 along with the accuracy, (iii) plot AUC-ROC and AUC-PR curve.

#### **Hint:**

- **Autoencoder**: You can implement a simple Autoencoder using libraries like Keras/TensorFlow or PyTorch.
- **Concrete Autoencoder**: Use the concrete-autoencoder library for this task. It requires TensorFlow 2.12.\*. Install it using:

pip install tensorflow==2.12.0 concrete-autoencoder

SVM: Use the following for reproducibility:
svm = SVC(kernel="rbf", probability=True, random state=42)

#### **Submission:**

You must submit the following items in CANVAS:

• Report (MS word or PDF)



- O Describe the algorithms/approaches/tools used: (a) What it is or What it does, (b) How it does, and (c) Application.
- O Describe results: (a) Put Figure/Table number and Title: On top of the table, and bottom of the figure. (b) Describe the figure and table. (c) Your observation about the figure and table. (d) Conclusion.
- Source code (\*.py or Jupyter notebook)
  - o Must be well organized (comments, indentation, ...)
- File name: HW1\_lastName

You must submit the files **SEPERATELY**. DO NOT compress into a ZIP file. If you fail to provide all required information or files, you may be given zero score without grading.

## **Deadline:**

The deadline is 11:59pm Wednesday, September 10, 2025. Late assignments will not be accepted.