

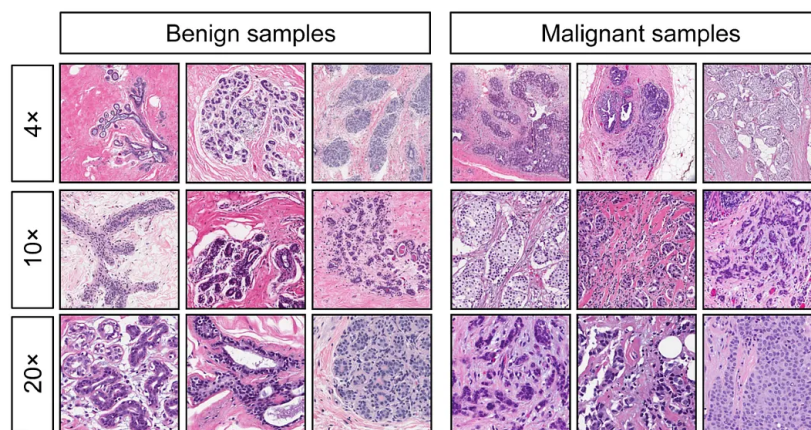


Quantum Computing and Machine Learning (614551008)

Practice 2 (2025-2026) Quantum Support Vector Machines

INSTRUCTIONS:

- **Deadline:** January 23rd, 23:59.
- **Objectives**
 - In this practice we will develop a quantum support vector machine classifier for the Breast Cancer Wisconsin dataset.
- **Dataset**



- “**Breast Cancer Wisconsin**” is a dataset that contains measurements of cell nuclei extracted from digitized images of breast masses.
- It includes **569 samples**, each described by **30 numerical features** that capture properties such as radius, texture, perimeter, area, smoothness, compactness, and symmetry, computed both as mean values and as variations within each sample.
- Each instance is labeled as either **malignant or benign**, providing a clean and well-balanced target for supervised learning.

- Because the features are continuous and moderately correlated, the dataset is especially suitable for tasks involving dimensionality reduction (like PCA), visualization, and testing classical and quantum machine-learning algorithms.
- The dataset can be easily downloaded from *scikit-learn* using the following instructions:

```
from sklearn.datasets import load_breast_cancer
data = load_breast_cancer()
X = data.data # Features
y = data.target # Labels
```

■ Tasks to be carried out

1. Preprocessing.

- Preprocess the dataset to prepare it to feed the quantum neural network: Since all the data is numeric we have only to normalize then using, for example, the `MinMaxScaler` or the `StandardScaler` from `scikit-learn`.
- Reduce the dimensionality using the PCA reduction from `scikit-learn` to reduce the number of qubits needed to represent the dataset in a quantum computer.
- Divide the data into train and test, using, for example, the `train_test_split()` method from `scikit-learn`.

2. Develop two SVM models to predict class of each case.

- Develop a classical model (i.e. use classical kernel function), using the `SVC` implementation in `scikit-learn` (<https://scikit-learn.org/stable/modules/generated/sklearn.svm.SVC.html>).
- Develop a quantum model (i.e. use quantum kernel function), using the `QSVC` implementation in `qiskit-machine-learning` (https://qiskit-community.github.io/qiskit-machine-learning/stubs/qiskit_machine_learning.algorithms.QSVC.html).

3. Run the models with different kernels/feature maps.

- Classical: linear, polynomial, RBF.
- Quantum: `ZFeatureMap`, `ZZFeatureMap`, `PauliFeatureMap`.

4. Results.

- Compare the results obtained.
- Use different metrics in addition to accuracy, keeping in mind that detecting a malignant case is more important than detecting a benign case.

5. Conclusions.

- Reach some final conclusions for the experiment carried out.

■ Submission

- The exercises will be developed using Jupyter Notebooks.
 - **The notebook should include:**
 - The practice can be carried out alone or in pairs, so the first cell of the notebook must be the full names of the authors.
 - Include all the code developed.
 - The code shall be accompanied by cells with an explanatory report containing a description of the process followed, detailing the results obtained and justifying the decisions taken.
 - The notebook will be saved with the results of its execution included.
 - **Submission process**
 - The exercises will be submitted to professor (Samuel Magaz Romero) by email (<mailto:s.magazr@udc.es>), with the subject of the email as per the following structure:
[MQIST - QSVM] Last Name(s), First Name(s)
For example:
[MQIST - QSVM] Magaz Romero, Samuel
- Submissions failing to meet this criteria may be automatically discarded.**
- There is a strict deadline for the assignment. Past due submissions will be rejected.