

# Code & Data Appendix

## Background

This appendix describes how to reproduce our experiments and include the source code, datasets, software & hardware setting, dependencies, and instructions for running.

## Repository Files

- |—data/
  - |—[dataset]/
    - |—[dataset]\_features.npy: The preprocessed features of [dataset]
    - |—[dataset]\_labels.npy: The labels of [dataset]
    - |—[dataset]\_pairs\_X.npy: The features of the pairs used for training the Siamese network of [dataset]
    - |—[dataset]\_pairs\_y.npy: The labels of the pairs used for training the Siamese network of [dataset]
- |—src/: The source code implementation of our approach

## Datasets

These are the datasets we used in our experiments and described in the paper.

As noted, the datasets were taken from [ODDS](#).

Dataset	#Samples	#Dim	Outliers
<a href="#">Anthyroid</a>	7200	6	7.42 (%)
<a href="#">Cardio</a>	1831	21	9.6 (%)
<a href="#">Mammo</a>	11183	6	2.32 (%)
<a href="#">Satellite</a>	6435	36	32 (%)

Dataset	#Samples	#Dim	Outliers
Seismic	2584	11	6.5 (%)
Thyroid	3772	6	2.5 (%)
Vowels	1456	12	3.4 (%)
Yeast	1364	8	4.7 (%)

## Dependencies

The required dependencies are specified in `environment.yml`.  
For setting up the environment, use [Anaconda](#):

```
$ conda env create -f environment.yml  
$ conda activate adtta
```

## Running the Code

### NOTES:

- A valid dataset name (instead of the mammo dataset) could be from the one described above.
- It is important to run the scripts from being inside `src/` (i.e., `$ cd src/`)

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### • Preprocessing

A preprocessing phase on the desired dataset.

```
$ cd src/  
$ python preprocess.py --dataset mammo
```

From this script, 4 files are going to be generated in the desired dataset `data/` folder:

- `mammo_features.npy`
- `mammo_labels.npy`

- mammo\_pairs\_X.npy
- mammo\_pairs\_y.npy

With these files, you can run the train and test.

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## • Train & Test

To train the estimator, as well as the Nearest Neighbors models (both Euclidean and Siamese), run:

```
$ cd src/  
$ python main.py --dataset mammo --neighbors 10 --augmentations 7
```