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- 1 Introduction
- 2 The dataset
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We developed different models of neural networks to work with the given dataset from scratch and we did different test to find the better hyperparameters to build the final model. This section is divided in three parts: in the first one we describe the main preprocessing applied to the data before the training of the models. Then we show how we built a model for classifying the dataset images between mass abnormality and calfication abnormality. In the last part, we describe the model built to classify between benign and malignant abnormality.

4.1 Data Preprocessing

Starting from the numpy arrays of the dataset images we were given, first of all we deleted all the samples associated with the *baseline patch* label. Since this labeled images were placed in the even positions of the dataset, we just selected all the odd-index samples al discard the others. This is done both for the training data and the test data.

4.2 Task 2.1

For the first classification, we

- 4.3 Task 2.2
- 5 Task 3: Pretrained network
- 6 Task 4: Baseline patches
- 7 Task 5: Ensemble network