

# Configuration Manual

MSc Research Project  
Data Analytics

Mansi Atul Chowkkar  
Student ID: x18134599

School of Computing  
National College of Ireland

Supervisor:  
Cristina Muntean

**National College of Ireland**  
**MSc Project Submission Sheet**  
**School of Computing**



**Student Name:** Mansi Chowkkar  
**Student ID:** X18134599  
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**Lecturer:** Cristina Muntean  
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**Project Title:** Breast Cancer Detection from Histopathological Images using Deep Learning and Transfer Learning.  
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I hereby certify that the information contained in this (my submission) is information pertaining to research I conducted for this project. All information other than my own contribution will be fully referenced and listed in the relevant bibliography section at the rear of the project.

ALL internet material must be referenced in the bibliography section. Students are required to use the Referencing Standard specified in the report template. To use other author's written or electronic work is illegal (plagiarism) and may result in disciplinary action.

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# 1 Hardware/Software Requirements

## 1.1 Hardware Requirements

The below hardware would be ideal for running the experiment smoothly.

|                 |   |
|-----------------|---|
| OS              | Windows 10                              |
| RAM             | Minimum 8GB (2.14 GB from Colaboratory) |
| Hard Disk Space | Minimum 100GB (100 GB drive space)      |

## 1.2 Software Requirements

|                            |   |
|----------------------------|---|
| Programming Language Tools | Google Colaboratory (Cloud based Jupyter notebook environment), Python version 3, Microsoft Excel, Overleaf |
| Web Browser                | Google Chrome or Mozilla Firefox  |
| Email                      | Access to a Gmail account   |

# 1 Google Colaboratory Environment Setup

This section will explain how to setup Google Colaboratory environment in order to perform this experiment. Screenshots are included in order to facilitate a better understanding. A new Gmail address as a mansithesis.unet@gmail.com account is created in order to access Google Colaboratory.

1. Sign in with your Gmail account:

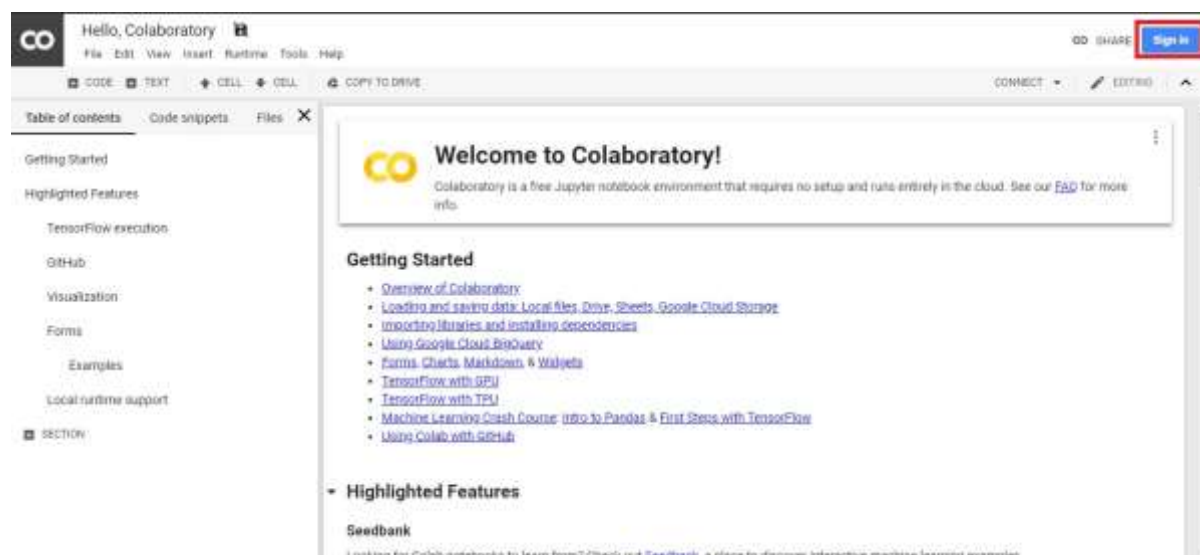


Figure 1: Sign in to Google Colaboratory

2. Once in the notebook, all required libraries are imported. The libraries required for each algorithm are written at the start of the coding of the algorithm.

## 2 Data preparation for Experiments

This section will explain how to upload data on Google Drive and access it in Collaboratory notebook.

1. Upload the image data folder on Google Drive:  
Histopathological image dataset of the Breast Biopsy is taken from the Kaggle as shown in the figure 2. Images of all magnifications are mixed in one folder for executing Experiment 1 and image data folder is uploaded on Google drive as shown in figure 3.

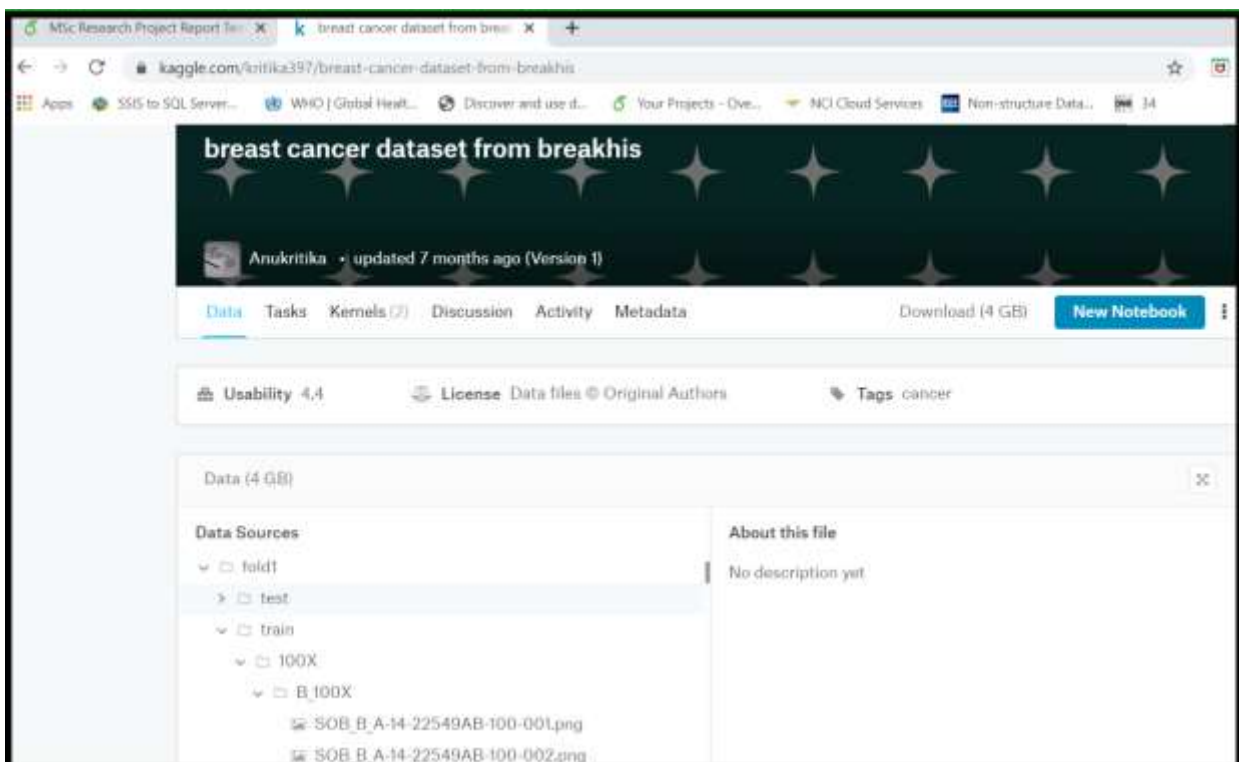


Figure 2:DataSet from Kaggle

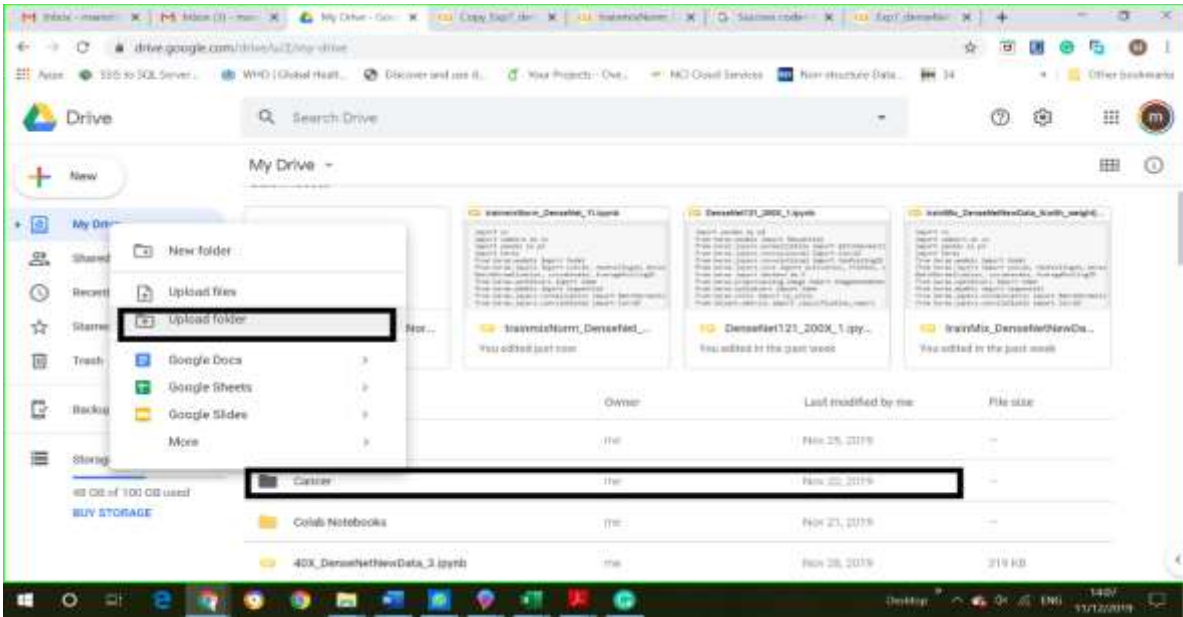


Figure 3: Upload data on Drive

2. Mount Google drive in the colab notebook:

Figure 4 shows the steps for mounting drive in the python code. Click on the url and select the gmail account for the colab and enter the authorization code as shown in figure 5.

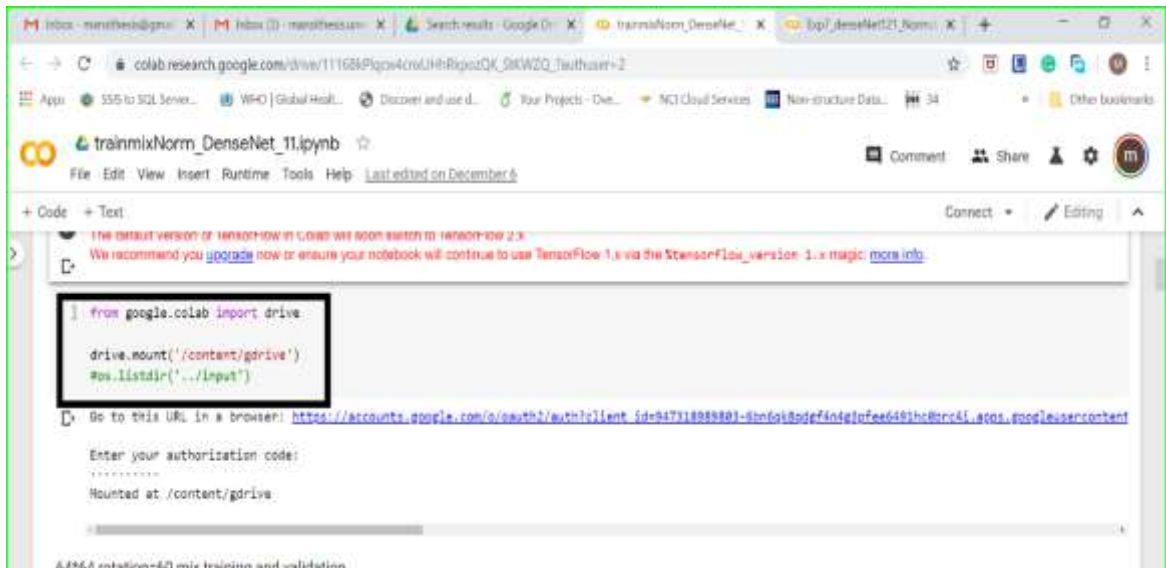


Figure 4: Mounting of Google Drive



Please copy this code, switch to your application and paste it there:

4/uAFkc0-  
hLcOszubPo241EWnPqencCwub4xdD33kz1UIbAMyNqBcsZXs





**Figure 5: Enter Authorization code**

3. Access the image data using below command in the notebook:

Figure 6 shows the code for accessing images from folder present on google drive.

```
[ ] train_gen=ImageDataGenerator(rotation_range=60,
                                width_shift_range=0.1,
                                height_shift_range=0.1,
                                rescale = 1./255,
                                #rescale = 0.8,
                                shear_range = 0.2,
                                zoom_range = 0.5,
                                horizontal_flip = True,
                                #featurewise_std_normalization=True,
                                fill_mode="nearest")

train = train_gen.flow_from_directory("/content/gdrive/My Drive/breast_cancer/fold1/train_mix_Norm/",
                                     class_mode='categorical',
                                     target_size=(64,64),
                                     color_mode="rgb",
                                     shuffle=True,
                                     batch_size=32)
```

**Figure 6: Accessing images in the python notebook**

4. Import Libraries for image normalization and augmentation:

Libraries required for Histogram normalisation are imported in python notebook as shown in figure7. ImageDataGenerator is used from Keras library to augment the images. Images are saved in the same directory for further use. Figure 7 shows the libraries required for the upsampling method.

```
import os
import cv2
import numpy as np
from PIL import Image
import pandas as pd
import matplotlib.pyplot as plt
from PIL import ImageOps
from numpy import expand_dims
from matplotlib import pyplot as plt
from IPython.display import display, Math, Latex
import matplotlib.pyplot as plt
```

Figure 7: Libraries for Histogram Normalization

```
import os
from numpy import expand_dims
from keras.preprocessing.image import load_img
from keras.preprocessing.image import img_to_array
from keras.preprocessing.image import ImageDataGenerator
```

Figure 8: Image Augmentation

### 3 Deep Learning and Transfer Learning Model Execution

This section will explain which libraries need to be imported for the execution of CNN and DenseNet-121

1. CNN Execution:

For CNN execution keras and tensorflow libraries are used from python. Figure 9 shows all the required libraries for image augmentation, model definitions, model compiling and model execution. Also, libraries are required for statistical analysis and graph plotting. The random seed is set every time so that every time different output is generated for different experiments.

- All layers in the CNN model are imported from keras.layers libraries.
- The model is built using tensorflow as tf library.
- The model layers are plotted using tensorflow.keras.utils library

```

import os
import keras
from keras.models import Model
from keras.layers import Conv2D, MaxPooling2D, Dense, Input, Activation, Dropout, GlobalAveragePooling2D, \
BatchNormalization, concatenate, AveragePooling2D
from keras.optimizers import Adam
from keras.models import Sequential
from keras.layers.core import Activation, Flatten, Dropout, Dense, Reshape
from keras import backend as K
from keras.preprocessing.image import ImageDataGenerator
from keras.utils import np_utils
from sklearn.metrics import confusion_matrix
from keras.preprocessing import image
from keras.preprocessing.image import img_to_array
from sklearn.preprocessing import MultiLabelBinarizer
import matplotlib.pyplot as plt
import numpy as np
import tensorflow as tf
from pathlib import Path
import cv2
from mlxtend.plotting import plot_confusion_matrix
tf.set_random_seed(1238)
from sklearn.metrics import f1_score

```

**Figure 9: Libraries for CNN execution**

## 2. DenseNet-121 Execution:

- For DenseNet-121 execution keras and tensorflow libraries are used from python. Figure 10 shows all the required libraries for image augmentation, model definitions, model compiling and model execution. Also, libraries are required for statistical analysis and graph plotting.
- DenseNet121 library is imported from keras package.
- Plotting of testing and training plots: matplotlib.pyplot as plt, seaborn as sns libraries are used for plotting training and validation graphs.





+ Code + Text

```
[6] import os
    from keras.layers.normalization import BatchNormalization
    from keras.layers.convolutional import Conv2D, MaxPooling2D
    from keras.layers.core import Activation, Flatten, Dropout, Dense, Reshape
    from keras import backend as K
    from keras.preprocessing.image import ImageDataGenerator
    from keras.optimizers import Adam
    from keras.utils import np_utils
    from keras.preprocessing import image
    from keras.preprocessing.image import img_to_array
    from sklearn.preprocessing import MultiLabelBinarizer
    import matplotlib.pyplot as plt
    import numpy as np
    import argparse
    import tensorflow as tf
    import cv2
    from mlxtend.plotting import plot_confusion_matrix
    from keras.applications import densenet
    from keras.models import Sequential, Model, load_model
    import seaborn as sn
    from sklearn.metrics import confusion_matrix, classification_report
    tf.set_random_seed(1234)
    from keras import layers
    from keras.applications import DenseNet121
```

Figure 10: Libraries for execution of DenseNet-121

## 4 Settings done for accelerating Computation time

This section will explain about how the drive storage is extended and GPU setting is done from the google colaboratory.

### 1. Drive Storage:

Extra 100 GB drive storage is subscribed to store normalized and augmented images. Extra storage was required for storing different python files with the executed models. Figure 11 shows the utilization of google drive for this project.

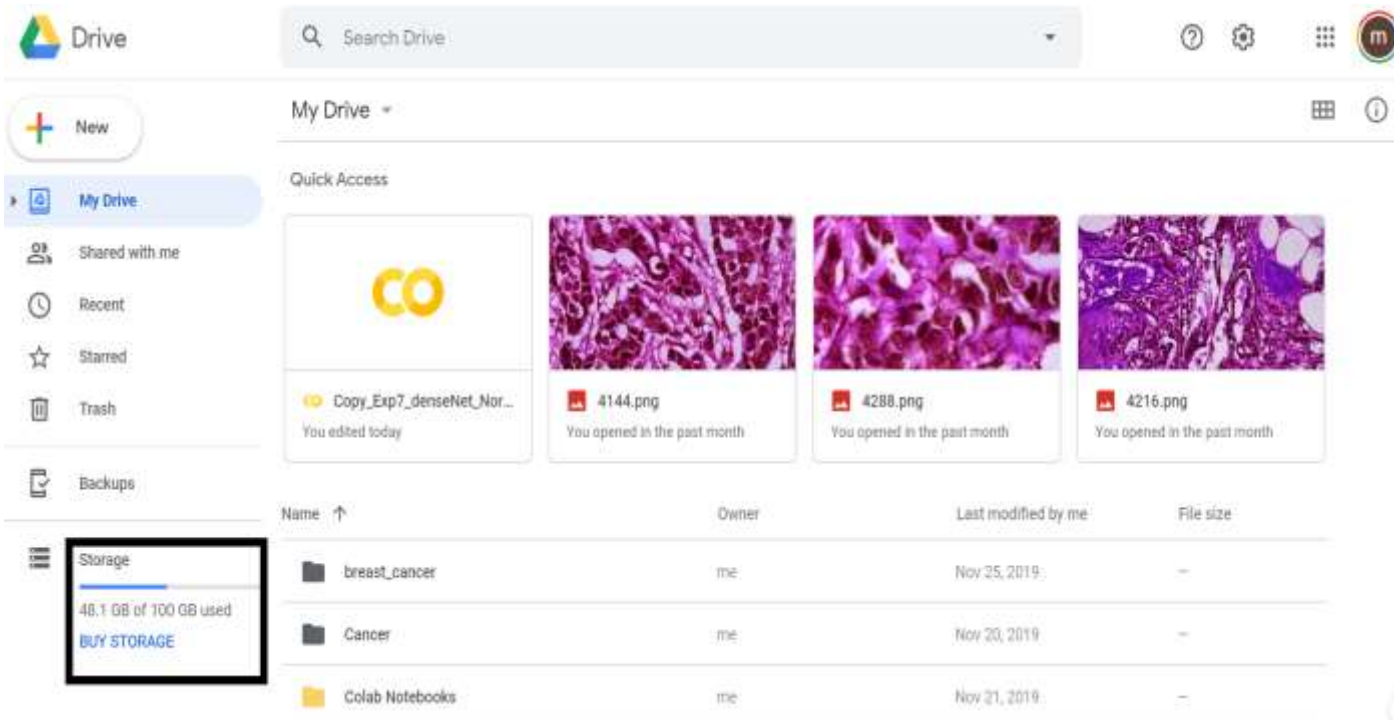


Figure 11: Storage

## 2. GPU:

For boosting the execution speed for both the models GPU accelerator hardware is selected from runtime option as shown in figure 12 and figure 13.

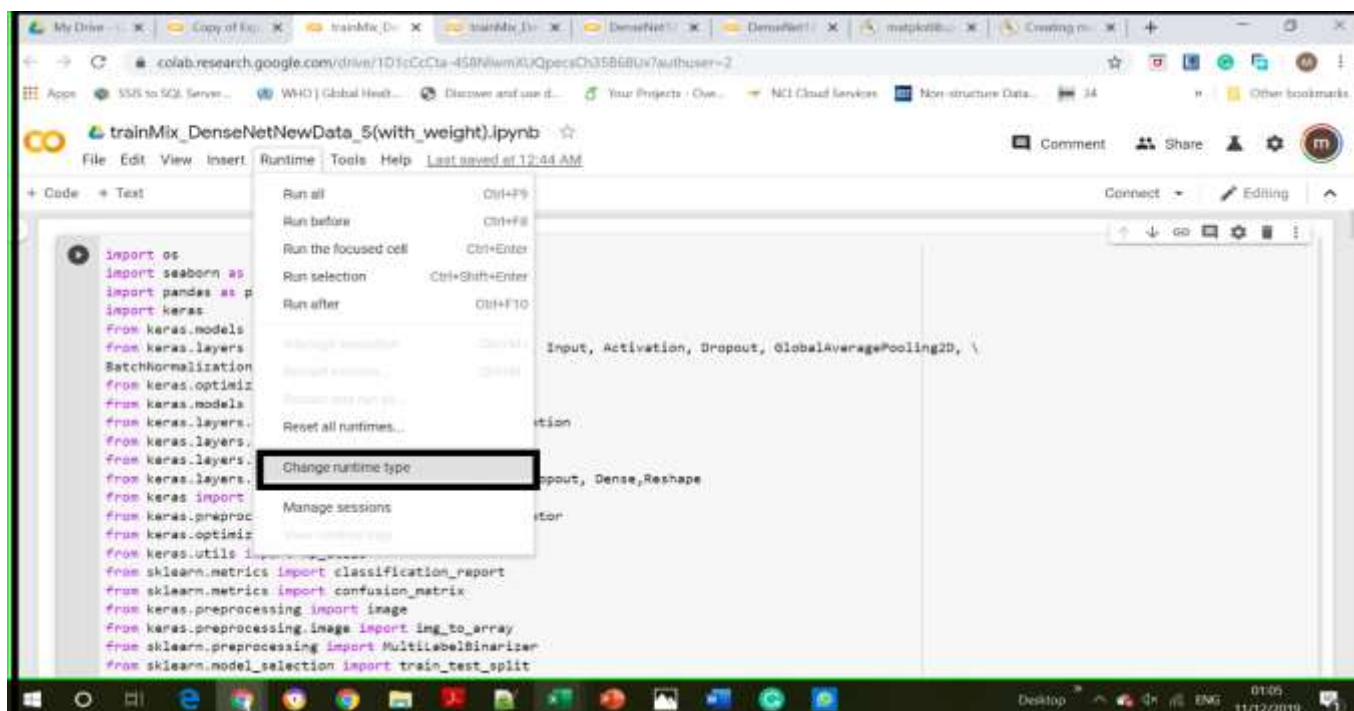


Figure 12: Runtime selection of hardware

## Notebook settings

Runtime type

Python 3

Hardware accelerator

GPU

Omit code cell output when saving this notebook

CANCEL

SAVE

Figure 13: GPU selection

## 5. Other Software used

For visualization of evaluation results Microsoft Excel is used. Research report is written using Overleaf tool. Figure 14 shows the utilization of Overleaf for this project.

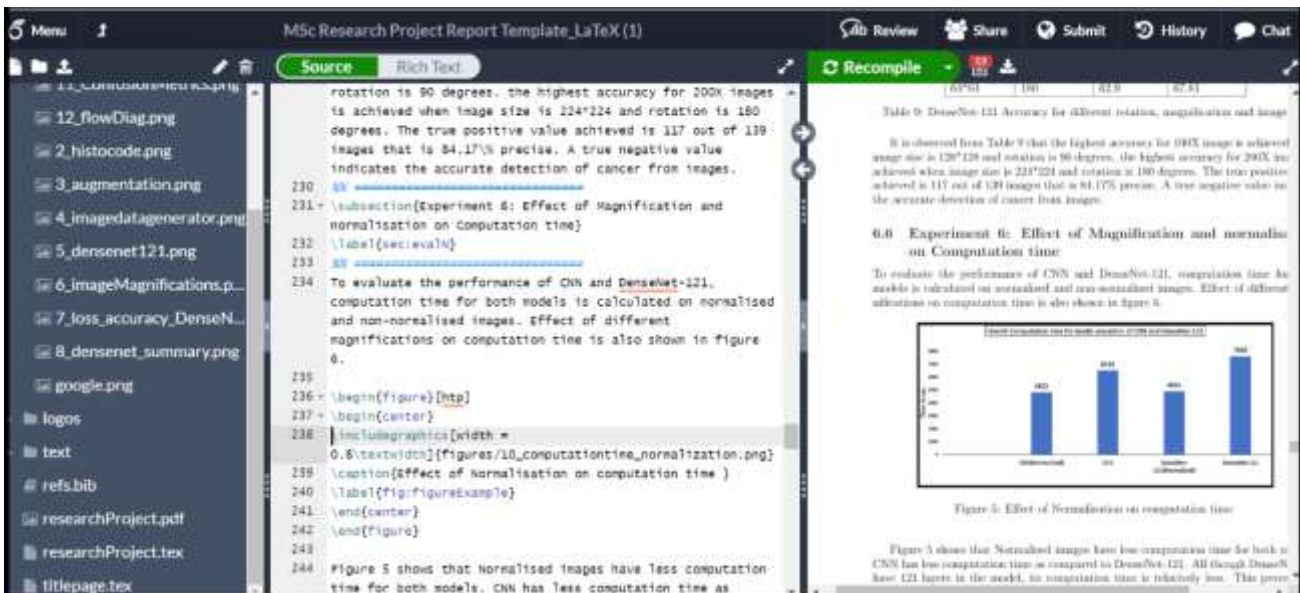


Figure 14: Overleaf Project

## References

<https://keras.io/preprocessing/image/>

<https://medium.com/hugoferreiras-blog/confusionmatrixandothermetricsinmachinelearning-894688cb1c0a>

<https://keras.io/getting-started/sequential-model-guide/>

<https://matplotlib.org/3.1.1/tutorials/introductory/pyplot.html>