

Configuration Manual

MSc Research Project Data Analytics

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Configuration Manual

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1 Overview

This document provides the step to step manual configuration for "PolyCystic Ovary Syndrome Detection Using CapsuleNet and Synthetic Data". This document will help to setup and install the prerequisites required to execute this research in future.

2 Hardware and software Requirement

2.1 Hardware Requirements

The research used the below-listed hardware in order to execute the code used in this research.

- Operating system used was Windows 10 Pro Language version 22H2.
- Processor: Intel(R) Core(TM) i5-8265U CPU @ 1.60GHz 1.80 GHz
- storage: RAM 8.00 GB
- System type: 64-bit operating system, x64-based processor

2.2 Software Requirements

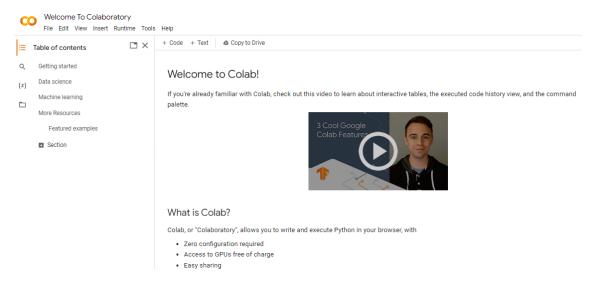
The below are software used to complete the research proposed

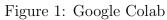
- Google Colab and Jupyter Notebook
- Python 3.7 Scripting Language
- Google Drive for Storage
- Notepad++. Word, Excel, Overleaf

3 Setting Environment

3.1 Google Colab

Firstly to start with the research project Google Colab was set up. In order to set up the Colab we need a Gmail account. The purpose to use Google Colab was as it provides free and high GPU for processing. Select the GPU before execution.



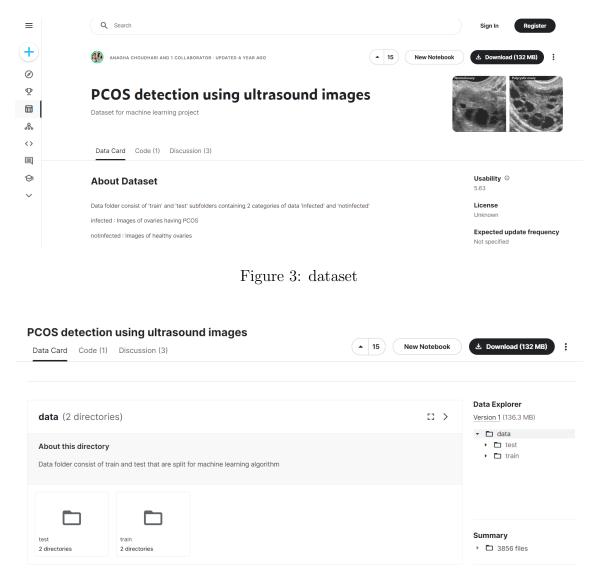


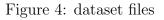
ardware accelerator 🥎 O CPU 💿 T4 GPU O A100 GPU O V100 GPU O TPU
Want access to premium GPUs? Purchase additional compute units

Figure 2: Selecting GPU

4 Data Selection

The data used in this research work was retrieved from the Kaggle data repository. The dataset consists of 3856 image files which were about 132 MB.





5 Data Storing and Model Building 5

5.1 Storing the data into Google Drive

The data fetched from google drive was uploaded to the drive in order to read and use the data easily while model building and compilation.

5.2 Prerequisite installation and library importing

This research used the below-mentioned libraries for model building, evaluation and data cleaning.

- numpy
- tensorflow.keras.utils
- PIL
- matplotlib
- torch
- random
- sklearn
- itertools
- keras
- google.colab
- ImageDataGenerator
- seaborn
- sklearn.metrics



Figure 5: Library Used

5.3 Connecting Google Drive and Colab

Google Colab was linked to Drive in order to retrieve and use the data as shown below 6. This piece of code was executed to mount the Google Colab and Drive

```
from google.colab import drive
drive.mount('/content/drive')
```

Mounted at /content/drive

Figure 6: Mounting Google Drive

5.4 Data loading, Pre-processing and splitting of data

5.4.1 Data Loading and reading

After mounting the drive data was read as shown below 7.



Figure 7: Data Reading

5.4.2 Data Pre-proceesing

After reading the data the data is in image format. It was mandatory to transfer the image in greyscale and should be of the same size before using it with the model. The image processing was done as shown below 8

```
image = Image.open(image_path).convert('RGB')
image = image.resize((128, 128))
image = np.array(image) / 255.0
X.append(image)
```

Figure 8: Pre-Processing

5.4.3 Setting the path for train and test

The path for train data and test data was set as shown below 9

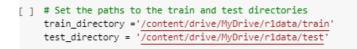


Figure 9: Setting path for Train and Test

6 Model building

6.0.1 Capsule Network

The Figure Below depicts a Capsule Network with 3 layers, primary and digital capsule layers with 256 convolution layers, 9x9 filters and routing agreement.

def	<pre>f CapsHet(input_shape, num_classes): model = tf.keras.Sequential()</pre>
	<pre># First layer: Convolutional layer with 9x9 filter and stride of 1 model.add(layers.Conv20(256, kernel_size=(9, 9), strides=(1, 1), activation='relu', input_shape=input_shape))</pre>
	# Second layer: Primary Capsule Network formed by 9x9 convolutions and stride of 2 model.add(layers.Conv2D(256, kernel_size=(9, 9), strides=(2, 2), activation='relu'))
	<pre># Third layer: Routing by Agreement process model.add(layers.Flatten()) model.add(layers.Gense(512, attivation='relu'))</pre>
	model.add[layers.Dense(256, activation='relu')]
	<pre># Last layer: Fully connected layer with softmax activation model.add(layers.Dense(num_classes, activation='softmax'))</pre>
	return model

Figure 10: Base Model Building

The model was compiled and used along with test data for evaluation purposes. 3

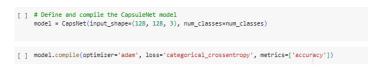


Figure 11: compile model

6.0.2 Data Augmentation

To generate synthetic data. Traditional Augmentation was implemented as shown below.

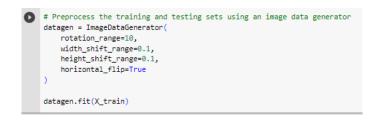


Figure 12: Data augmentation

6.0.3 Fine tune the model

Model tuning was performed in this research to enhance the results. The code and design of the model were defined as shown in fig 13

<pre>def CapsNetwork(input_shape, number_of_classes): tune_base_model = tf.keras.Sequential()</pre>
<pre># First layer: Convolutional layer with 9x9 filter and stride of 1 tune_base_model.add(layers.Conv2D(256, kernel_size=(9, 9), strides=(1, 1), activation='relu', input_shape=input_shape))</pre>
<pre># Second layer: Primary Capsule Network formed by 9x9 convolutions and stride of 2 tune_base_model.add(layers.Conv2D(256, kernel_size=(9, 9), strides=(2, 2), activation='relu'))</pre>
<pre># Third layer: Routing by Agreement process tume_bsse_model.add(layers.Flatten()) tume_bsse_model.add(layers.Demse(S12, activation='relu')) tume_bsse_model.add(layers.Demse(256, activation='relu'))</pre>
<pre># Last layer: Fully connected layer with softmax activation tune_base_model.add(layers.Dense(number_of_classes, activation='softmax'))</pre>
return tune_base_model
<pre># Load the pretrained model tune_base_model = CassHetwork(input_shape=(128, 128, 3), number_of_classes=number_of_classes)</pre>
<pre># Stage 1: Train only the last layer for layer in tune_base_model.layers[:-1]: layer.trainable = False</pre>
<pre>tune_base_model.compile(optimizer='adam', loss='categorical_crossentropy', metrics=['accuracy'])</pre>
Train the last layer on the rotated data tune_base_model.fit(X_train_rotated_image, y_train_rotated_image, batch_size=32, epochs=5, validation_data=(Xtest, ytest))
Stage 2: Train the entire model with a lower learning rate for layer in tune_base_model.layers: layer.trainable = True
<pre>tune_base_model.compile(optimizer=tf.keras.optimizers.Adam(1r=0.0001), loss='categorical_crossentropy', metrics=['accuracy'])</pre>

Figure 13: Tune Model

7 Evaluation

This research consists of different experiments to evaluate the model. The model is evaluated on the base of accuracy, precision, recall, sensitivity, and specificity and by comparing train and test accuracy. # predictions for the test set y_prediction = base_model.predict(Xtest) y_prediction_labels = np.argmax(y_prediction, axis=1) # Convert one-hot encoded test labels back to original class labels y_test_labels = np.argmax(ytest, axis=1) # calculate confusion matrix conf_matrix = confusion_matrix(y_test_labels, y_prediction_labels) # calculate specificity specificity_base_model = conf_matrix[0, 0] / (conf_matrix[0, 0] + conf_matrix[0, 1]) # calculate sensitivity (recall) sensitivity_base_model = recall_score(y_test_labels, y_prediction_labels, average='weighted') # calculate F1 score f1 = f1_score(y_test_labels, y_prediction_labels, average='weighted') print(f"Specificity: {specificity_base_model:.4f}") print(f"I score: {f1:.4f}")

Figure 14: Accuracy of Experiment 1



Figure 15: Confusion for Experiment 1

```
# Plot loss graph
plt.figure(figsize=(8, 6))
plt.plot(base_model_history.history['loss'], label='Training Loss')
plt.plot(base_model_history.history['val_loss'], label='Validation Loss')
plt.xlabel('Epoch')
plt.ylabel('Loss')
plt.legend()
plt.title('Training and Validation Loss')
plt.show()
# Plot accuracy graph
plt.figure(figsize=(8, 6))
plt.plot(base_model_history.history['accuracy'], label='Training Accuracy')
plt.plot(base_model_history.history['val_accuracy'], label='Validation Accuracy')
plt.xlabel('Epoch')
plt.ylabel('Accuracy')
plt.legend()
plt.title('Training and Validation Accuracy')
plt.show()
```



```
# Accuracy values for each dataset
accuracy_values = [accuracy_1_to_1, accuracy_1_to_2, accuracy_1_to_3]
# Dataset labels
dataset_labels = ['accuracy 1:1', 'accuracy 1:2', 'accuracy 1:3']
# Create line graph
plt.plot(dataset_labels, accuracy_values, marker='o', linestyle='-', color='b')
# Set axis labels and title
plt.xlabel('Dataset')
plt.ylabel('Accuracy (%)')
plt.title('Model Accuracy on Different Datasets')
# Show gridlines
plt.grid(True)
# Show the line graph
```

plt.show()

Figure 17: Experiment 2 Graph

<pre># Assuming you have already trained the model and have predictions for the test set y_predictions_synthetic_data1 = tune_base_model.predict(X_test_rotated_image) y_pred_labels_synthetic_data1 = np.argmax(y_predictions_synthetic_data1, axis=1)</pre>
<pre># Convert one-hot encoded test labels back to original class labels y_test_labels_synthetic1 = np.argmax(y_test_rotated_image, axis=1)</pre>
<pre># Calculate confusion matrix conf_matrix = confusion_matrix(y_test_labels_synthetic1, y_pred_labels_synthetic_data1)</pre>
<pre># Calculate specificity specificity = conf_matrix[0, 0] / (conf_matrix[0, 0] + conf_matrix[0, 1])</pre>
<pre># Calculate sensitivity (recall) sensitivity = recall_score(y_test_labels_synthetic1, y_pred_labels_synthetic_data1, average='weighted'</pre>
<pre># Calculate F1 score f1 = f1_score(y_test_labels_synthetic1, y_pred_labels_synthetic_data1, average='weighted')</pre>
<pre>print(f"Specificity_synthetic_data: {specificity:.4f}") print(f"Sensitivity (Recall)_synthetic_data: {sensitivity:.4f}") print(f"F1 Score_synthetic_data: {f1:.4f}")</pre>
9/9 [=======] - 1s 109ms/step Specificity_synthetic_data: 1.0000 Sensitivity (Recall)_synthetic_data: 0.9674 F1 Score_synthetic_data: 0.9675

Figure 18: Accuracy of Experiment 3

import numpy as np	
import pandas as pd	
import seaborn as sns	
import matplotlib.pyplot as plt	
from sklearn.metrics import confusion_matrix, accuracy_score, precision_score, r	ecall_score, f1_score
# Print confusion matrix as a heatmap	
class_names = ['infected', 'non-infected']	
<pre>conf_df = pd.DataFrame(conf_matrix, index=class_names, columns=class_names)</pre>	
<pre>plt.figure(figsize=(10, 8))</pre>	
<pre>sns.heatmap(conf_df, annot=True, fmt="d", cmap='Blues')</pre>	
<pre>plt.xlabel('Predicted Labels')</pre>	
plt.ylabel('True Labels')	
plt.title('Confusion Matrix original data')	
plt.show()	

Figure 19: Confusion for Experiment 3

```
# Plotting the graphs
import matplotlib.pyplot as plt
# Plot loss graph
plt.figure(figsize=(8, 6))
plt.plot(original_data_tune_model.history['loss'], label='Training Loss')
plt.plot(original_data_tune_model.history['val_loss'], label='Validation Loss')
plt.xlabel('Loss')
plt.title('Training and Validation Loss')
plt.show()
# Plot accuracy graph
plt.figure(figsize=(8, 6))
plt.plot(original_data_tune_model.history['accuracy'], label='Training Accuracy')
plt.ylabel('Epoch')
plt.ylabel('Accuracy')
plt.ylabel('Accuracy')
plt.legend()
plt.title('Training and Validation Accuracy')
plt.show()
```

Figure 20: Graph for Experiment 3