

Configuration Manual

MSc Research Project Data Analytics

Rohit Salvi Student ID: x21208832

School of Computing National College of Ireland

Supervisor:

Paul Stynes

National College of Ireland Project Submission Sheet School of Computing



Student Name:	Rohit Salvi
Student ID:	x21208832
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Configuration Manual

Rohit Salvi x21208832

1 Overview

The configuration manual is the step-by-step guide for setting up the environment, prerequisites and execution of the research project "Lightweight Deep Learning Framework for Brain Tumour Classification".

2 Hardware/Software Requirements

2.1 Hardware Requirements

The research project was executed on the system with the following hardware configurations.

- Operating System: Windows 10 Home Single Language(Version: 22H2).
- Processor: Intel(R) Core(TM) i5-8250U CPU @ 1.60GHz 1.80 GHz.
- Storage: 2TB.
- **RAM:** 8GB(Extendable 20.4GB Virtual Memory).
- Graphical Processing Unit: NVIDIA GeForce MX150.

2.2 Software Requirements

The software that aided the designing, implementation and execution of the research project are as follows.

- **Development Environment:** Jupyter Notebook(Version: 6.4.12).
- **Programming Language:** Python(Version: 3.10.7).
- Tools: Lucidchart and Overleaf.

3 Set Up

3.1 Python

3.1.1 Installing Python

Following are the steps to install Python in the system.

- 1. Go to offical website of Python¹ and download python source code and installer with version 3.10.7.
- 2. On the python source code and installer with version 3.10.7 is downloaded. Install Python by running the installer as shown in Figure 1.



Figure 1: Installing Python

3.1.2 Starting and Verifying Python

The following steps will help to verify the availability of Python and Python version in the system.

- 1. Open the command prompt by searching "Command Prompt" in the Windows search bar.
- 2. Once the command prompt is opened type python and execute it. The version of Python is also displayed on execution of it as shown in Figure 2



Figure 2: Python and its Version

¹https://www.python.org/

3.2 Jupyter Notebook

3.2.1 Installing and Running Jupyter Notebook

Following are the steps to install and run Jupyter Notebook in the system.

1. Go to offical website of Jupyter². Scroll down to the section of Jupyter Notebook and click on "Install the Notebook" as shown in Figure 3.

○ Project Jupyter Home ×	+	 ✓ - □ × ★ □ ♠ Inconto : 	
Uppter Weckense by Uppter Weckense by Uppter Weckense by Uppter Upp		Jupyter Notebook: The Classic Notebook Interface The Jupyter Notebook is the original web application for creating and sharing computational documents. It offers a simple, streamlined, document-centric experience. Try it in your browser	
		Space C	¥

Figure 3: Install Jupyter Notebook

2. After clicking on "Install the Notebook", several ways to install Jupyter will be displayed. Open the command prompt and execute the command as shown in the Figure 4 to install Jupyter Notebook.

Jupyter Notebook

Install the classic Jupyter Notebook with:

pip install notebo	k		
To run the noteboo	<:		
jupyter notebook			

Figure 4: Command to Install and Running Jupyter Notebook

3. On executing the commands mentioned in the above step, a web interface will be popped up in the default browser. Click on "New" and then "Python3" to open the new Jupyter Notebook as shown in Figure 5. Then try executing the simple Python command to ensure the integration and execution of Python in Jupyter Notebook as shown in Figure 6.

²https://jupyter.org/

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$\leftrightarrow \rightarrow$ C Δ (3) localhost.8889/tree			
💭 jupyter		Ouit Logout	
Files Running Clusters Nbextensi	ons		
Select items to perform actions on them.		Upload New - 2	
0 - 1		Name Python 3 (pykernel)	
	The notebook list is empty.	ĸ	
		Text File	
		Folder	
		Terminal	

Figure 5: Opening New Jupyter Notebook

Cjupyter Research Project Last Checkpoint: 2 minutes ago (uns	weed changes)		🐣 Logost
File Edit View Insert Cell Kernel Navigate Widge	ds Help	Trusted	Python 3 (ipykernel) O
S + 3< 2 E + ₽ Run ■ C + Code			
	b: [1]: print("wills words") manager data (way word prints) and wills words" b: [1]:		

Figure 6: Execution of Python Code in Jupyter Notebook

3.3 Data

3.3.1 Data Selection

The data used in the research project is accessible on Kaggle(Nickparvar; 2021). The dataset consists of 7023 '.*jpg*' images of MRIs of 4 different classes of brain tumours.

3.3.2 Importing Data

To import the dataset into the system, navigate to the Brain Tumour MRI Dataset³ on Kaggle. Hit the download button as shown in Figure 7. Post that select the repository created for the implementation of the research project in the system. Click on download, then download will begin and data will be imported into the system as '.*zip*' file. Extract the data from '.*zip*' file.



Figure 7: Importing Data from Kaggle

³https://www.kaggle.com/datasets/masoudnickparvar/brain-tumor-mri-dataset

3.4 Libraries

Following is the list of the packages and libraries required for pre-processing of data, implementation of the model and evaluation of it.

- PyTorch
- Torchvision
- NumPy
- Collections
- Matplotlib
- Pillow
- Scikit-learn
- Seaborn
- Scikit-optimize

3.4.1 Installing Libraries

Install the necessary libraries required for the execution of the research project using the following command.

```
pip install torch
pip install torchvision
pip install matplotlib
pip install Pillow
pip install scikit-learn
pip install scikit-optimize
```

3.4.2 Importing Libraries

Once the required libraries are installed, those libraries are imported, as shown in the Figure 8, for implementation and execution of the research framework.

4 Project Development

4.1 Seed SetUp

For the purpose of reproducibility of research results, set the seed of the environment as shown in the Figure 9.

4.2 Directories SetUp

Directories in which the data is present are set as shown in the Figure 10. Note: The research data and Jupyter Notebook file are in the same directory.

```
import os
import torch
import torchvision.transforms as transforms
from torchvision.datasets import ImageFolder
from torch.utils.data import DataLoader
import numpy as np
from collections import Counter
from torch.utils.data import Subset
executed in 6.62s, finished 16.37.49 2023-08-13
```

For Data Visualization

import matplotlib.pyplot as plt from PIL import Image executed in 1.51s, finished 16:37:50 2023-08-13

For Model

import torch.nn as nn import torch.optim as optim executed in 11ms, finished 16:37:50 2023-08-13

For Evaluation

from sklearn.metrics import accuracy_score, confusion_matrix
import seaborn as sns
executed in 5.34s, finished 16.37:55 2023-08-13

For Optimization

from skopt.space import Real
from skopt import gp_minimize
executed in 597ms, finished 16:37:56 2023-08-13

For Results

from sklearn.metrics import classification_report
executed in 12ms, finished 16:37:56 2023-08-13

For K-Fold Validation

from sklearn.model_selection import KFold
executed in 11ms, finished 16:37:56 2023-08-13

Figure 8: Importing Libraries

	seed = 12
	torch.manual_seed(seed)
	torch.cuda.manual_seed(seed)
	torch.backends.cudnn.deterministic = True
	<pre>torch.backends.cudnn.benchmark = False</pre>
	np.random.seed(seed)
1	xxecuted in 29ms, finished 10:14:54 2023-08-13

Figure 9: Setting Seed



Figure 10: Directories SetUp

4.3 Data Loading and Pre-Processing

After setting up the data directories, the data is loaded and pre-processed. The snippet of code for data loading and pr-processing can be seen in the Figure 11.

Data augmentations of Training data

```
trainDataTransform = transforms.Compose([
    transforms.RandomHesizedCrop(imageSize),
    transforms.RandomHorizontalFlip(),
    transforms.RandomVerticalFlip(),
    transforms.ToTensor(),
    transforms.Normalize(mean = [0.485, 0.456, 0.406], std = [0.229, 0.224, 0.225])
])
executed in 13ms, finished 10:14:54 2023-08-13
Normalizing Testing Data
```

```
v testDataTransform = transforms.Compose([
    transforms.Resize(imageSize),
    transforms.ToTensor(),
    transforms.Normalize(mean = [0.485, 0.456, 0.406], std = [0.229, 0.224, 0.225])
])
executed in 14ms. finished 10.14.54 2023-08-13
```

Loading Data

```
trainData = ImageFolder(trainingDirectory, transform=trainDataTransform)
testData = ImageFolder(testingDirectory, transform=testDataTransform)
executed in 3.72s. finished 10:14:58 2023-08-13
```

Figure 11: Code for Data Loading and Pre-Processing

4.4 CNN Model

The CNN model of the research project has similar architecture of the model by Soewu et al. (2022). The architecture and code for the implementation of CNN can be seen in the Figure 12.



```
model = CNNModel(num_classes=len(categories))
model.to(device)
executed in 75ms, finished 10:14:59 2023-08-13
CNNModel(
  (conv1): Conv2d(3, 16, kernel_size=(3, 3), stride=(1, 1), padding=(1, 1))
  (pool): MaxPool2d(kernel_size=2, stride=2, padding=0, dilation=1, ceil_mode=False)
  (conv2): Conv2d(16, 32, kernel_size=(3, 3), stride=(1, 1), padding=(1, 1))
  (conv3): Conv2d(32, 16, kernel_size=(3, 3), stride=(1, 1), padding=(1, 1))
  (fc1): Linear(in_features=16384, out_features=256, bias=True)
  (fc2): Linear(in_features=256, out_features=4, bias=True)
  )
```

Figure 12: CNN Model

CNN model is trained with the training data as shown in the Figure 13.



Figure 13: Training CNN

4.5 Pruning CNN Model

CNN Model is pruned using a magnitude-based weight training technique. The function for pruning the CNN model can be seen in the Figure 14.

Funct	ion to prune CNN Model
def	<pre>pruneHodel(model, pruningRatio): allParameters = [] for paramName, param in model.named_parameters(): if 'weight' in paramName: # OnLy prume weights, not biases allParameters.append((paramName, param.data.view(-1))) # Flatten and concatenate all weights allWeights = torch.cat[[param for _, param in allParameters])</pre>
	# Calculate the threshold value for pruming numBransToPrume = int(prumingBatio * len(allWeights)) threshold = torch.topk(torch.abs(allWeights), numParamsToPrune).values.min()
,	<pre># Apply pruning mask to each parameter tensor for paramWame, param in allParameters: mask = torch.abs(param) > threshold param.data *= mask.float() print(f*Pruning (pruningRatio*100:.2f)% of model parameters.")</pre>
	return model
execute	a m 20116, imilanda 12, 10, 20, 20, 20, 10
def	<pre>getPrunedModel(originalModel, pruningRatio): # Create a new model instance prunedModel instance prunedModel - CNMModel(num_classes=len(categories)) prunedModel.to(device)</pre>
	<pre># Load the state_dict from the original model to the new model prunedModel.load_state_dict(originalModel.state_dict())</pre>
	# Apply pruning to the new model prunedModel = pruneModel(prunedModel, pruningRatio)
	return prunedModel
execute	d in 26ms. finished 12:10:23 2023-08-13

Figure 14: Pruning Function

Later the pruned model is re-trained on the training data.

4.6 Optimal Pruning Ratio

The optimal pruning ratio is identified using the objective function as shown in Figure 15.



Figure 15: Objective Function for identifying Optimal Pruning Ratio

Then lightweight deep learning framework is built using the CNN model, pruning functional and optimal pruning ratio.

5 Project Testing

The developed framework is validated with the help of testing data. The code for the testing of the framework is in the Figure 16.

<pre>optimisedPrunedModel.eval() correct = 0 total = 0 predictedLabelsOfOptimisedPrunedModel = [] trueLabelsOfOptimisedPrunedModel = [] with torch.org_rad(): for inputs, labels = inputs.to(device), labels.to(device) outputs = optimisedPrunedModel(inputs) , predicted = torch.max(outputs.data, 1) total + labels.size(0) correct += (predicted == labels).sum().item()</pre>	
# Save predicted and true labels for calculating metrics predicted.abels0F0ptimisedPrunedWodel.extend(predicted.cpu().numpy()) truelabels0F0ptimisedPrunedWodel.extend(labels.cpu().numpy())	
ecuted in 38.75, finished 01:43:01 2023-08-11	

Figure 16: Validating the Framework

To validate the robustness of the model, k-fold cross-validation technique is applied as shown in the Figure 17.



Figure 17: K-Fold Cross Validation

The performance metrics of the framework are calculated as shown in the Figure 18, 19, 20 and 21

accuracyOfOptimisedPrunedModel = accuracy_score(trueLabelsOfOptimisedPrunedModel, predictedLabelsOfOptimisedPrunedModel)
print(f"Accuracy of Optimised Pruned CNN: {accuracyOfOptimisedPrunedModel*100}")
executed in 30ms, finished 01:43:01:2023-08-11

Figure 18: Computing Accuracy

confusionMatrixOfOptimisedPrunedModel = confusion_matrix(trueLabelsOfOptimisedPrunedModel, predictedLabelsOfOptimisedPrunedModel, predictedLabels
executed in 39ms, finished 01:43:01 2023-08-11
<pre>plt.figure(figsize = (8, 6))</pre>
sns.heatmap(confusionMatrixOfOptimisedPrunedModel, annot = True, fmt = "d", cmap = "Blues", xticklabels = categories, yticklab
plt.xlabel('Predicted')
plt.ylabel('True')
plt.title('Confusion Matrix')
plt.show()
executed in 2.52s, finished 01:43:04 2023-08-11

Figure 19: Computing Confusion Matrix



Figure 20: Computing Sensitivity and Specificity

```
plt.plot(range(1, epochs + 1), lossValuesOfOptimisedPrunedModel, label = 'Loss')
plt.plot(range(1, epochs + 1), accuracyValuesOfOptimisedPrunedModel, label = 'Accuracy')
plt.xlabel('Epoch')
plt.ylabel('Value')
plt.title('Loss and Accuracy per Epoch')
plt.legend()
plt.show()
executed in 256ms, finished 01.43.04 2023-06-11
```

Figure 21: Plotting Loss and Accuracy per Epoch

References

Nickparvar, M. (2021). Brain tumor mri dataset. URL: https://www.kaggle.com/datasets/masoudnickparvar/brain-tumor-mri-dataset

Soewu, T., Singh, D., Rakhra, M., Chakraborty, G. S. and Singh, A. (2022). Convolutional neural networks for mri-based brain tumor classification, 2022 3rd International Conference on Computation, Automation and Knowledge Management (ICCAKM), IEEE, pp. 1–7.