



Computer Science Department

2021/2022

CS 396 Selected Topics in CS-2 Research Project

Report Submitted for Fulfillment of the Requirements and ILO's for Selected Topics in CS-2 course for Fall 2021

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Paper Details

- Paper Name: Multiclass classification of brain tumours using a novel CNN architecture.
- Authors: Hareem Kibriya & Momina Masood & Marriam Nawaz & Tahira Nazir.

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Project Description

Brain tumors are a deadly condition that radiologists have a tough time diagnosing. It is critical to make treatment-related decisions based on accurate and timely categorization of malignant cancers. Several approaches for detecting brain tumors have been presented in recent years. These strategies, however, necessitate handmade feature extraction and manual tumor segmentation prior to classification, which is error-prone and time-consuming.

Datasets for it: Brain Tumor MRI Dataset

It includes 8222 files for (Training & Testing)

Training (4 directories)

- Glioma
- Meningioma
- pituitary
- no tumor

Testing (4 directories)

- Glioma
- Meningioma
- pituitary
- no tumor

dimension of images = 512*512px

• Implementation details

According to the paper,

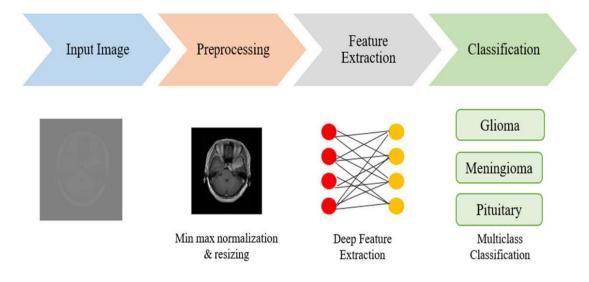
First we removed "No tumor" file from training and testing

Now number of files is: 7787

Then we split training data to (training and validation) with 30%

Now we have:

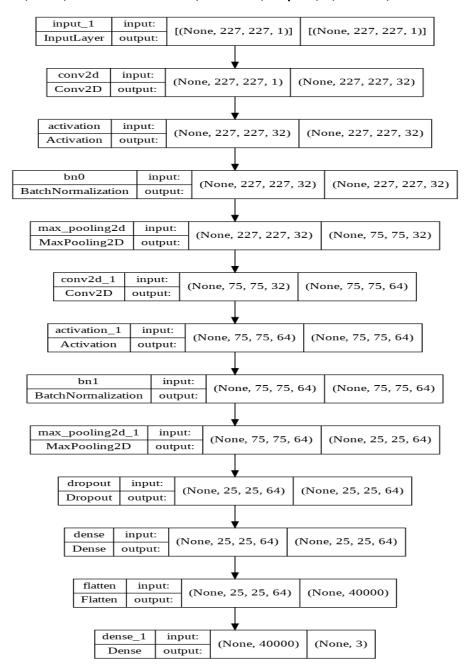
- Training (4614)
- Validation(1978)
- Testing (1195)



Our Model

13 layers

(input,2Conv,2Relu,2BatchNormalization,2MaxPool,Dropout,Fc,Softmax,Classification)

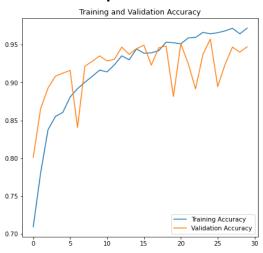


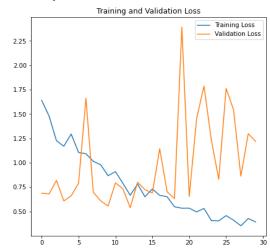
Hyperparameters and architectures tested before reaching the final model

Factor (s)	Value (s)
Convolutional Layer+ReLU	1, 2 , 3
Batch Normalization Layer	1, 2 , 3
Dropout Layer	1, 2
Cross channel normalization layer	0 , 1, 2
Fully Connected Layer	1, 2
Grouped Convolution Layers	0 ,1, 2
Pooling Layers	Max, Average
Dropout Rate	0.25, 0.5
Learning Rates	0.0001, 0.0002, 0.0003 , 0.0005
Optimizers	Adam, SGDM
No. of Epochs	10, 20, 30 , 40
Mini batch size	4, 8, 10
No. of convolution kernels	32, 64 , 128, 256
Convolution kernel size	2, 3, 5
Image Size	128×128 , 227 × 227

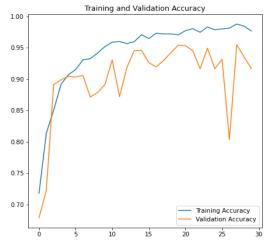
• Testing results

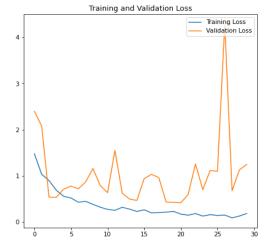
When we keep Lr default and use Adam Optimizer:



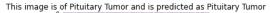


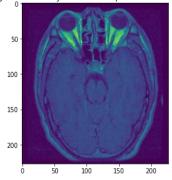
When we change the learning rate to 0.0003 with adam Optimizer:



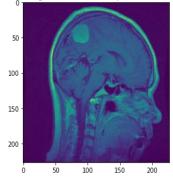


Outputs:

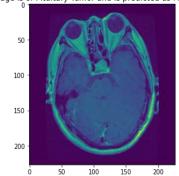




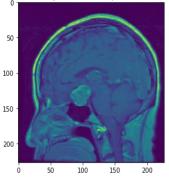
This image is of Meningioma Tumor and is predicted as Meningioma Tumor



This image is of Pituitary Tumor and is predicted as Pituitary Tumor



This image is of Pituitary Tumor and is predicted as Pituitary Tumor



Accuracy: 96 %

Recall, Percision, f1-Score:

```
[16] pred = model.predict(test_X)
     pred = np.argmax(pred,axis=1)
     print(classification_report(test_y,pred))
                                recall f1-score
                   precision
                                                   support
                        0.93
                                  0.77
                                            0.84
                                                       400
                        0.79
                                  0.94
                                            0.86
                        0.96
                                  0.93
                                            0.94
                                                       374
         accuracy
                                            0.88
        macro avg
                                            0.88
                        0.89
                                  0.88
     weighted avg
                        0.89
                                  0.88
                                            0.88
```

Confusion matrix:

