

Brain Tumor Detection From MRI Images With IBM Watson Studio

1. INTRODUCTION

1.1 Overview

Brain tumor identification is a really challenging task in the early stages of life. These days the issue of brain tumor automatic identification is of great interest. A tumor is the unusual growth of the tissues. A brain tumor is a number of unnecessary cells growing in the brain or central spinal canal. It is the unrestrained progress of cancer cells in any portion of the body.

Deep learning techniques can be used in order to detect the brain tumor of a patient using the MRI images of a patient's brain. In this application, we are helping the doctors and patients to classify the type of scan for the specific image given with the help of Neural Networks and store the patient's data.

1.2 Purpose

Brain tumor detection from MRI images is one of the most challenging task in today's modern Medical imaging research. Brain tumor detection really challenging task in early stages of life. These days issue of brain tumor automatic identification is of great interest. Tumor is the unusual growth of the tissues.

A brain tumor is a quantity of unnecessary cells growing in the brain or central spine canal. It is an unrestrained progress of cancer cells in any portion of the body. In order to detect the brain tumor of a patient, MRI images of a patient's brain are considered. This application helps you detect the tumor and store the patient's data.

2. LITERATURE SURVEY

2.1 Existing problem

In this study, using MR images of the brain, we segmented brain tissues into normal tissues such as white matter, gray matter, cerebrospinal fluid (background), and tumor-infected tissues.

To classify the tumor stage by analyzing feature vectors and area of the tumor. In this study, we investigated texture based and histogram based features with a commonly recognized classifier for the classification of brain tumor from MR brain images. From the experimental results performed on the different images, it is clear that the analysis for the brain tumor detection is fast and accurate when compared with the manual detection performed by radiologists or clinical experts.

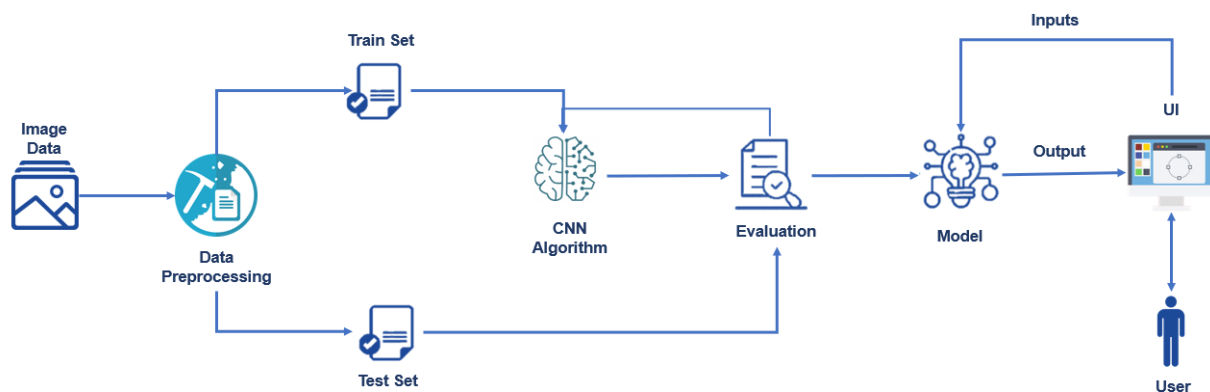
2.2 Proposed solution

Tumor is the unusual growth of the tissues. A brain tumor is a quantity of unnecessary cells growing in the brain or central spine canal. It is an unrestrained progress of cancer cells in any portion of the body.

In order to detect the brain tumor of a patient, MRI images of a patient's brain are considered. This application helps you detect the tumor and store the patient's data. The result of the analysis is carried out whether the brain is normal one or diseased one by applying deep learning techniques.

3. THEORITICAL ANALYSIS

3.1 Block Diagram



3.2 Hardware / Software designing

Software Requirements:

- Anaconda Navigator
- Keras
- Flask

Hardware Requirements:

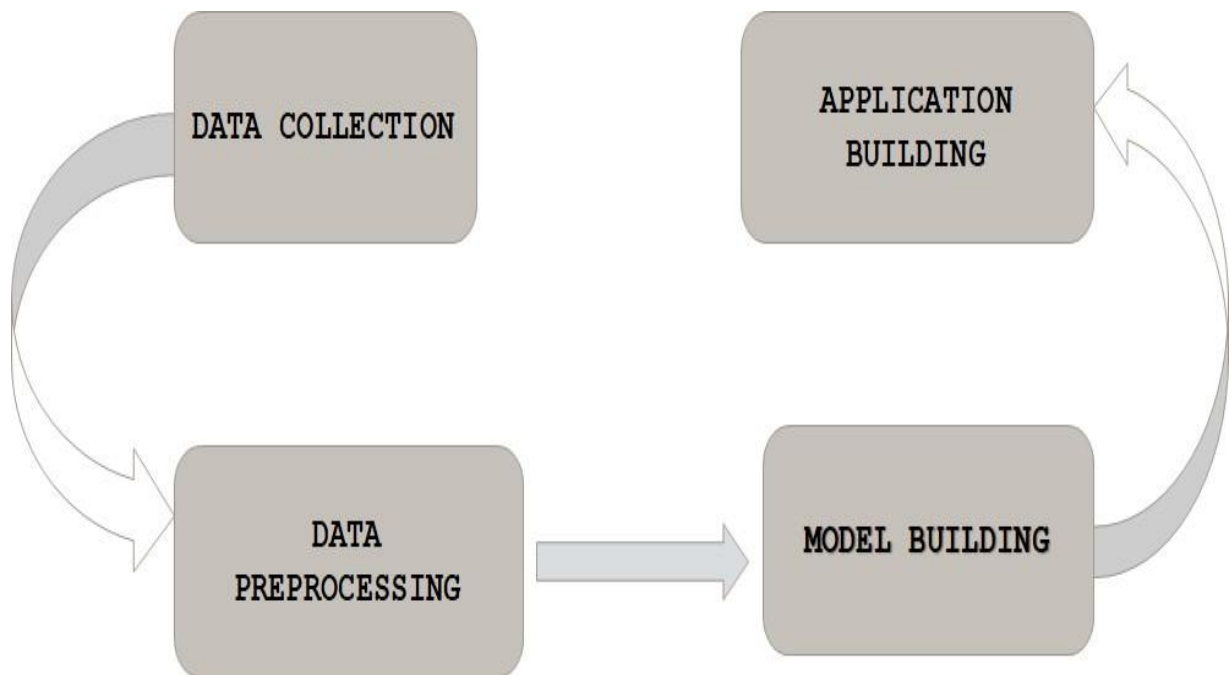
- Processor : Intel Core i3
- Hard Disk Space : Min 100 GB
- Ram : 4 GB
- Display : 14.1 “Color Monitor(LCD, CRT or LED)
- Clock Speed : 1.67 GHz

4. EXPERIMENTAL INVESTIGATIONS

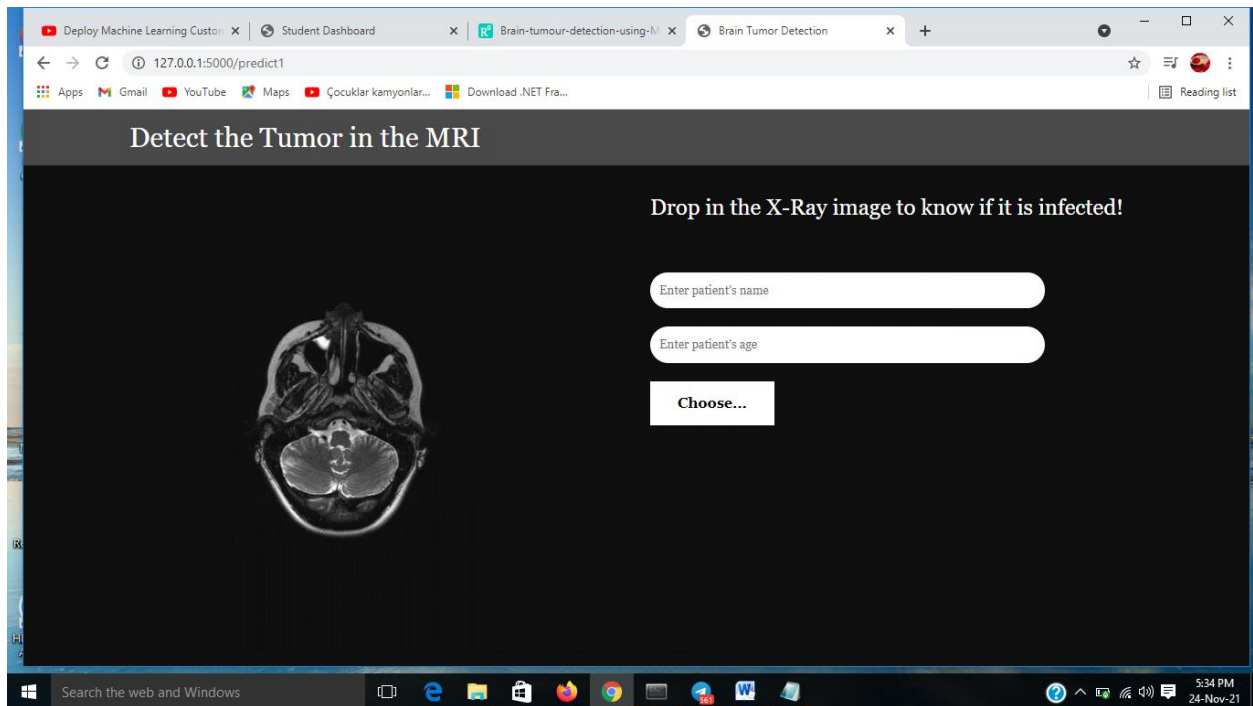
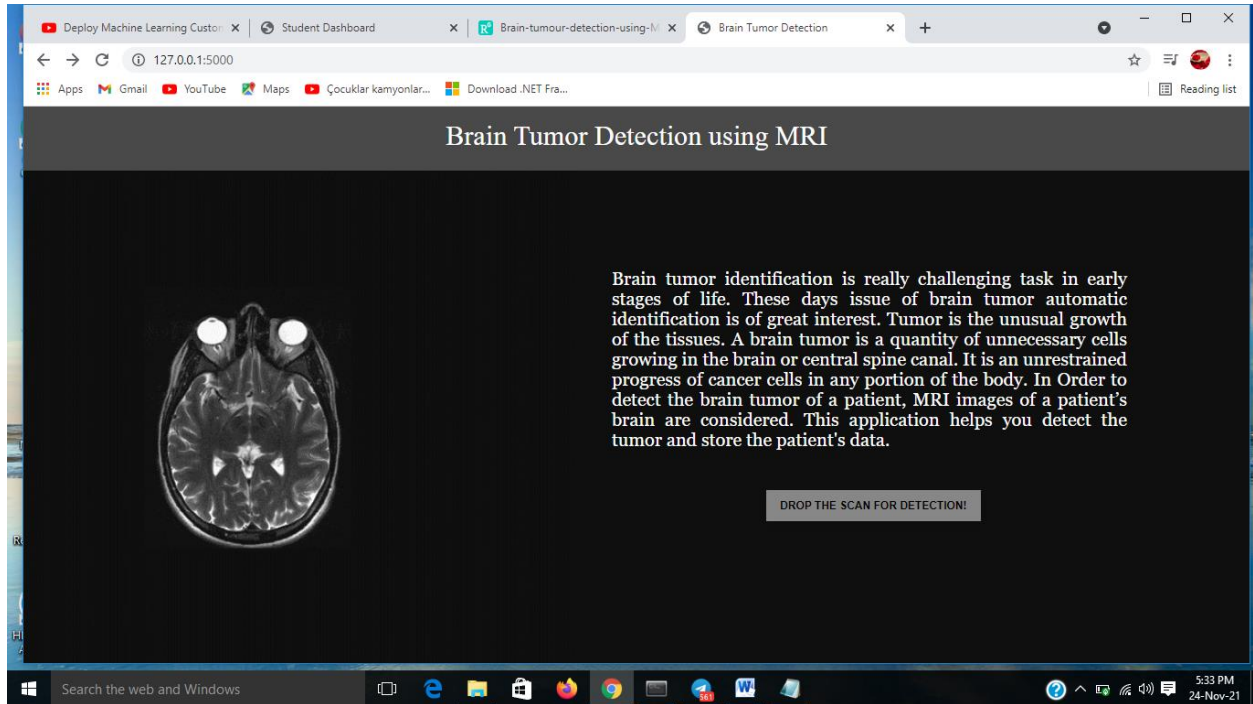
Our experimental results show that the proposed approach can aid in the accurate and timely detection of brain tumor along with the identification of its exact location. Thus, the proposed approach is significant for brain tumor detection from MR images.

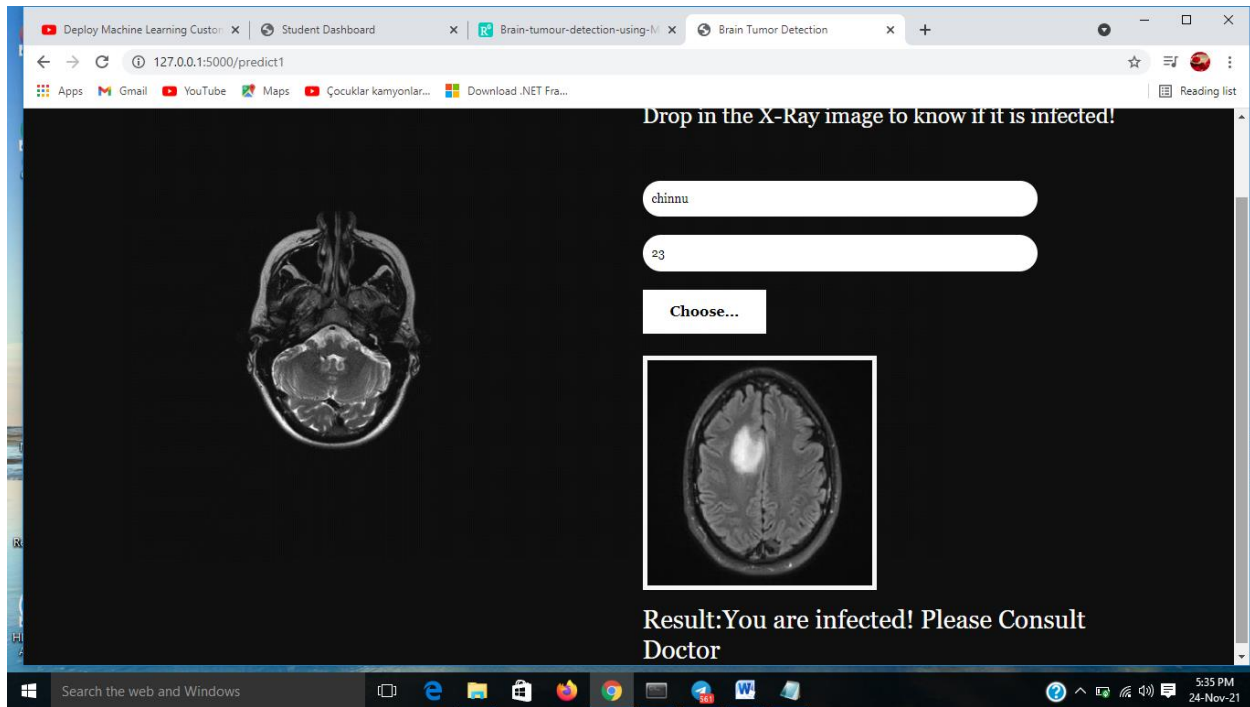
The experimental results achieved 96.51% accuracy demonstrating the effectiveness of the proposed technique for identifying normal and abnormal tissues from MR images. The result of the analysis is carried out whether the brain is normal one or diseased one by applying deep learning techniques.

5. FLOWCHART



6. RESULT





7. ADVANTAGES & DISADVANTAGES

Advantages:

- MRI provide long information concerning the human delicate tissue, that help within the conclusion of brain tumor.
- The experimental results show high accuracy in detecting brain tumors with the help of datasets.
- Inorder to detect the brain tumor of a patient ,MRI images of a patient's brain are considered. This application helps you detect the tumor and store the patient's data.

Disadvantages:

- The Convolutional Neural Network (CNN) was implemented, which drives an overall accuracy of 91.3% and a recall of 88%, 81% and 99% in the detection of meningioma, glioma and pituitary tumor respectively.
- Deep learning architecture by leveraging 2D convolutional neural networks for the classification of the different types of brain tumor from MRI image slices.

8. APPLICATIONS

- The main aim of the application is tumor identification, this application is to provide proper treatment as soon as possible and protect the human life which is in danger.
- This application is helpful to doctors as well as patients. It is a user friendly application

9. CONCLUSION

The proposed algorithms for Brain Tumor Detection based on MRI Images will have higher accuracy and low error rates. The statistical analysis of the experimental results has indicated that the developed algorithm can segment brain MR images with good accuracy. We find ways to detect these tumors at faster rate to improve survival rates. The result of the analysis is carried out whether the brain is normal one or diseased one by applying deep learning techniques.

10. FUTURE SCOPE

Brain tumor identification is a really challenging task in the early stages of life. These days the issue of brain tumor automatic identification is of great interest. Deep learning techniques can be used in order to detect the brain tumor of a patient using the MRI images of a patient's brain.

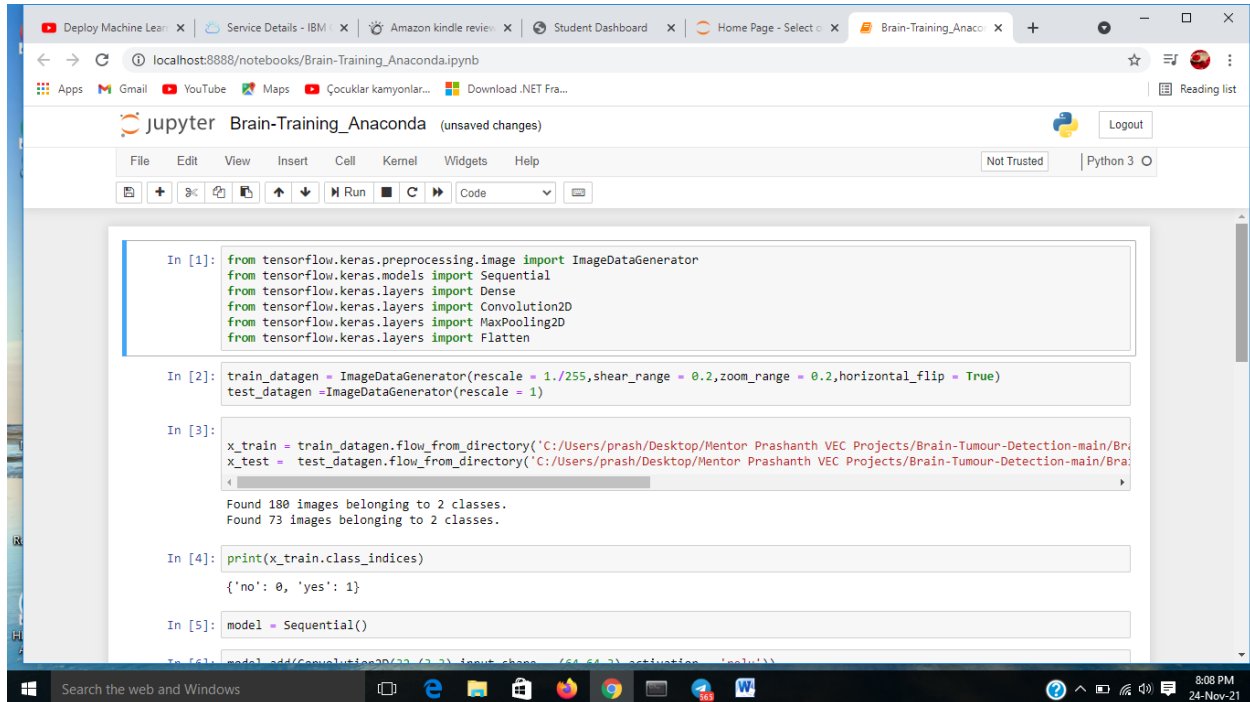
In this application, we are helping the doctors and patients to classify the type of scan for the specific image given with the help of Neural Networks and store the patient's data. This application is helpful to doctors as well as patients. It is a user friendly application.

11. BIBLIOGRAPHY

- https://en.wikipedia.org/wiki/Convolutional_neural_network
- <https://www.geeksforgeeks.org/introduction-convolution-neural-network/>
- https://en.wikipedia.org/wiki/Brain_tumor

APPENDIX

Source Code



The screenshot shows a Jupyter Notebook interface with the following code cells:

```
In [1]: from tensorflow.keras.preprocessing.image import ImageDataGenerator
        from tensorflow.keras.models import Sequential
        from tensorflow.keras.layers import Dense
        from tensorflow.keras.layers import Convolution2D
        from tensorflow.keras.layers import MaxPooling2D
        from tensorflow.keras.layers import Flatten

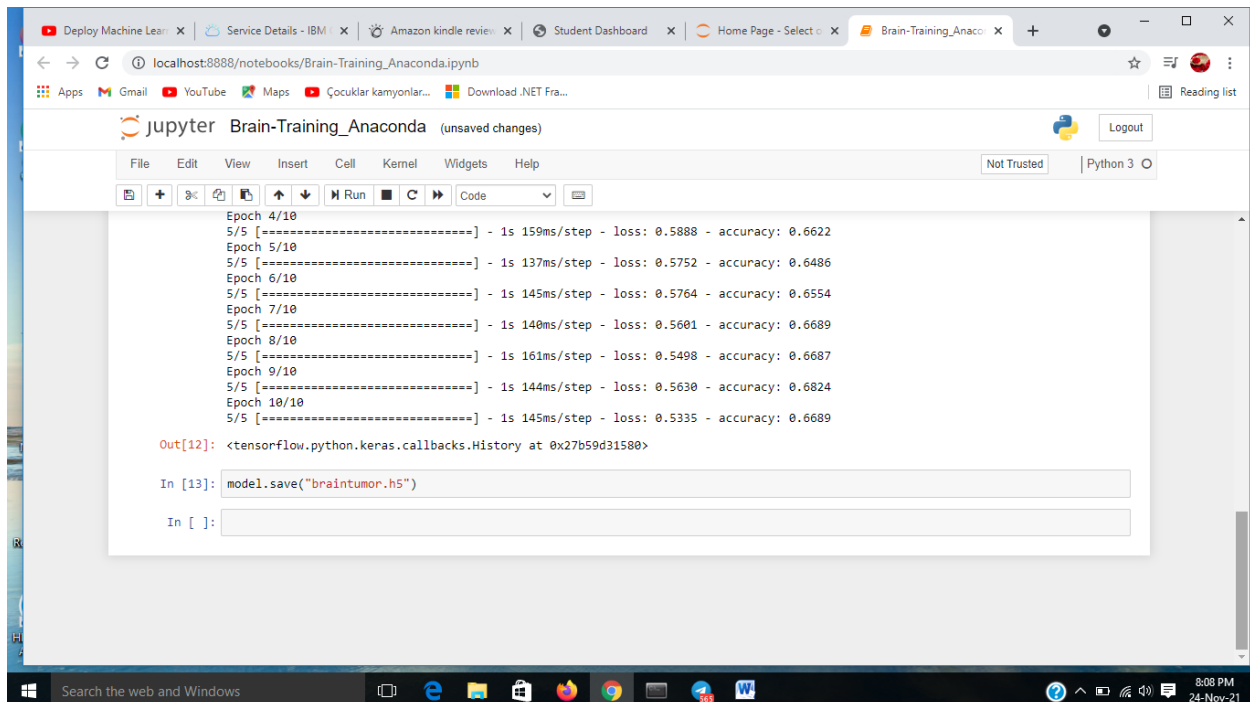
In [2]: train_datagen = ImageDataGenerator(rescale = 1./255, shear_range = 0.2, zoom_range = 0.2, horizontal_flip = True)
        test_datagen = ImageDataGenerator(rescale = 1)

In [3]: x_train = train_datagen.flow_from_directory('C:/Users/prash/Desktop/Mentor Prashanth VEC Projects/Brain-Tumour-Detection-main/Brain-Tumour-Detection-main/train')
        x_test = test_datagen.flow_from_directory('C:/Users/prash/Desktop/Mentor Prashanth VEC Projects/Brain-Tumour-Detection-main/Brain-Tumour-Detection-main/test')

Found 180 images belonging to 2 classes.
Found 73 images belonging to 2 classes.

In [4]: print(x_train.class_indices)
{'no': 0, 'yes': 1}

In [5]: model = Sequential()
```



The screenshot shows the continuation of the Jupyter Notebook with the following code cells:

```
Epoch 4/10
5/5 [=====] - 1s 159ms/step - loss: 0.5888 - accuracy: 0.6622
Epoch 5/10
5/5 [=====] - 1s 137ms/step - loss: 0.5752 - accuracy: 0.6486
Epoch 6/10
5/5 [=====] - 1s 145ms/step - loss: 0.5764 - accuracy: 0.6554
Epoch 7/10
5/5 [=====] - 1s 140ms/step - loss: 0.5601 - accuracy: 0.6609
Epoch 8/10
5/5 [=====] - 1s 161ms/step - loss: 0.5498 - accuracy: 0.6687
Epoch 9/10
5/5 [=====] - 1s 144ms/step - loss: 0.5630 - accuracy: 0.6824
Epoch 10/10
5/5 [=====] - 1s 145ms/step - loss: 0.5335 - accuracy: 0.6609

Out[12]: <tensorflow.python.keras.callbacks.History at 0x27b59d31580>

In [13]: model.save("braintumor.h5")

In [ ]:
```

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localhost:8888/notebooks/Brain_tumor_IBM_Download_model.ipynb

jupyter Brain_tumor_IBM_Download_model Last Checkpoint: 11/16/2021 (unsaved changes)

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```
In [1]: from ibm_watson_machine_learning import APIClient

In [2]: wml_credentials = {
        "url": "https://us-south.ml.cloud.ibm.com",
        "apikey": "RXglttWmDdOu-CjVMGmHmZzewsg6bHswNgnH5QKktXL"
    }
    client = APIClient(wml_credentials)

In [3]: client = APIClient(wml_credentials)

In [4]: def guid_from_space_name(client, space_name):
        space = client.spaces.get_details()
        return(next(item for item in space['resources'] if item['entity']['name'] == space_name)['metadata']['id'])

In [5]: space_uid = guid_from_space_name(client, 'deployment_brain_t')
        print("Space UID="+space_uid)

        Space UID=e25a99a4-a8bd-40ad-90e5-0c40b5f79a5f

In [6]: client.set_default_space(space_uid)

Out[6]: 'SUCCESS'

In [7]: client.software_specifications.list()
```

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Deploy Mach x Service Deta x Amazon kind x Student Dash x Home Page x Brain-Test x Brain_tumor x Brain-Trainin x Brain-Trainin x +

localhost:8888/notebooks/Brain-Testing.ipynb

jupyter Brain-Testing (autosaved)

File Edit View Insert Cell Kernel Widgets Help Not Trusted Python 3

```
WARNING:tensorflow:From C:\Users\Sai Nidhi\Anaconda3\lib\site-packages\keras\backend\tensorflow_backend.py:181: The name tf.ConfigProto is deprecated. Please use tf.compat.v1.ConfigProto instead.

WARNING:tensorflow:From C:\Users\Sai Nidhi\Anaconda3\lib\site-packages\keras\backend\tensorflow_backend.py:186: The name tf.Session is deprecated. Please use tf.compat.v1.Session instead.

WARNING:tensorflow:From C:\Users\Sai Nidhi\Anaconda3\lib\site-packages\keras\optimizers.py:790: The name tf.train.Optimizer is deprecated. Please use tf.compat.v1.train.Optimizer instead.

WARNING:tensorflow:From C:\Users\Sai Nidhi\Anaconda3\lib\site-packages\tensorflow\python\ops\nn_impl.py:180: add_dispatch_support.<locals>.wrapper (from tensorflow.python.ops.array_ops) is deprecated and will be removed in a future version.
Instructions for updating:
Use tf.where in 2.0, which has the same broadcast rule as np.where

In [3]: img = image.load_img('Y3.jpg', target_size = (64,64))

In [4]: x = image.img_to_array(img)
        x = np.expand_dims(x, axis = 0)

In [5]: pred = model.predict_classes(x)

In [6]: pred

Out[6]: array([[1]])

In [ ]:
```

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