

BRAIN TUMOUR IMAGE SEGMENTATION USING DEEP NETWORKS

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Abstract

Automated segmentation of brain tumour from multimodal MR images is pivotal for the analysis and monitoring of disease progression. As gliomas are malignant and heterogeneous, efficient and accurate segmentation techniques are used for the successful delineation of tumours into intra-tumoural classes. Deep learning algorithms outperform tasks of semantic segmentation as opposed to the more conventional, context-based computer vision approaches. Extensively used for biomedical image segmentation, Convolutional Neural Networks have significantly improved the state-of-the-art accuracy on the task of brain tumour segmentation. In this paper, we propose an ensemble of two segmentation networks: CNN and a U-Net, in a significant yet straightforward combinative technique that results in better and accurate predictions. Both models were trained separately on the BraTS-19 challenge dataset and evaluated to yield segmentation maps which considerably differed from each other in terms of segmented tumour sub-regions and were ensembled variably to achieve the final prediction. The suggested ensemble achieved dice scores of 0.750, 0.906 and 0.846 for enhancing tumour, whole tumour, and tumour core, respectively, on the validation set, performing favourable in comparison to the state-of-the-art architectures currently available.

Keywords—Deep learning, BraTS, Medical segmentation, U-Net, CNN, ensembling.

1. INTRODUCTION

Accurate segmentation of tumours through medical images is of particular importance as it provides information essential for analysis and diagnosis of cancer as well as for mapping out treatment options and monitoring the progression of the disease. Brain tumours are one of the fatal cancers worldwide and are categorised, depending upon their origin, into primary and secondary tumour types. The most common histological form of primary brain cancer is the glioma, which originates from the astroglial cells and attributes towards 80% of all malignant brain tumour. Gliomas can be of the slow-progressing low-grade (LGG) subtype with a better patient prognosis or are the more aggressive and infiltrative high-grade glioma (HGG) or glioblastoma, which require immediate treatment. These tumours are associated with substantial morbidity, where the median survival for a patient with glioblastoma is only about 14 months with a 5-year survival rate near zero despite maximal surgical and medical therapy. A timely diagnosis, therefore, becomes imperative for effective treatment of the patients. Magnetic Resonance Imaging (MRI) is a preferred technique widely employed by radiologists for the evaluation and assessment of brain tumours. It provides several complementary 3D MRI modalities acquired based on the degree of excitation and repetition times, i.e. T1-weighted, post-contrast T1-weighted (T1ce), T2-

weighted and Fluid Attenuated Inversion Recovery (FLAIR). The highlighted sub regions of the tumour across different intensities of these sequences, such as the whole tumour (the entire tumour inclusive of infiltrative oedema), is more prominent in FLAIR and T2 modalities. In contrast, T1 and T1ce images show the tumour core exclusive of peritumoural oedema. It allows for the combinative use of these scans and the complementary information they deliver towards the detection of different tumours sub regions.

2. RELATED WORK

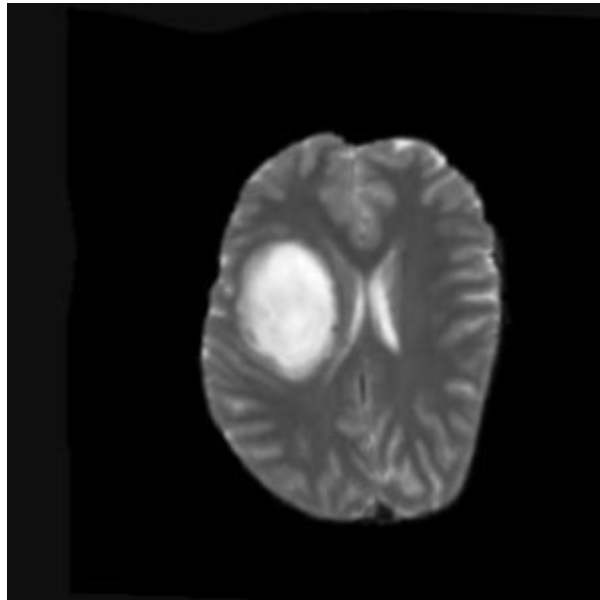
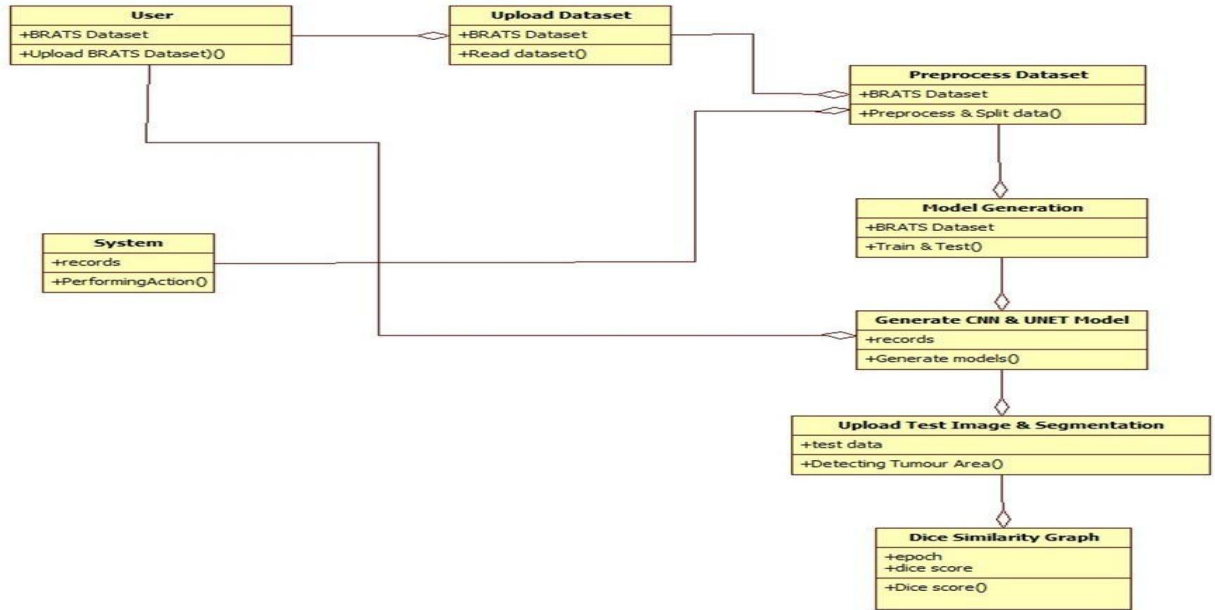
In this project we report the set-up and results of the Multi modal Brain Tumor Image Segmentation Benchmark (BRATS) organized in conjunction with the MICCAI 2012 and 2013 conferences. Twenty state-of-the-art tumor segmentation algorithms were applied to a set of 65 multi-contrast MR scans of low- and high-grade glioma patients – manually annotated by up to four raters – and to 65 comparable scans generated using tumor image simulation software. Quantitative evaluations revealed considerable disagreement between the human raters in segmenting various tumor sub-regions (Dice scores in the range 74-85%), illustrating the difficulty of this task. We found that different algorithms worked best for different sub-regions (reaching performance comparable to human inter-rater variability), but that no single algorithm ranked in the top for all sub regions simultaneously.

3. PROPOSED FRAMEWORK

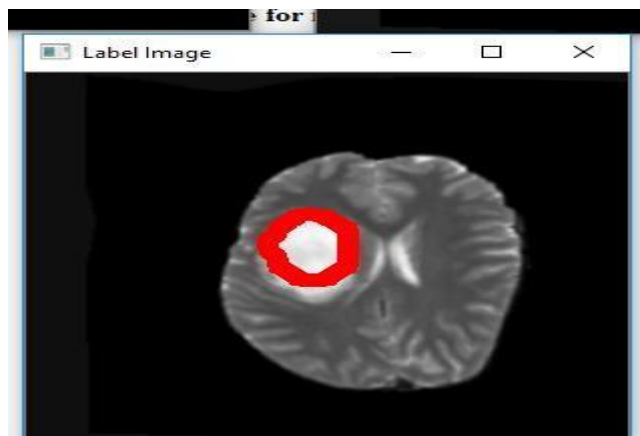
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CONVOLUTIONAL NEURAL NETWORKS (CNN)

To demonstrate how to build a convolutional neural network based image classifier, we shall build a 6 layer neural network that will identify and separate one image from other. This network that we shall build is a very small network that we can run on a CPU as well. Traditional neural networks that are very good at doing image classification have many more parameters and take a lot of time if trained on normal CPU. However, our objective is to show how to build a real-world convolutional neural network using TENSOR FLOW. Neural Networks are essentially mathematical models to solve an optimization problem. They are made of neurons, the basic computation unit of neural networks. A neuron takes an input (say x), do some computation on it (say: multiply it with a variable w and add another variable b) to produce a value (say; $z = wx + b$). This value is passed to a non-linear function called activation function (f) to produce the final output (activation) of a neuron. There are many kinds of activation functions. One of the popular activation function is Sigmoid. The neuron which uses sigmoid function as an activation function will be called sigmoid neuron.



WITHOUT IDENTIFYING BRAIN TUMOUR



WITH IDENTIFYING BRAIN TUMOUR

4. METHODOLOGY

MODULES:

Upload BRATS Dataset:

In this module user upload dataset of BRATS.

Generate CNN & UNET Model

We can see models are using different size images to filter the mand to get best features from it to build efficient model and now model is generate.

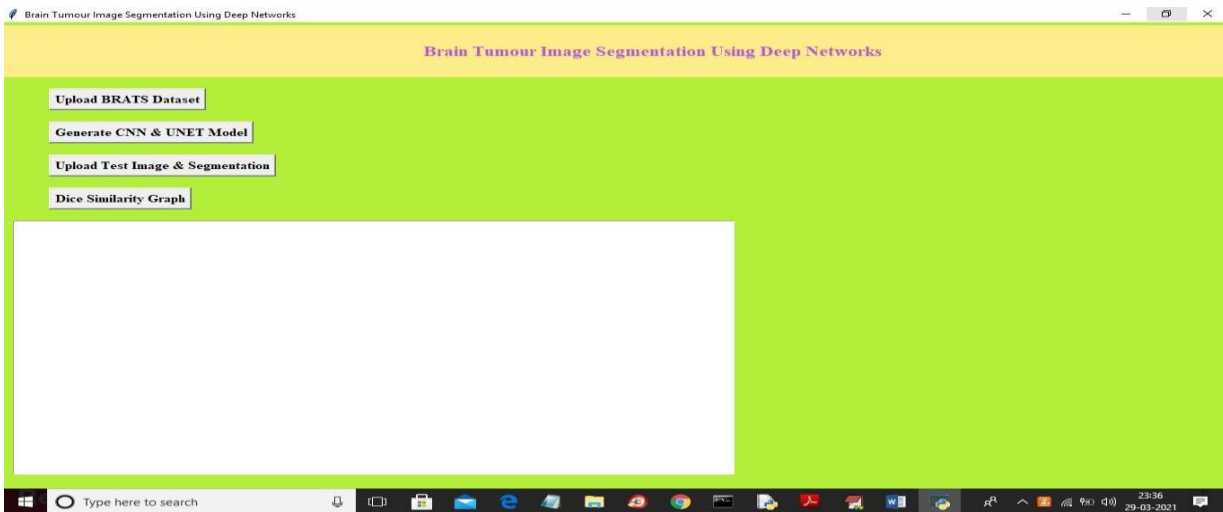
Upload Test Image & Segmentation

In this module user upload test image and top 4 images are the input images such as FLAIR, T1, T2 and T1CE and 5th image is the predicted image with segmented part showing in red colour and this algorithm correctly detecting and marking tumour area.

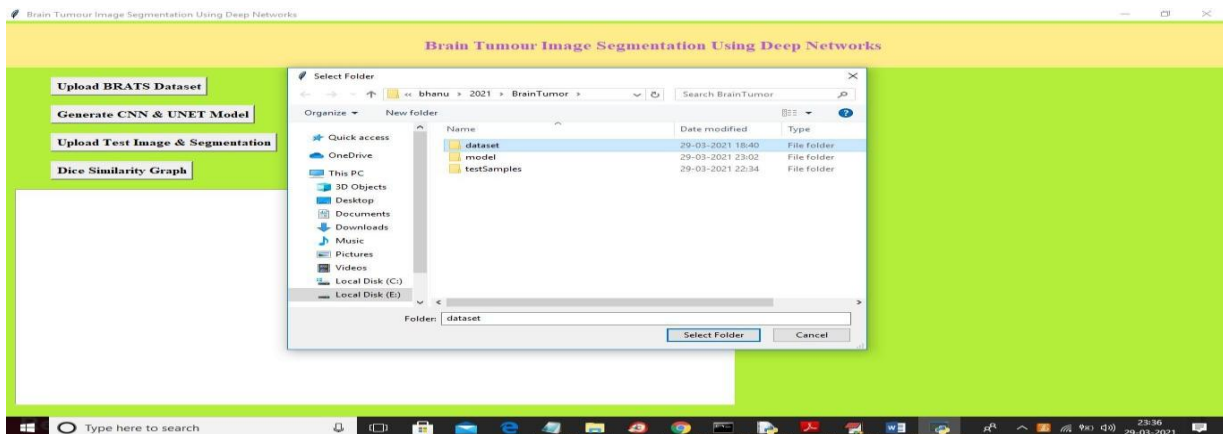
Dice Similarity Graph

In this Module, we get final dice score as $0.8 * 100 = 80\%$. In above graph x-axis represents epoch and y-axis represents dicescore.

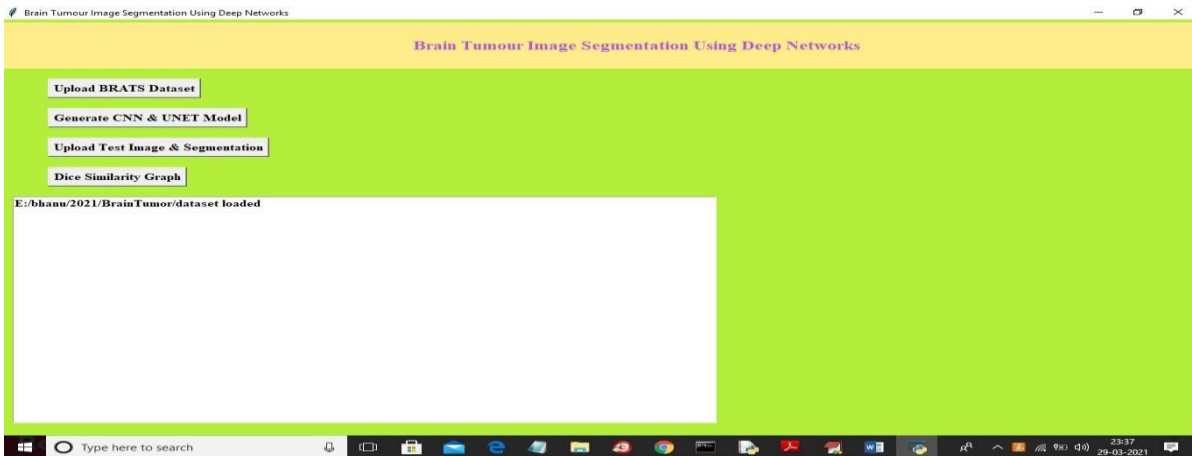
5. RESULTS AND DISCUSSION



5.1 MAIN ACTIVITY



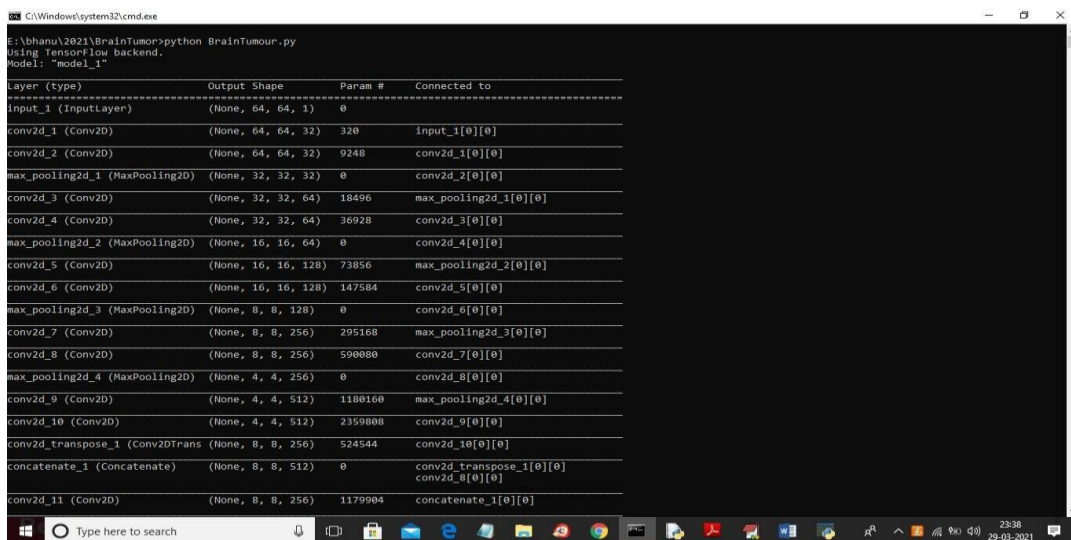
5.2 UPLOAD THE DATASET



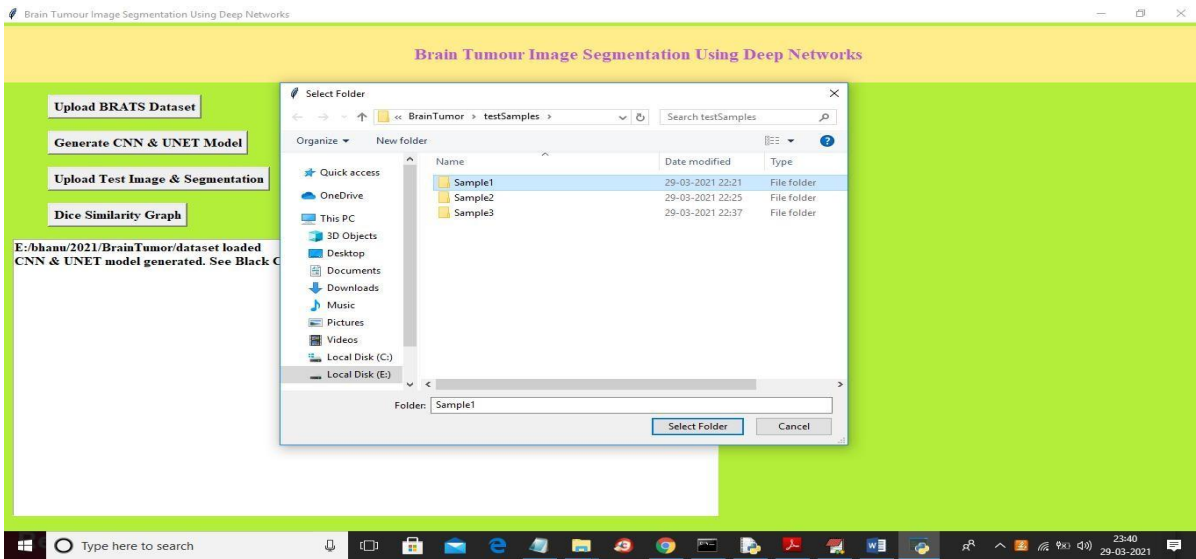
5.3 DATASET LOADED



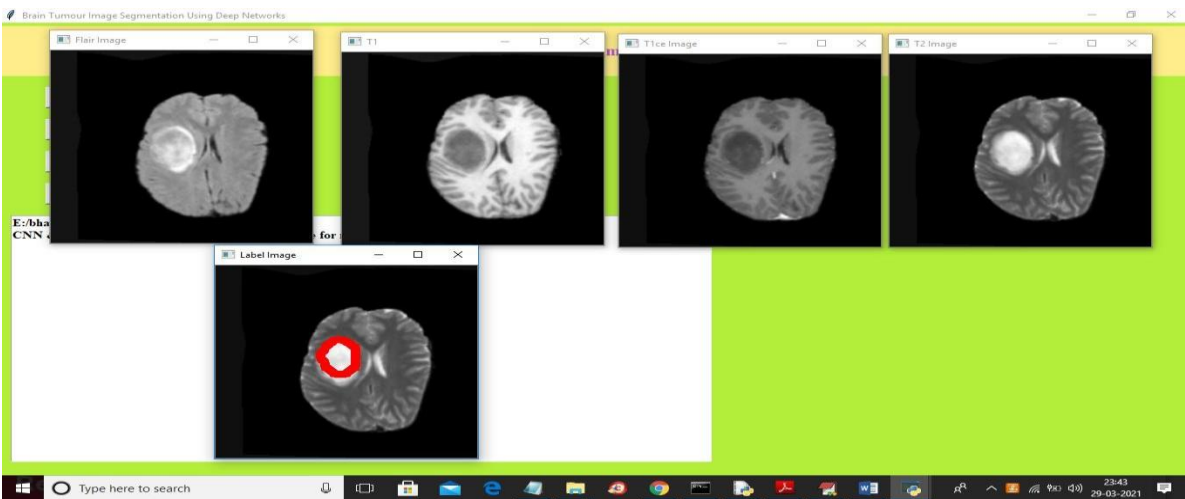
5.4 BOTH CNN AND U-NET MODELS ARE GENERATED



5.5 MODELS ARE USING DIFFERENT SIZE IMAGES



5.5 SELECTING AND UPLOADING SAMPLE1 FOLDER



5.6 In above screen first 4 images are the input images and fifth image is the predicted label image with segmented parts around tumour area



5.7 ACCURACYGRAPH

6. CONCLUSIONANDFUTUREWORK

In this paper we are using Image Segmentation method. We have used a hybrid of two different techniques, i.e. Watershed and Contrast Technique. This technique is well suited for detection of tumor in the image. This segmentation method gives high accuracy as compared to other methods. MRI images are best suitable for brain tumor detection. In this study Digital Image Processing Techniques are important for brain tumor detection by MRI images. The pre processing techniques include different methods like Filtering, Contrast enhancement, Edge detection is used for image smoothing. The preprocessed images are used for post processing operations like; threshold, histogram, segmentation and morphological, which is used to enhance the images.

7. REFERENCES

J.Vijay et.al [3], propose the work on automated brain tumor detection by using segmentation by k-means algorithm and object labeling algorithm. They identified that a well known segmentation problem within MRI is the task of labeling the tissue type which include White Matter (WM), Grey Matter (GM), Cerebrospinal Fluid (CSF) and sometimes pathological tissues like tumor. S.Koley et.al[4], propose the efficient work on tumor detection and segmentation of brain MRI for the purpose of determining the exact location of brain tumor using CSM based partitioned K-means clustering algorithm. CSM has attracted much attention as it has given efficient result as a self merging algorithm compared to other merging processes and the effect of noise is also less and the probability of obtaining the exact location of tumor is more. Their approach is much simpler and computationally less complex and computation time is very less.