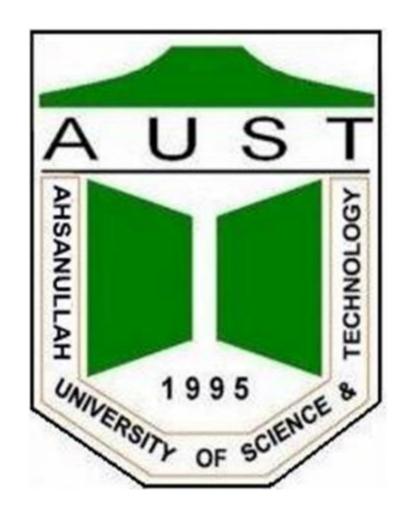
International Conference on Advances in Science, Engineering and Robotics Technology (ICASERT-2019)



Ahsanullah University of Science and Technology

Dept. of Computer Science and Engineering

Brain Tumor Detection Using Convolutional Neural Network

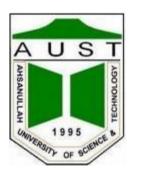
Tonmoy Hossain, Fairuz Shadmani Shishir, Mohsena Ashraf MD Abdullah Al Nasim, Faisal Muhammad Shah

HELLO!

I AM TONMOY HOSSAIN

4th Year 2nd Semester
Department of CSE
Ahsanullah University of Science and Technology

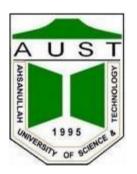
INTRODUCTION



INTRODUCTION

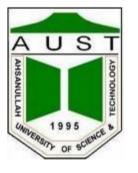
- ✓ In the field of Medical Image Analysis, research on Brain tumors is one of the most prominent ones
- Tumor segmentation is one of the most arduous task
- $\sqrt{}$ Primary brain tumors occur in around 250,000 people a year globally, making up less than 2% of cancers^[1]

[1]. "Chapter 5.16" World Cancer Report 2014. World Health Organization. 2014. ISBN 978-9283204299. Archived from the original on 02 May 2019.



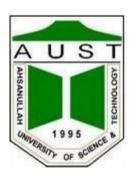
MOTIVATION

- √ Well adaptation of automated medical image analysis in the perspective of Bangladesh
- Early detection of Brain Tumors
- Reducing the pressure on Human judgement
- Build a User Interface which can identify the cancerous cells

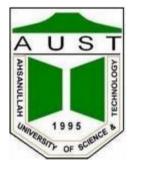


CHALLENGES

- Device Independent
- Real-time in erratic background
- Segmenting tumors conjoined with the skull
- Reducing processing time by scaling the hidden layers



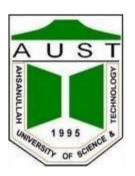
RESEARCH DOMAIN



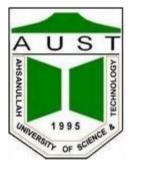
Problem

- Segmentation of the tumorous cells
- Detection of the Tumor
- Extract extensive features from the tumor

- **How we can implement the problem?**
 - Basic Image Processing techniques was used for segmentation
 - Using Convolutional Neural Network based detection

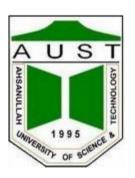


BACKGROUNDS

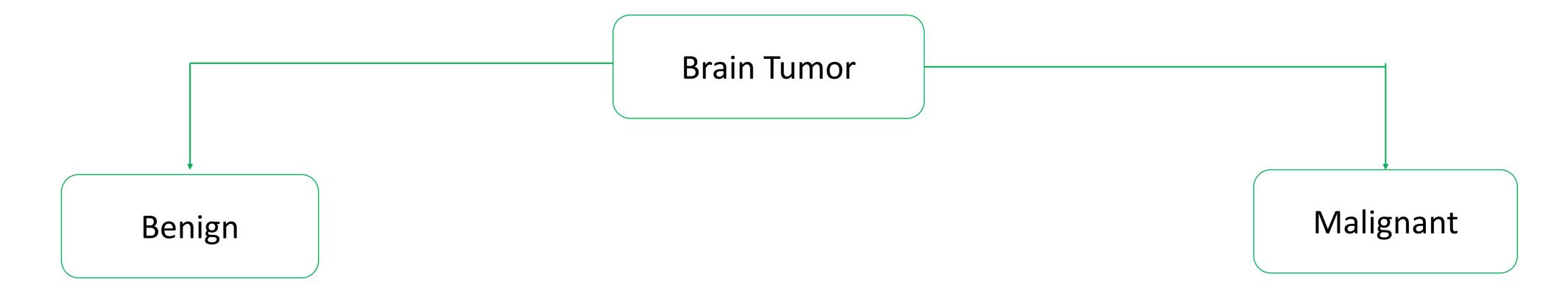


BRAIN TUMOR

- (v) tumor cells remain undifferentiated in the image
- (cells contain abnormal nuclei
- (v) abnormal cells form within the brain
- (many dividing cells: disorganized arrangement
- destroy healthy brain cells by invading them
- (\checkmark) tumor may grow from neuroma, meningioma, craniopharyngioma or glioma

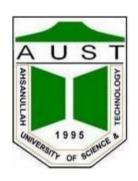


Types of Brain Tumor

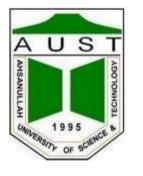


- (non cancerous
- grows slowly: do not spread into other tissues
- have clear borders

- (brain cancers
- grows rapidly and invades healthy brain tissues
- distorted borders

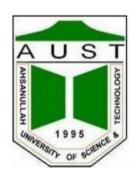


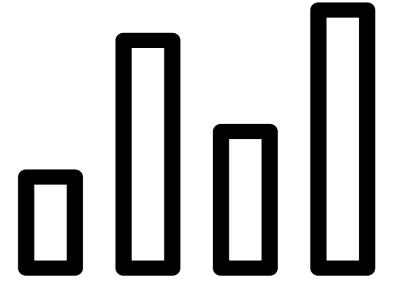
BACKGROUND STUDIES



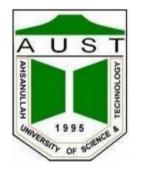
Existing Works

- O Devkota et al. 2017
 - "Image Segmentation for Early Stage Brain Tumor Detection using Mathematical Morphological Reconstruction"
- Song et al. 2016
 - "A Novel Brain Tumor Segmentation from Multi-Modality MRI via A Level-Set-Based Model"
- (Dina et al. 2012
 - "Automated Brain Tumor Detection and Identification using Image Processing and Probabilistic Neural Network Techniques"
- Zahra et al. 2018
 - "Brain Tumor Segmentation Using Deep Learning by Type Specific Sorting of Images"





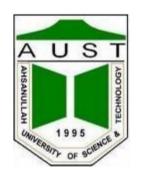
AREVIEW



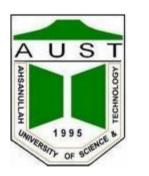
Brain Tumor Segmentation Techniques on Medical Images - A Review^[2]

- A total of 52 papers had been reviewed including Machine learning and Deep learning methods
- The whole review divided in Layer based, Region based, Edge based, Thresholding based segmentation techniques etc.
- Clustering technique was used in majority of the articles
- For Classification, K-Means, Fuzzy C-Means algorithm had been used

[2]. Faisal Muhammad Shah, Tonmoy Hossain, Mohsena Ashraf, Fairuz Shadmani Shishir, MD Abdullah Al Nasim, Md. Hasanul Kabir, "Brain Tumor Segmentation Techniques on Medical Images - A Review", INTERNATIONAL JOURNAL OF SCIENTIFIC & ENGINEERING RESEARCH, VOLUME 10, ISSUE 2, FEBRUARY-2019, ISSN 2229-5518.



Dataset

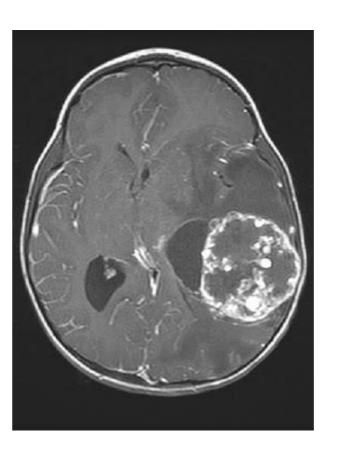


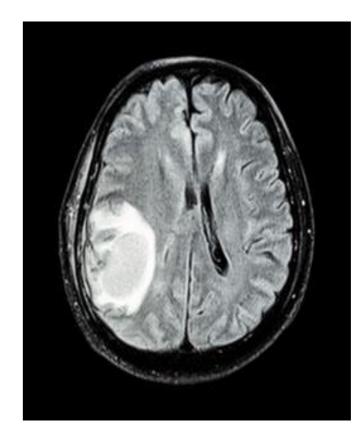
Dataset

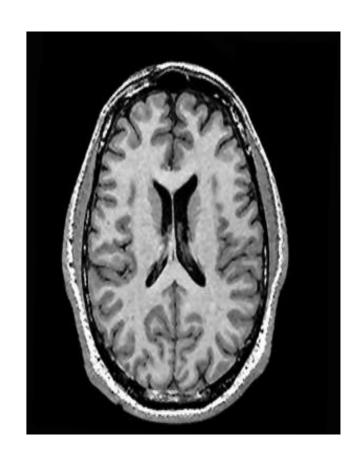
- (V) BraTS'13 data^{[3][4]}
- Total MRI Image: 217
- (Break down intro two category: class-0 and class-1
- All the MRI images are clinically-acquired preoperative multimodal scans of HGG and LGG
- O Described as- T1, T1Gd, T2 and FLAIR volumes



Some Examples



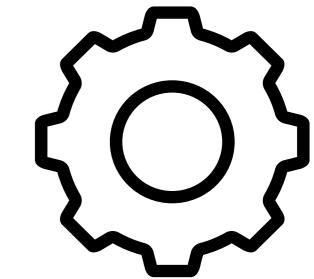




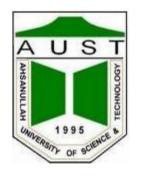
[3] Menze BH, Jakab A, Bauer S, Kalpathy-Cramer J, Farahani K, Kirby J, Burren Y, Porz N, Slotboom J, Wiest R, Lanczi L, Gerstner E, Weber MA, Arbel T, Avants BB, Ayache N, Buendia P, Collins DL, Cordier N, Corso JJ, Criminisi A, Das T, Delingette H, Demiralp Γ, Durst CR, Dojat M, Doyle S, Festa J, Forbes F, Geremia E, Glocker B, Golland P, Guo X, Hamamci A, Iftekharuddin KM, Jena R, John NM, Konukoglu E, Lashkari D, Mariz JA, Meier R, Pereira S, Precup D, Price SJ, Raviv TR, Reza SM, Ryan M, Sarikaya D, Schwartz L, Shin HC, Shotton J, Silva CA, Sousa N, Subbanna NK, Szekely G, Taylor TJ, Thomas OM, Tustison NJ, Unal G, Vasseur F, Wintermark M, Ye DH, Zhao L, Zhao B, Zikic D, Prastawa M, Reyes M, Van Leemput K. "The Multimodal Brain Tumor Image Segmentation Benchmark (BRATS)", IEEE Transactions on Medical Imaging 34(10), 1993-2024 (2015) DOI: 10.1109/TMI.2014.2377694

[4] Bakas S, Akbari H, Sotiras A, Bilello M, Rozycki M, Kirby JS, Freymann JB, Farahani K, Davatzikos C. "Advancing The Cancer Genome Atlas glioma MRI collections with expert segmentation labels and radiomic features", Nature Scientific Data, 4:170117 (2017) DOI: 10.1038/sdata.2017.117





METHODOLOGY (Segmentation)



Proposed Method for tumor segmentation and classification using traditional classifiers

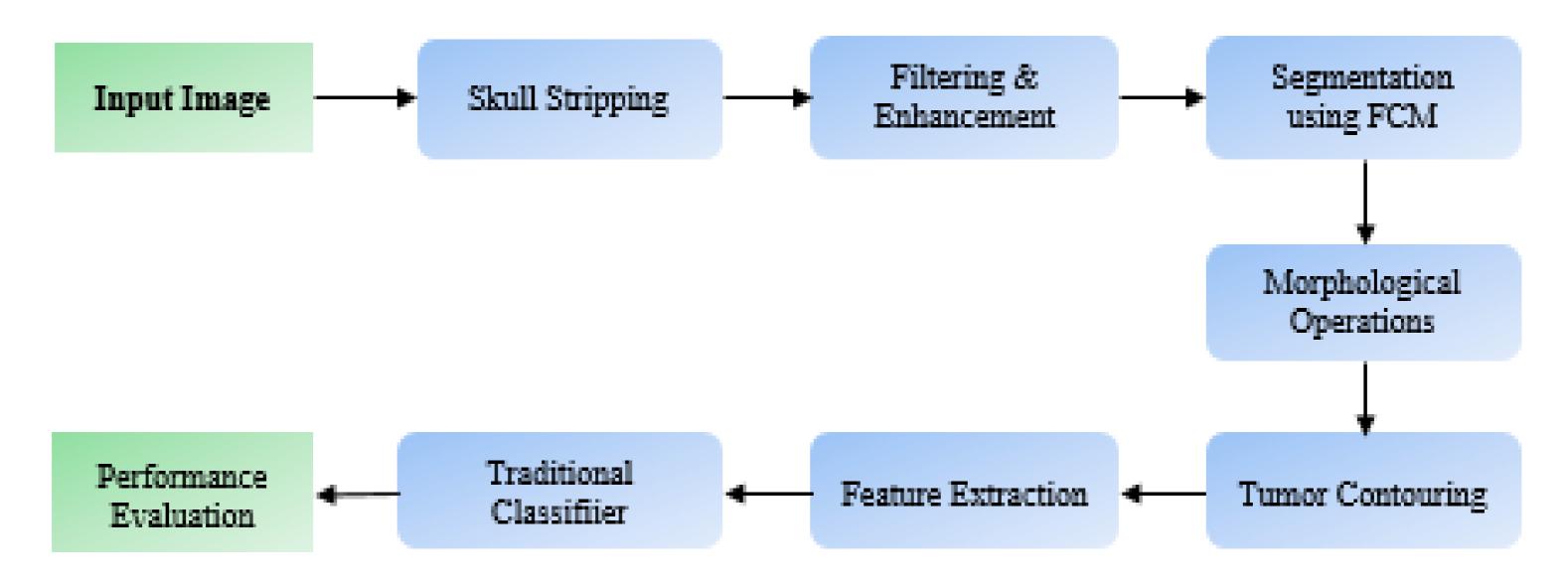
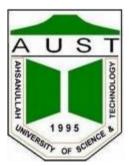


Fig 1: Proposed methodology for classification using Traditional Classifiers



Elaborated proposed methodology

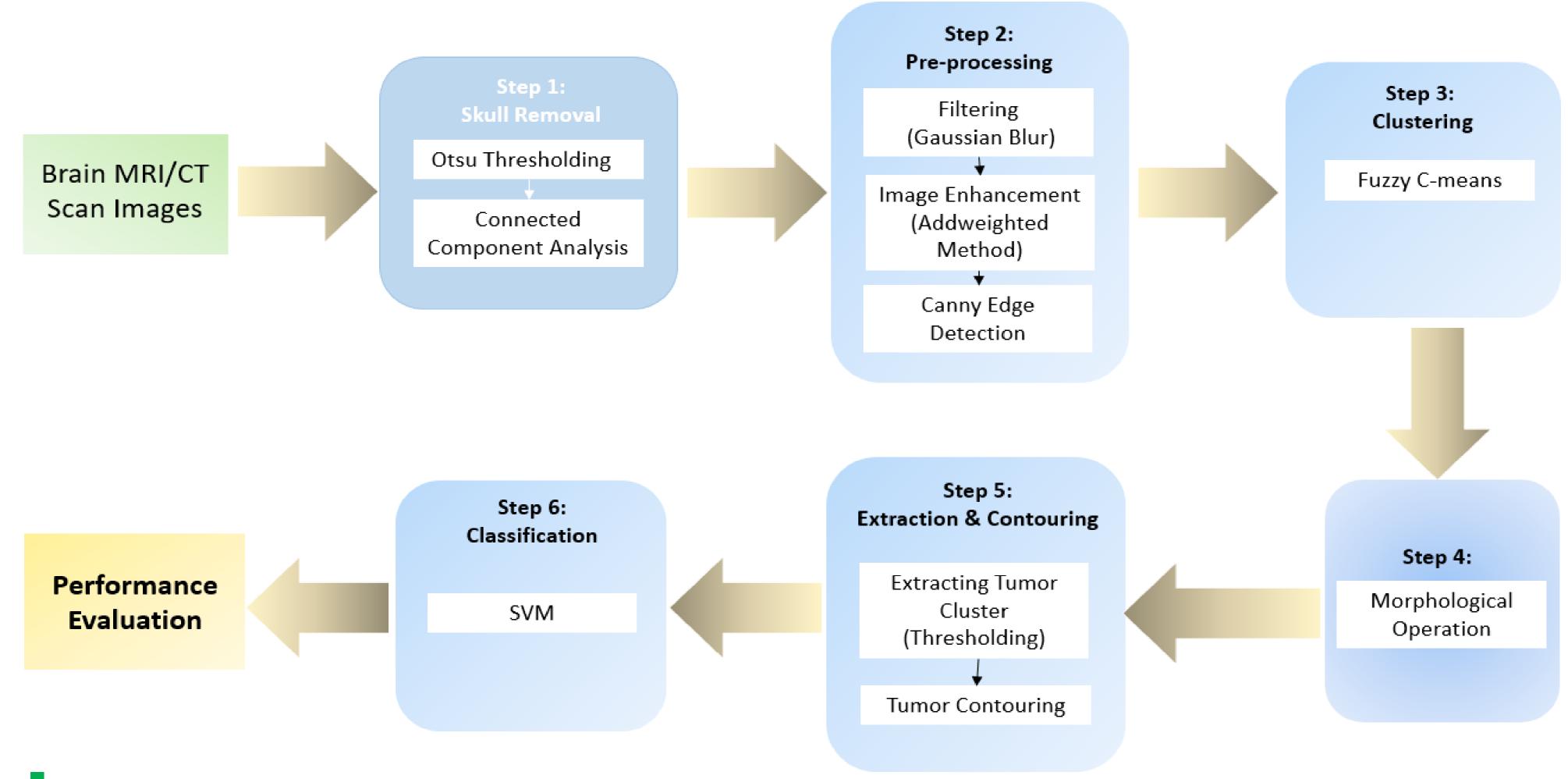




Fig 2: elaborated proposed methodology

Skull Stripping

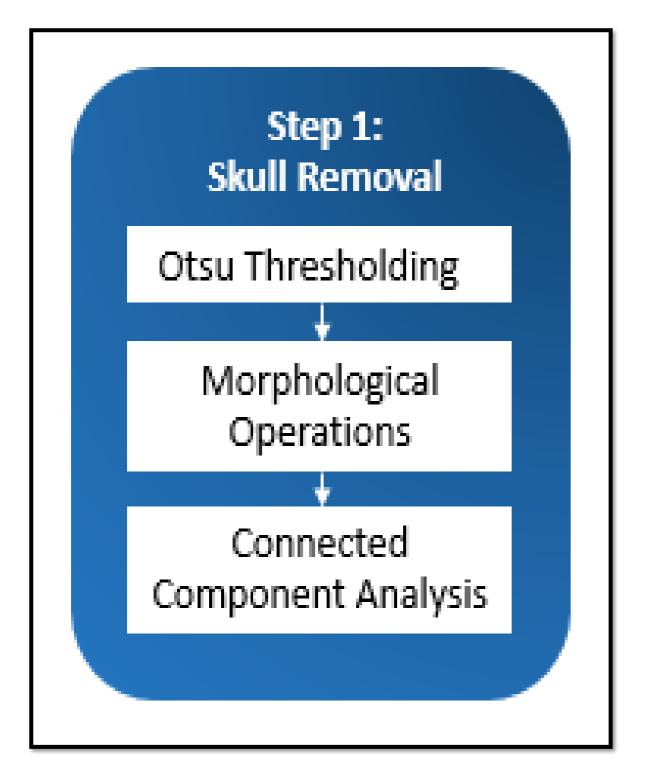


Fig 3: process of skull removal

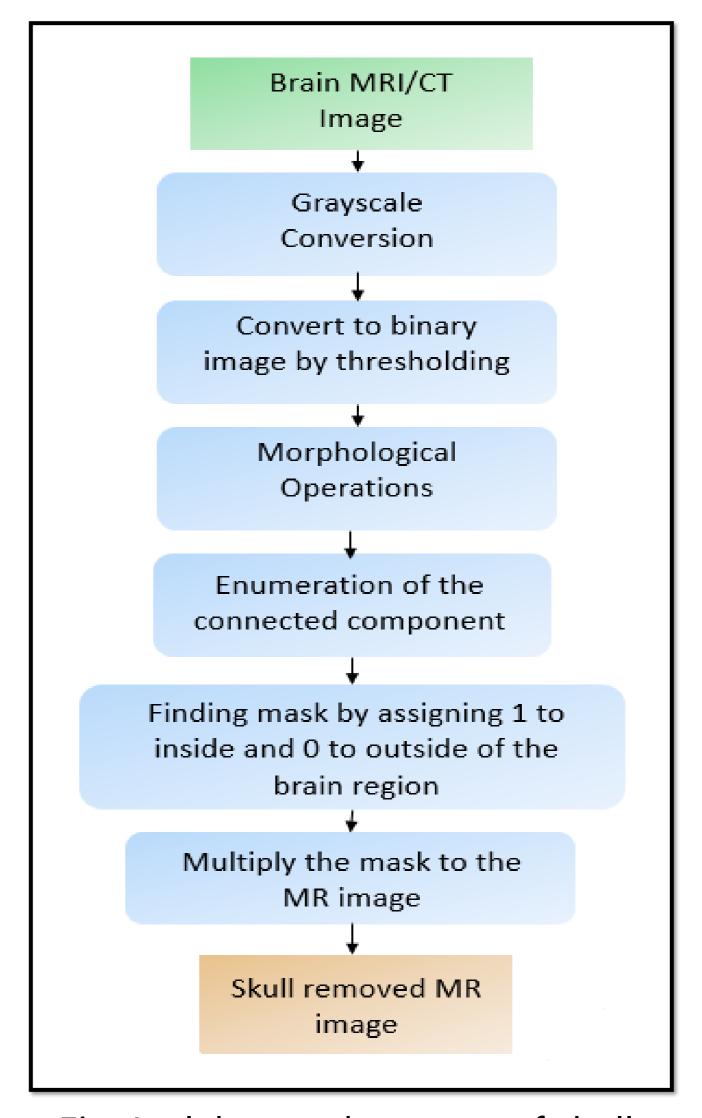
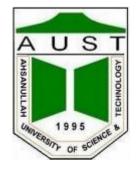


Fig 4: elaborated process of skull removal





Skull Stripping

- Converted our MRI Images into Grayscale
- OTSU Thresholding was applied for binarization
- Erosion operation had been performed before applying connected component analysis
- Each maximal region of connected pixels (not separated by boundary) is called a connected component. We found the largest component which is the skull
- We found the mask by assigning 1 to inside and 0 to outside of the brain region
- Multiplied the mask to T1, T2 and FLAIR images

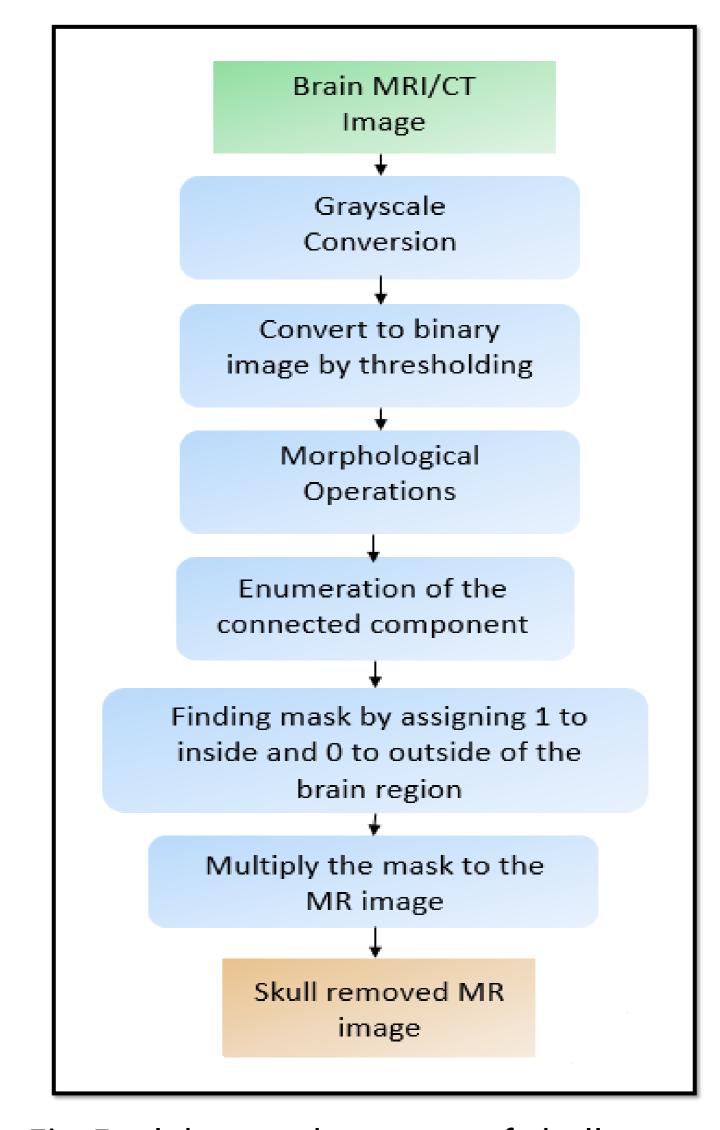
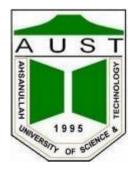


Fig 5: elaborated process of skull removal



Skull Stripping

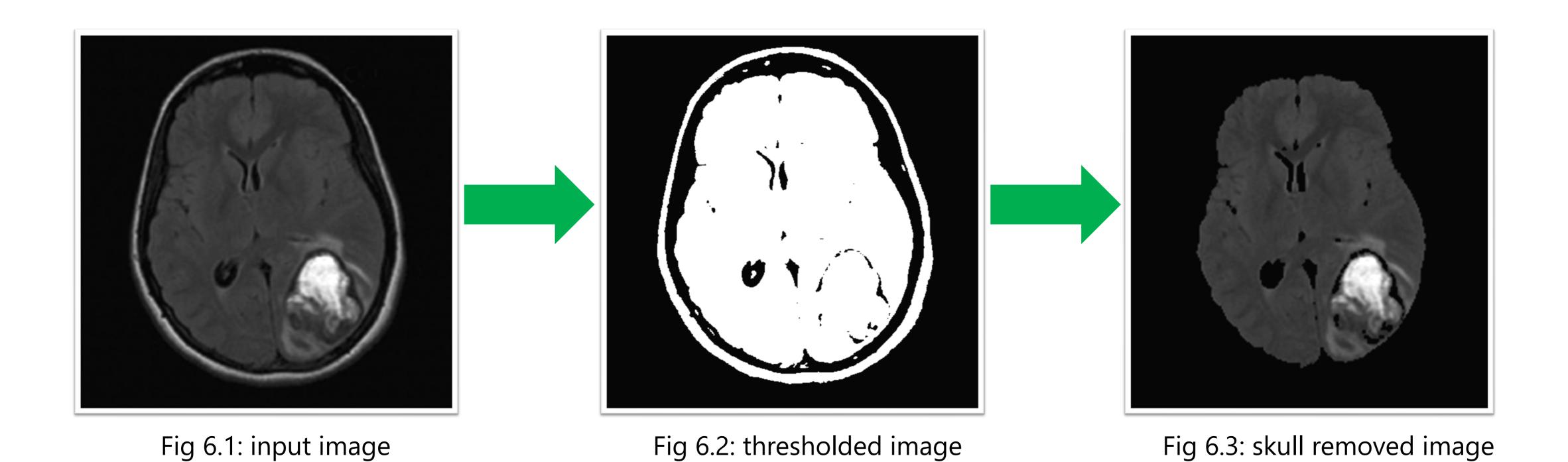
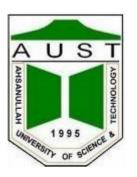
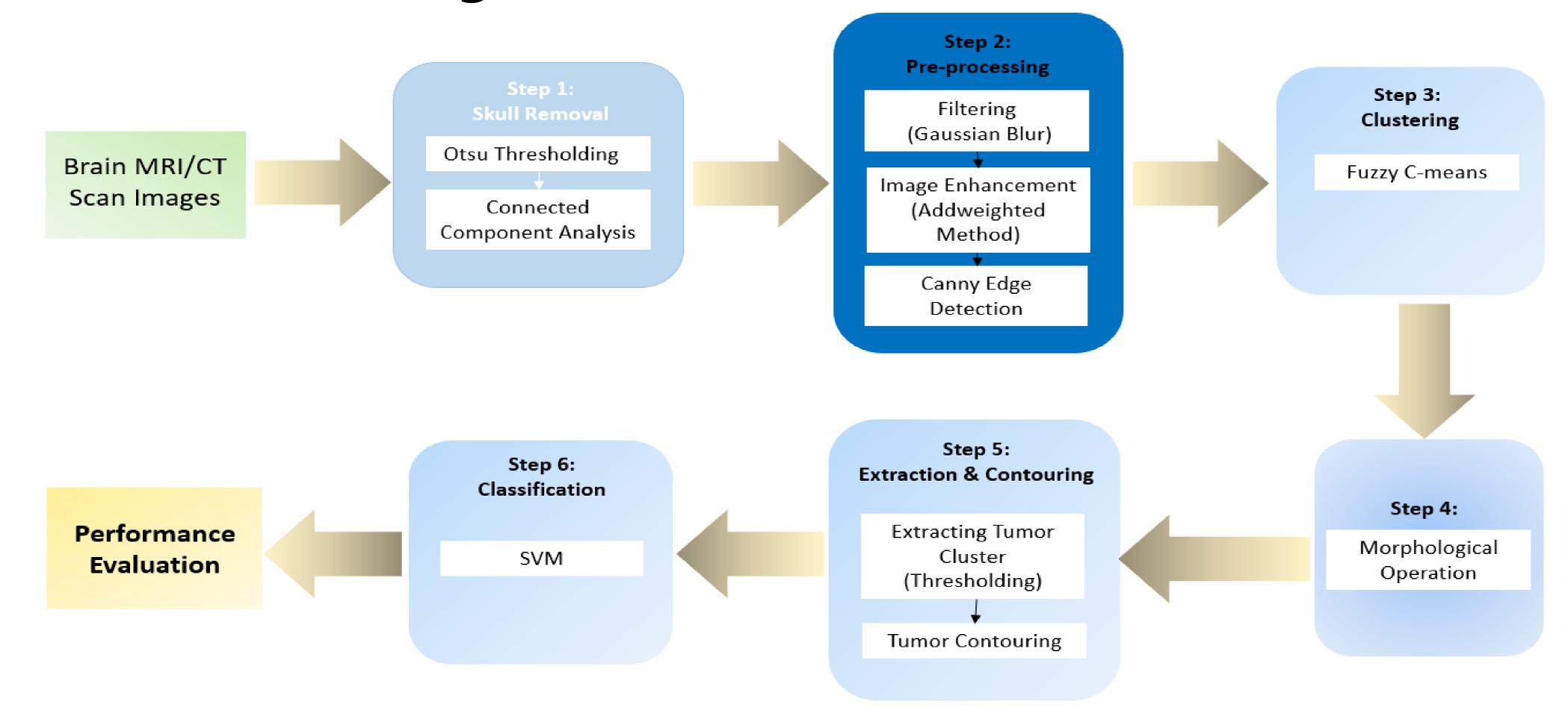
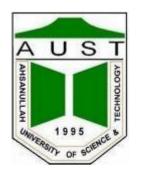


Fig 6: steps of skull stripping



Pre-Processing





Pre-Processing

- (Median filter gives us the most prominent result among the filters
- (v) For enhancing the image quality, we used the add-weighted method
- Applied the Canny Edge Detection method for detecting the edges

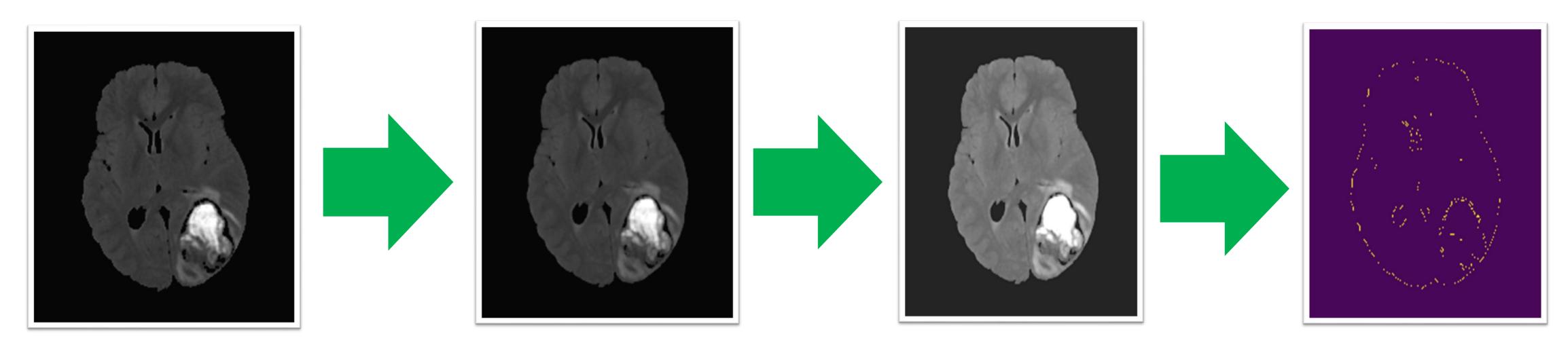


Fig 7.1: skull removed MRI

Fig 7.2: gaussian Blur Filter

Fig 7.3: enhanced MRI

Fig 7.4: edge detection MRI

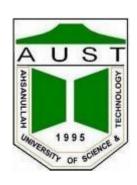
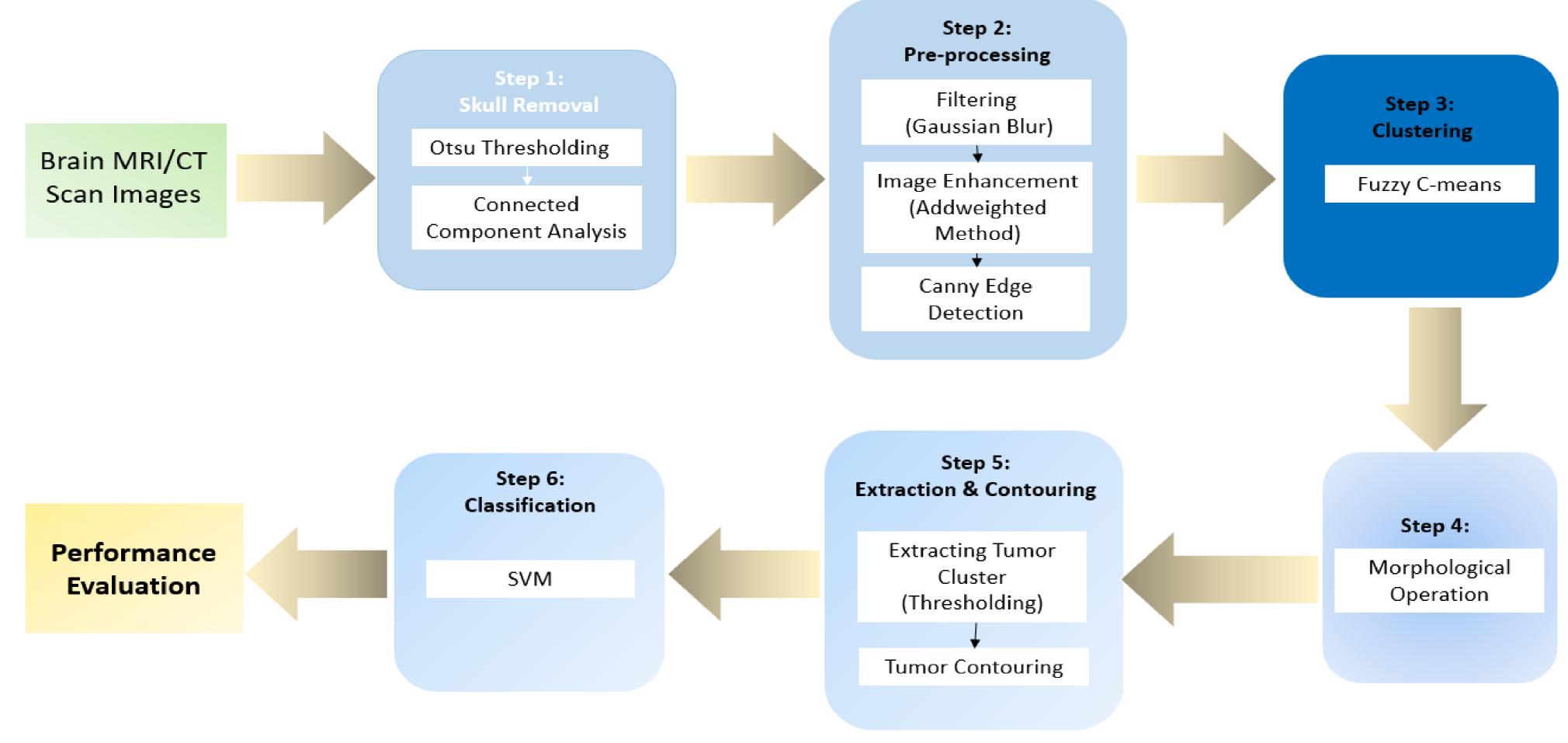
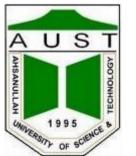


Fig 7: steps of pre processing

Segmentation Using FCM







Segmentation Using FCM

- A method of clustering which allows one piece of data to belong to two or more clusters
- Involves assigning data points to clusters
- Items in the same cluster are as similar as possible
- Items belonging to different clusters are as dissimilar as possible

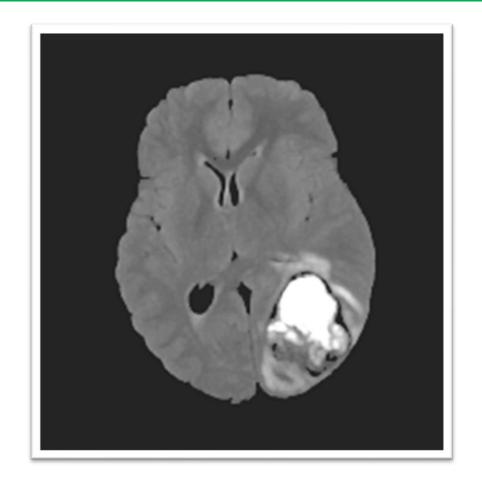
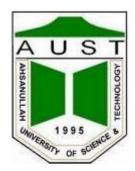


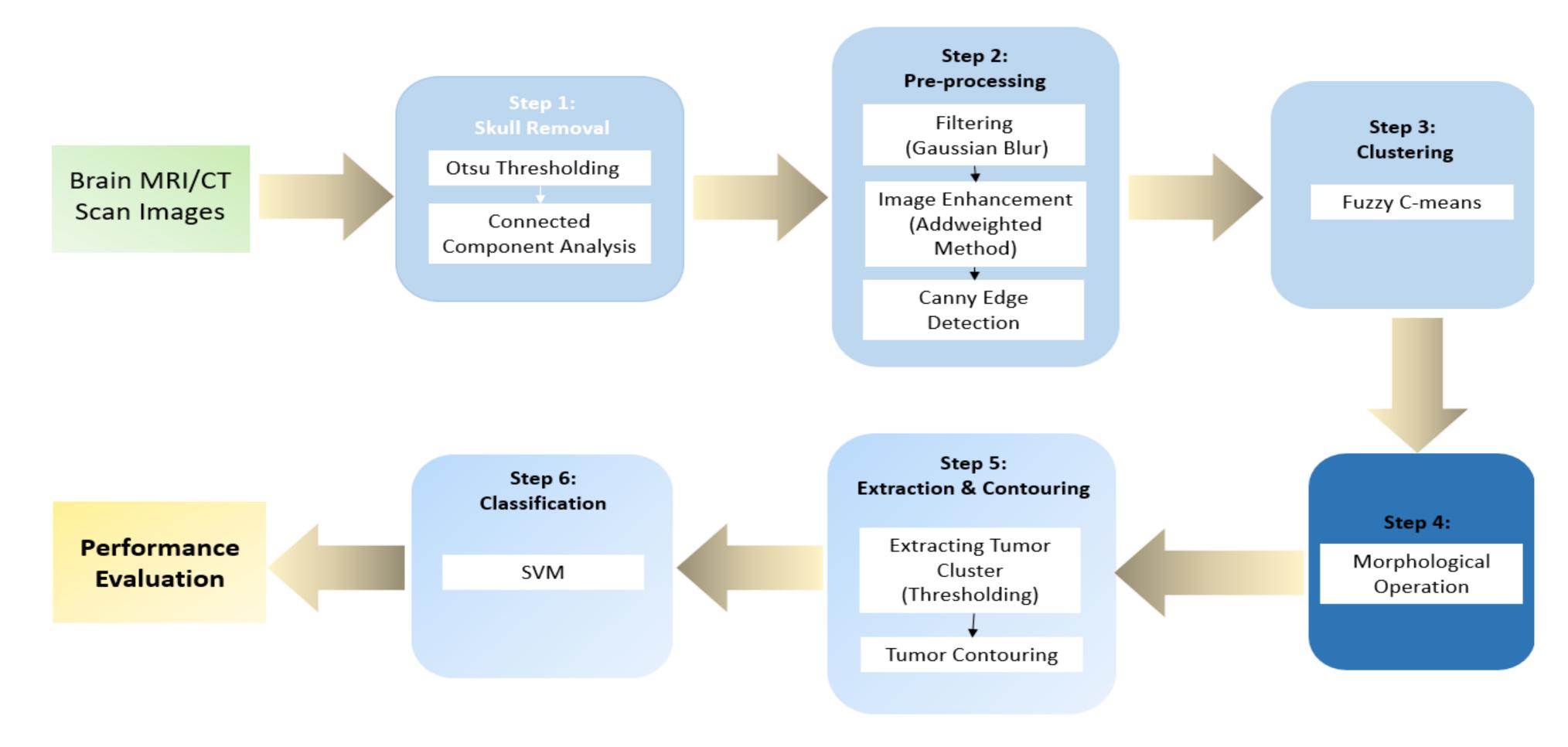


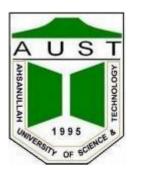


Fig 8: segmented tumor

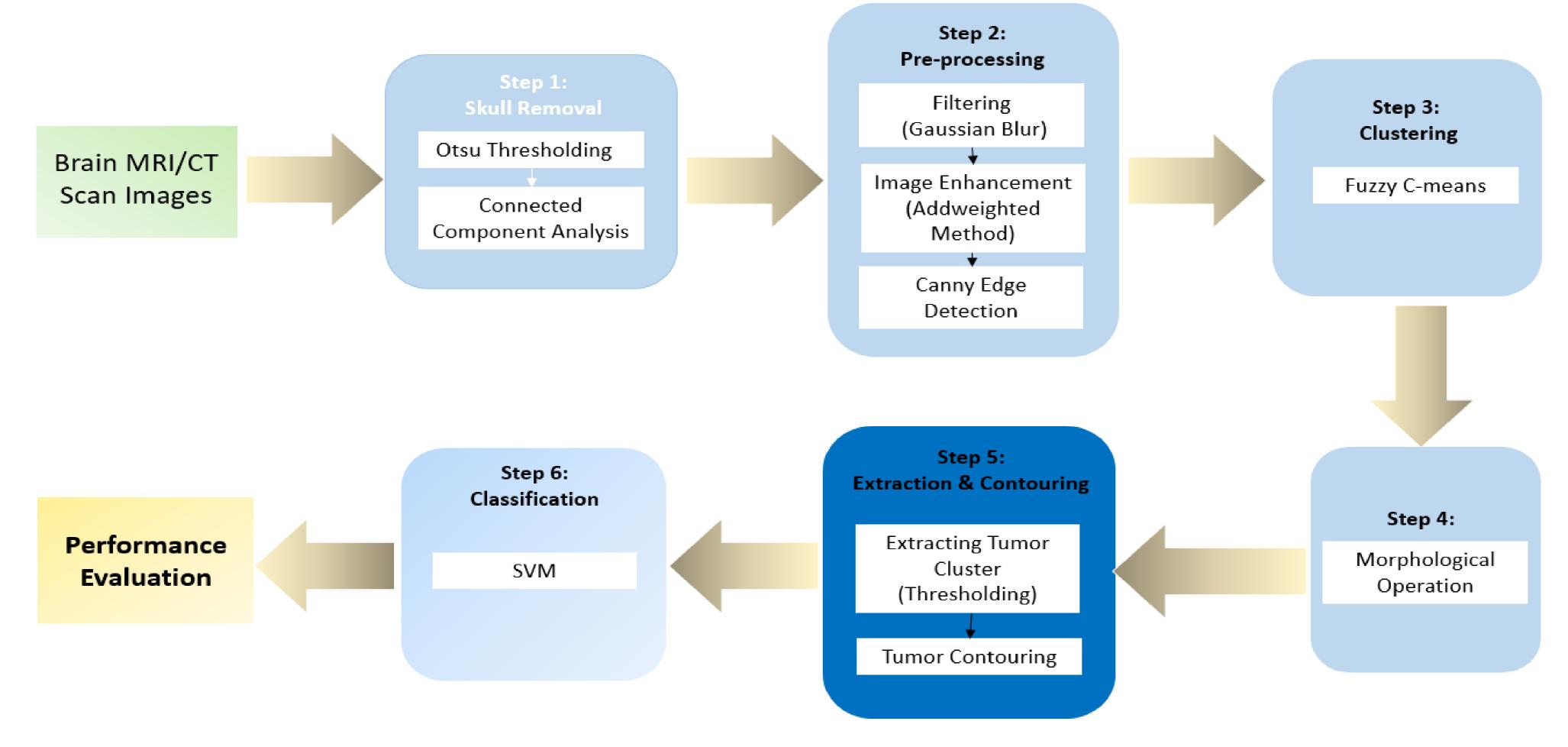


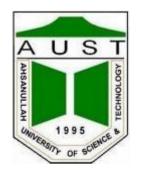
Morphological Operation





Tumor Contouring







Tumor Contouring

- Contours can be explained simply as a curve joining all the continuous points (along the boundary), having same color or intensity
- Used the cv2.findContours() method for finding the contours

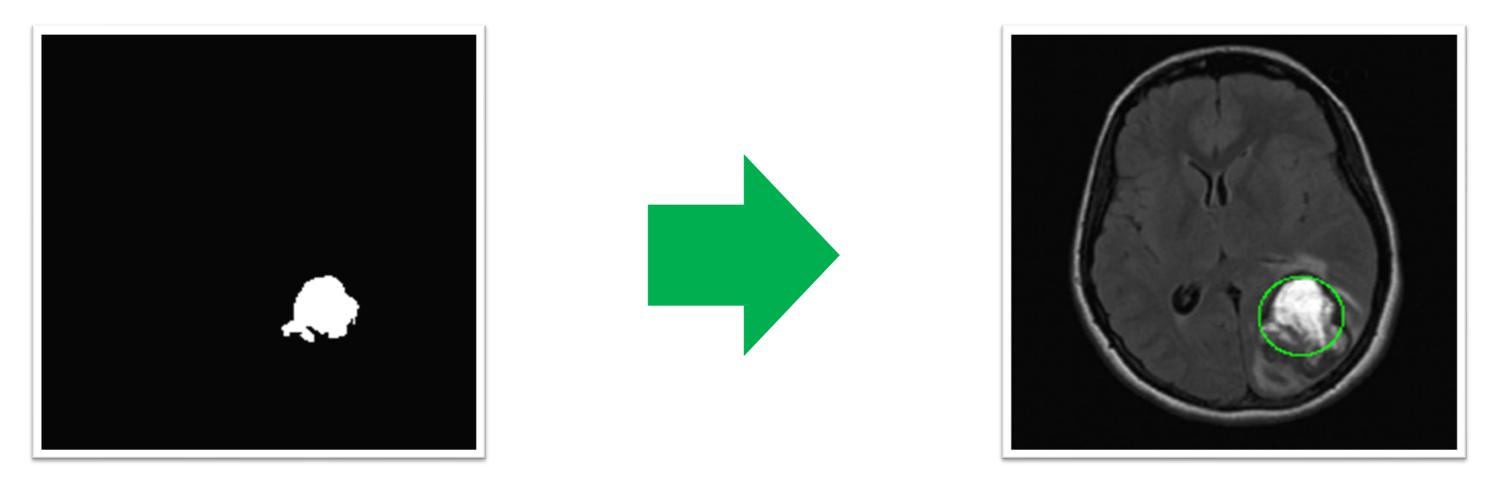


Fig 9.1: segmented MRI

Fig 9.2: contoured tumor MRI

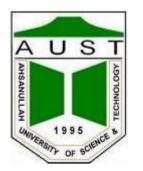


Fig 9: steps of tumor contouring

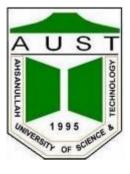


Traditional Classifier



We adopt six traditional Classifier

- K-Nearest Neighbor
- Logistic Regression
- Multilayer Perceptron
- Naïve Bayes
- Random Forest
- Support Vector Machine



Traditional Classifier

	TP	TN	FP	FN	Accuracy
K-Nearest Neighnour	56	3	4	3	89.39
Logistic Regression	56	2	5	3	87.88
Multilayer Perception	59	0	7	0	89.39
Naïve Bayes	47	5	2	12	78.79
Random Forest	58	1	6	1	89.39

Table I: confusion metrics of the classifiers

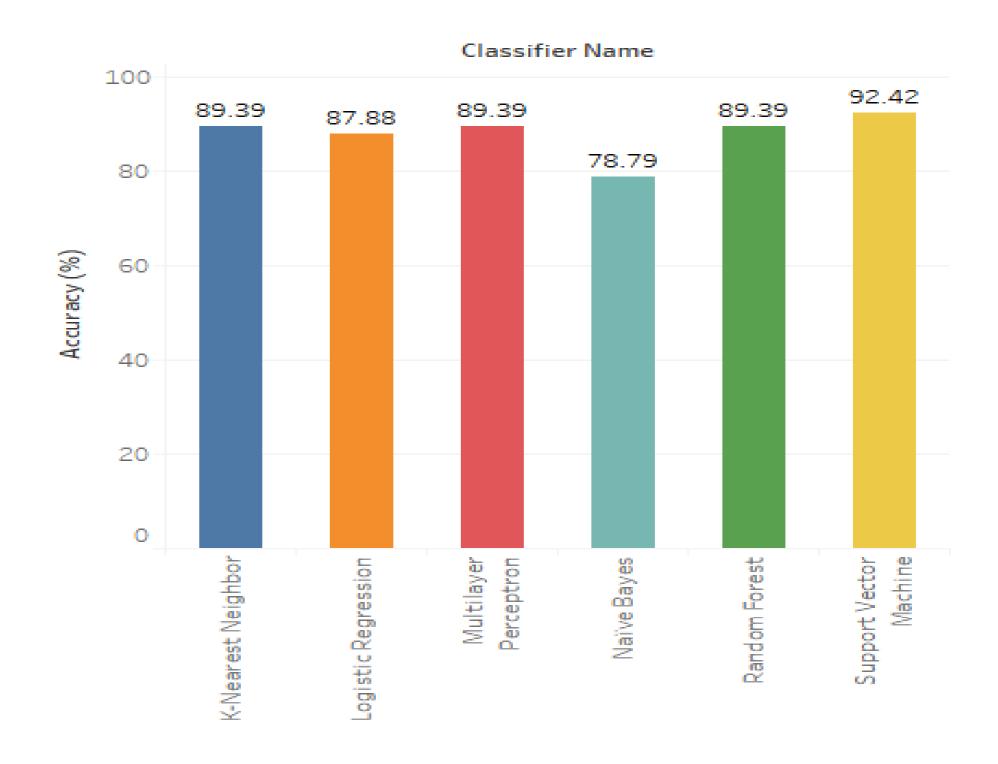
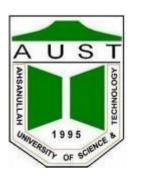


Fig 10: accuracy of the classifiers

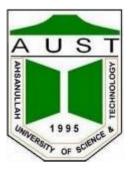


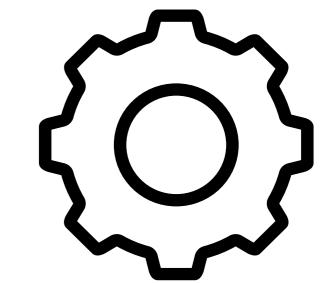


Traditional Classifier

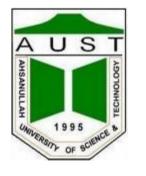
Classifier Name	Dice Score	Jaccard Index	Precision	Recall
K-Nearest Neighnour	0. 941	0.889	0.933	0.949
Logistic Regression	0.933	0.875	0.918	0.949
Multilayer Perception	0.944	0.894	0.894	1.000
Naïve Bayes	0.870	0.770	0.959	0.797
Random Forest	0.943	0.892	0.903	0.983
SVM	0.959	0.921	0.935	0.983

Table II: Performance Metrics of the classifiers





METHODOLOGY (CNN)



A Five-Layer CNN developed for tumor detection

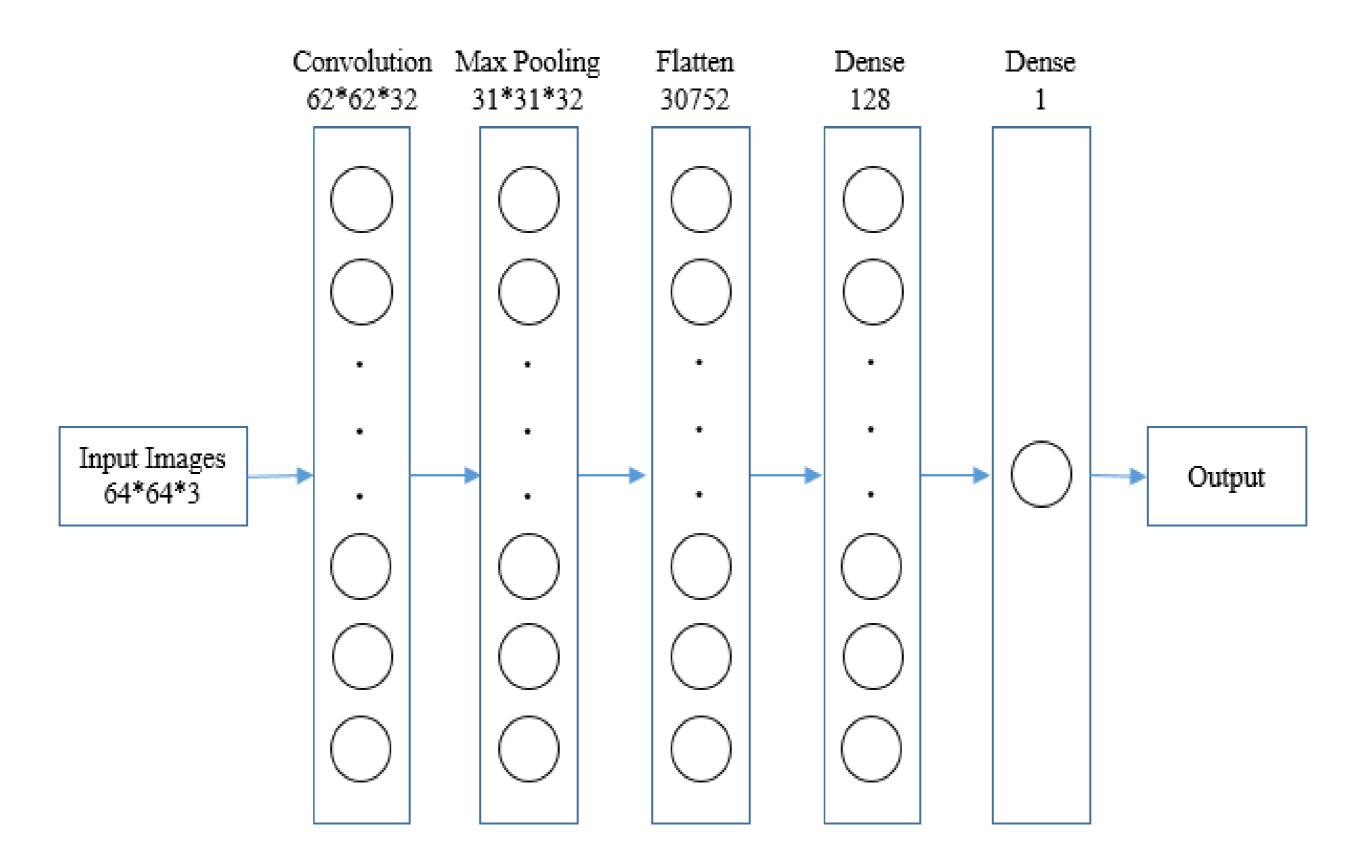
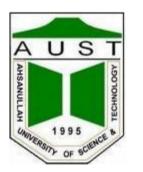


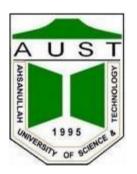
Fig 11: Proposed Methodology for tumor detection using 5-Layer Convolutional Neural Network





Convolution Layer

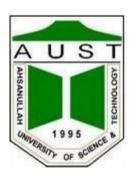
- The Beginning Layer
- Converting all the images into 64*64*3 homogeneous dimension
- Convolutional kernel of 32 convolutional filters of size 3*3 with the support of 3 tensor channels
- Activation function: ReLU





Max Pooling Layer

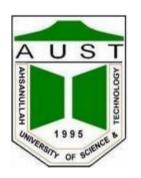
- Because of overfitting Max Pooling layer was introduced
- MaxPooling2D for the model
- Runs on 31*31*32 dimension
- \bigcirc Pool size is (2, 2)
- Output: Pooled feature map





Flatten

- Pooled feature map is work as the input
- Transformed the whole matrix into a single column vector
- Fed to the neural network for processing





Fully Connected Layers

- Two fully connected layers were employed Dense-1 and Dense-2 represented the dense layer
- The single obtained vector goes as an input
- O Dense function was applied in Keras
- 2 128 nodes in the hidden layer
- \bigcirc For better Convergence ReLU and sigmoid function is used as an Activation function in the 1st and 2nd dense layer respecticely

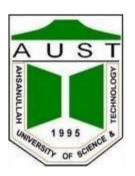




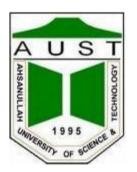
圖 Workflow of the Model

Complete workflow is divided into 7 steps

Working Flow Devised for Proposed Methodology

- 1. Load the input dataset
- 2. Adding a Convolution Layer with 32 convolutional filter
- 3. Passing the Convolutional kernel into the Max Pooling layer
- 4. Pooled feature map is used to get the single column vector
- 5. Processing of the vector in dense layer with 128 nodes
- 6. Final dense layer applying Sigmoid as the Activation function
- 7. Validation stage and Performance evaluation

Fig 12: working flow of the proposed CNN Model.





Hyper-parameter values

The hyper-parameters are divided into two stages-initialization and training

Stage	Hyper-parameter	Value	
Initialization	bias	Zeros	
initialization	Weights	glorot_uniform	
	Learning rate	0.001	
	beta_1	0.9	
Training	beta_2	0.999	
	epsilon	None	
	decay	0.0	
	amsgrad	False	
	epoch	10	
	Batch_size	32	
	steps_per_epoch	80	

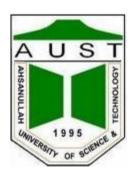


Table III: HYPER-PARAMETER VALUE OF CNN MODEL

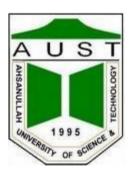


Evaluation Process

We devised an algorithm for the performance evaluation of our proposed model

```
Algorithm 1: Evaluation process of CNN model
 1 loadImage();
 2 dataAugmentation();
 3 splitData();
 4 loadModel();
 5 for each epoch in epochNumber do
      for each batch in batchSize do
          \hat{y} = \text{model(features)};
          loss = crossEntropy(y, \hat{y});
 8
          optimization(loss);
          accuracy();
10
          bestAccuracy = max(bestAccuracy, accuracy);
11
12 return
```

Fig 13: algorithm of the performance evaluation





Performance of the proposed model

Trained our model into two stage- 70:30 and 80:20 splitting ratio

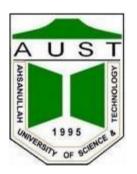
Accuracy: 97.87%

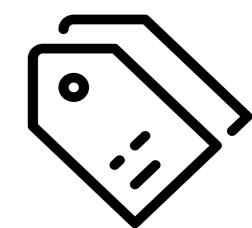
No	Training Image	Testing Image	Splitting Ratio	Accuracy (%)
1	152	65	70:30	92.98
2	174	43	80:20	97.87

Table IV: performance of the proposed CNN model



Fig 14: training and validation graph





FUTURE PLAN





Future Plan

- Work on 3D images
- Build our own dataset based on Bangladeshi patients
- Try to detect the grade and stage of the tumor
- Try to predict the location of the tumor from 3D images



THANK YOU!

Any Question!