Automatic Brain Cancer Detection using SVM Kernel Trick from MRI Imaging

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Abstract:- A brain tumour is an unchecked cell development that has the potential to spread to other tissues. It can be identified using Magnetic Resonance Imaging (MRI), a non-surgical method of organ research for the diagnosis of any illness associated with the symptoms. Tumours can be malignant or not, and they can also be hazardous or not to human life. A tumour can be classified as either benign or malignant, two separate categories. Benign tumours are seen as less hazardous or non-cancerous since they do not spread to other parts of the brain. It has distinct contours or limits that show the specific shade of the tumour, however malignant tumours are cancerous growths that may migrate to other parts of the brain on their own and are thus extremely hazardous. The malignant tumor's borders don't appear to be firm; rather, they have a faded look. Here, the suggested approach is capable of classifying both the tumour type and the diagnosis of a brain tumour. Here, the suggested system makes use of polynomial Support Vector Machine (SVM) to handle the impairments and identify the illness as necessary. The clustered cluster may be classified using SVM based on their patterns. The feature mapping of patterns might come from both diseased and normal cells. Drawing a hyper plane for non-linear data is challenging for other classification algorithms, however SVM can classify the data by converting it to linear data, which then makes it simple to design a hyper plane. It has been controlled using a kernel approach that allows for the mapping of high dimensional feature space. Utilising the polynomial feature of SVM, the optimisation problem may also be resolved. High degree of accuracy was perceived by the system in comparison to the prior model. 97.24, 93.94,

and 99.35 percent of accuracy, specificity, and sensitivity, respectively, were attained by the system.

Keywords:- Support Vector Machine, Brain Tumor, Segmentation, Malignant, Benign, MRI, Brain Cells.

I. INTRODUCTION

For the purpose of examining, locating, and treating brain tissues, magnetic resonance imaging is a common and safe method. The patient's life can be saved by receiving the right therapy if a brain tumour is discovered early. The accurate categorization and division of the tumour region should be made feasible by this suggested framework since the precise identification of tumours in MRI images becomes a crucial task to complete. The detection of a tumour from an MR image is also used in division and 3D reproduction. To avoid wasting time, professionals only conduct a limited amount of physical and visual research. The SVM Classifier may be used to restrict an abnormal cell mass in a slice of magnetic resonance imaging (MR) and divide the tumour cells to determine the size of the tumour that is present in that sectioned area. To demonstrate the kind of the tumour, the sectioned part's divided components will be produced using a fake brain organisation. These features will also be used in the Classification student application to compare the accuracy of several classifiers [1]. Perhaps the most precise and sensitive organ in the human body is the brain. Unimaginably fake brain tumours are to blame for the high fatality rate. The National Brain Tumour Society has shown that brain tumours are a highly harmful contaminant for people. It is a collection or mass of damaged brain cells. The cerebral cortex is located quite near to the brain. Brain tumours can either be nondestructive (harmless) or cancerous (compromising).



Fig. 1: Benign Tumor MRI Image

As dangerous or risky tumours grow, the pressure inside of your brain increases. Undermining tumours can be divided into two categories: primary and auxiliary tumours and benign tumours. The patient's health worsens as the undermining tumour quickly spreads to numerous brain structures. A mass of slowly proliferating brain cells is a brain tumour that is benign (without malignant development). It typically doesn't stand up and spread. The symptoms of a brain tumour vary depending on its size and location inside the brain. A few tumours that keep on growing progress silently at every turn. Not joking, persistent brain discomfort, seizures (sufficient), perpetual squeamishness, vomiting, and sluggishness are typical side effects. A situation where the phones mentioned in the danger of sickness are unexpectedly generated is known as an earlier condition or basic condition, also known as probable condition or fundamental condition. These conditions can eventually lead to illness if left untreated

[2].Destructive tumours that progress rapidly and cause mortality are referred to as risky tumours. Destructive tumours grow quickly, are envious, seek for a new location, and spread (metastasize) as opposed to cleanse tumours. A dangerous tumour is formed by damaged cells that are growing quickly. The magnetic field fragment utilised to finish radio recurrence to provide clear images of interior changes in the human body, including bones, sensitive tissues, and organs. The MRI approach is very sensible for identifying brain tumours. MRI images should make it possible to identify brain tumours. Image updating clusters utilised for clinical image diagnostics are employed in image processing to further improve picture quality. In order to identify and collect brain tumours, EDGE detection, histogram, and division limits are expected to play a significant role. The goal of this research is to identify various channels, detachment techniques, and assessments for the perception of brain tumours [2].



Fig. 2: Malignant Tumor MRI Image

II. RELATED WORKS

A. Literature Survey

Using CWT, DWT, and SVMs, Mircea Gurbin et al. [3] established a framework for classifying and detecting brain tumours. The suggested technique uses several levels of wavelets, with CWT being used to provide the high precision component. The CWT prevents divisions from lacking edges. The result demonstrates that SVMs with the proper information preparation arrangements can recognise unique and typical tumour sites and properly classify them as a benign tumour, malignant tumour, or normal brain. SVMs offer important computational advantages. The doctor needs this categorization to accurately describe the symptoms and recommend the best course of therapy. The obtained results demonstrate that, in comparison to DWT, CWT provides greater calculation. Even if the computation time is greater, using CWT is preferable if our main interests are perception, coordination, and include recognition. DWT is frequently more appropriate if we are interested in denoising, pressure, or rebuilding. The proper resolution of the challenges surrounding the localization and categorization

of brain tumours is advised using a mixed technique. The division and categorization of brain tumours suggested by T. A. Jemimma et al. [4] are carried out using the Water Shed Algorithm (WSA), Dynamic Angle Projection Pattern highlights, and these elements are arranged using CNN. The importance of the watershed division calculation effectively eliminates the tumour regions to enable competent DAPP highlight extraction. The histogram highlights are acquired after the DAPP eliminates the surface components from the fragmented tumour districts. These component vectors are raised to play a role in the classification process as it is carried out by the CNN classifier. For a successful diagnosis of a brain tumour, the division and categorization of the MRI brain picture are essential. The BRATS data set, which achieves greater dice score proficiency of 93.5% and awareness of 94.2%, is used to execute the trial outcomes. In further study, a few more varied factors may compare to obtain greater accuracy in the categorization and division of brain tumours. It may also be extended to differentiate between various tumour types, including fibromas, adenomas, and pancreatic tumours.



Fig. 3: Overview of the System [3]

The suggested technique by R.Lavanyadevi et al. [5] calls for precisely detecting the semantically significant entire areas in a picture. The doctor or radiologist might therefore identify hazard by connecting each and every pixel in the image together with which suggest a semantically important relevance. Brain pictures containing safe, risky, or usual images are removed from adjoining twofold models and dark level co-events. The PNN classifier is used in preparation mode to prepare the semantic components and the deleted highlights. Similar highlights from the test brain picture and mystery are subtracted in classification mode using preset models and a PNN classifier. When the test picture differs from any preparation image, the image can be recalled for creating set data. Regarding the relationship between PNN and CNN, PNN is seen to have several advantages. PNN benefits from gathering knowledge briefly because of reality. PNN can modify its learning gradually since it has a fast learning capability. The method suggested by Hein Tun Zaw et al. [6] can assist clinical staff members, such as specialists and radiologists, in analysing brain malignant development from MRI images, notably for GBM, which necessitates the finding of all potentially harmful spreading locations. The most severe entropy edge has been used in this method to identify brain tumours using Naive Bayes classification. This review makes use of the REMBRANDT data collection. The developed computation can accurately identify the tumour in all possible regions of the brain where the tumour may reside, including the projection of the outside world. The computation results in a 94% overall precision, an 81.25% identification rate on tumour pictures, and a 100% discovery rate on non-tumor images. A method for the straightforward division of the

brain tumour and identification of the tumour type was proposed by Ragib Shahariar Ayon et al. [7]. After preprocessing the picture using denoising and inclination correction, which was then handled to the handling stage as information image, brain tumour finding is completed. We used the spatial FCM to clip the probable tumour out of the brain MR picture. The tumour cut was then handled until the post-handling step, where it is channelled via a local region. The final graphic shows the projected tumour region separately. Similar features were used to create a variety of classifiers, and we chose the one that predicted the tumour type most accurately. After comparing and analysing various bunching and classification computations, we can state that the suggested method is superior to conventional strategies for tumour division and classification.

III. PROBLEM IDENTIFICATION

An implementation model based on FCM (Fuzzy C-Means Clustering) and CNN (Convolutional Neural Network) techniques was developed by L. Jagjeevan Rao et al. [8]. The algorithm in this case employs CNN as a classifier and FCM to extract the brain's characteristics and limitations. However, CNN is typically employed for feature extraction, whereas C-means clustering is utilised for classification. CNN is a convolutional neural network that has the capacity to obtain input features and enhance the features by using distinct filters, train the layers accordingly, and generate a model that can be able to diagnose or recognise the target object. CNN has not been designed for the classification process. SVM is regarded as the top classifier for diagnosing diseases.



Fig. 4: Graphical Representation of Result [8]

The FCM's drawback is that it takes a long time to converge the data, is more sensitive to noise, and has trouble with non-linear data. The traditional CNN model performs poorly when it comes to training and creating dense networks, which directly impacts the execution time. The accuracy of the system was 91%, which is somewhat less than what may be improved by utilising other techniques.

IV. PROPOSED WORK & IMPLEMENTATION

The brain tumour can be accurately diagnosed with the planned study. The system may also categorise tumours according to whether they are benign or malignant. The system's implementation has been broken down into three categories, including normal, malignant, and benign. When a tumour is benign, its borders appear to be solid or fixed in size and are thus less harmful than when a tumour is malignant, which is dangerous because it can spread to other parts of the brain. The malignant tumor's borders blur or fade, and it may be distinguished from the benign picture. The normal picture of the brain is thought to be a healthy brain image since it does not include any tumours. The suggested method in this case makes use of polynomial SVM, a kernel-based classifier that can handle non-linear data and produce results with more accuracy.



Fig. 5: Original Brain Tumor MRI Image

The original MRI picture of the brain tumour is shown in Fig. 5. The white portion of the image is thought to be the tumour portion of the image, and according to the theory and real idea, it belongs to the malignant class. Figure 6 depicts the original image that has been smoothed down so that clusters may be produced more readily for segmentation. In the segmentation process, certain clusters are separated into various groups based on the threshold value or colour of the picture, which is then classified using a classifier to get the desired results.

A. Smoothening



Fig. 6: Region Smoothening the Brain Tumor MRI Image

B. Segmentation

Picture segmentation is a technique that divides a digital picture into several subgroups known as Image segments. This approach helps to reduce the complexity of the image to make further handling or inspection of the image easier. Simply said, segmentation is the process of giving pixels names. A common name is assigned to each picture component or pixel that belongs to the same class. Take, for instance, a situation where a picture is required to be provided as part of a protest finding. The identification can be inputted with a location selected by a segmentation computation rather than processing the full picture. This will prevent the identify from processing the complete image, hence cutting down on guessing time.



Fig. 7: Brain Tumor Segmentation

The segmentation of a brain tumour picture is shown in Fig. 7, where the red portion depicts a cluster of damaged cells and the remaining portions are all thought of as the image's background that must be removed later for better categorization. The suggested system's flowchart, shown in Fig. 8, initially loads the dataset picture as input data. The pre-processing module was then started in order to improve the photos' visibility. One of them is the histogram, which is in charge of balancing the system's brightness and contrast. Once visibility increases, the image is smoothed to improve lesion visibility before features are extracted and feature selection, after which SVM classification can be started to categorise the data points. The system then determines the retrieved lesion's entropy. It determines the lesion's density for subsequent comparison with the cutoff value. Threshold value is the relative value at which a choice can be made. There are three categories of brain tumours; the first is malignant; as a result, if entropy exceeds and is equivalent to the higher threshold value, it will be regarded as malignant cancer. The second kind is benign; a healthy brain imaging can be taken into consideration if entropy is larger than the lower threshold value but less than the higher threshold value. As a result, the method also assigns a grade to each brain tumour according to its density.



Fig. 8: Flow Chart of Proposed System



Fig. 9: Polynomial SVM Classification [9]

Polynomial-SVM Algorithm (Kernel Trick)

Input: 2-D Image Matrix Output: Entropy Step 1: Input 2-D Image Matrix Step 2: Convert Matrix to Gray Levels Step 3: Apply Histogram Equalization

 $P_n = \frac{number \ of \ Pixel \ Intensity \ n}{Total \ number \ of \ pixels} \ n = 0, 1 \dots L - 1$

Where *Pn* is the affected pixel value after histogram equalization. Step 4: Apply Gaussian Filter for Smoothening Step 5: Collect data points

 $y = w_0 + w_1 x_1 + w_2 x_2 + w_3 x_3 + w_4 x_4 \dots \dots$ $= w_0 + \sum_{i=1}^m w_i x_i$

$$w_i = w_0, w_1, w_2 \dots \dots \dots w_m$$

Where w_i is the vector, b is the bias and x is the variable Step 6: Separate the data points by hyperplane

$$\vec{w}.\vec{x} - b = 1$$
$$\vec{w}.\vec{x} - b = 0$$
$$\vec{w}.\vec{x} - b = -1$$

where \vec{w} is the normal vector of the hyperplane Step 7: Classify Datapoints Step 8: Compute Entropy Step 9: if $E \ge T_1$ then Malignant; elseif $E < T_1 \&\& E > T_2$ then Benign; else Normal; end else end if Step 10: End

V. RESULT ANALYSIS

A. Result Simulation

Four metrics—True Positive (TP), False Positive (FP), True Negative (TN), and False Negative (FN)—are used to measure the success of an experiment. True Positive refers to cases in which a system correctly identified an image as either malignant or benign, whereas True Negative refers to situations in which a system incorrectly identified an image as either benign or malignant. False Positive means when a system diagnoses a normal-class picture as either malignant or benign, while False Negative means when a system diagnoses a malignant or benign image as normal. In the Kaggle benchmark, there are a total of 76 testing photos from the benign class, 89 from the malignant class, and 98 from the normal class.

Sensitivity =
$$\frac{TP}{TP + FN} * 100 \%$$

Specificity = $\frac{TN}{FP + TN} * 100 \%$

Precision = $\frac{TP}{TP + FP} * 100 \%$ Negative Prediction Rate = $\frac{TN}{FN + TN} * 100 \%$ False Positive Rate = $\frac{FP}{FP + TN} * 100 \%$ False Negative Rate = $\frac{FN}{FN + TP} * 100 \%$ Accuracy = $\frac{TP + TN}{TP + FP + TN + FN} * 100 \%$ F1 = $\frac{2TP}{2TP + FP + FN} * 100 \%$

Recall =
$$\frac{\text{TP}}{\text{FN} + \text{TP}} * 100 \%$$

						Features	
		1100				Contrast	0.642286
R	R	K	7 -			Correlation	0.860832
200			1.3			Cluster Prominence	106.939
	Y		y i		9	Cluster Shade	0.957809
				1	1	Dissimilarity	0.260273
Input Image		Smoo	thening	Segmentation		Energy	0.242085
Malignant	.94 jpg					Entropy	1.67639
	Result Anallysis		Classification		Homogeneity	0.923078	
Total True Positive	154	Total False Positive	6	SVM Cla	ssification	Maximum Probability	0.328709
Total Trup Nonation	93	Total False Negative	1	Charifastas	Dent	Variance	35.0444
roidi nige ivegalive				Lassincation	Result	variance	30.0444
Total Total Class	254	Obtained Accuracy	04.4992	Mafarant	Tumor Datacted		

Fig. 10: Proposed API

SEN: Sensitivity, SPE: Specificity, P: Precision, NPV: Negative Predictive Value, FPR: False Positive Rate, FNR: False Negative Rate, ACC: Accuracy, F1: F1 Score and R: Recall.

Table 1: Result Outcomes (Confusion Matrix)				
Terms & Parameters	Proposed			
Total Testing Class	254			
TP	154			
TN	93			
FP	6			
FN	1			
SEN in %	99.35			
SPE in %	93.94			
P in %	96.25			
NPV in %	98.94			
FPR in %	6.06			
FNR in %	0.65			
ACC in %	97.24			
F1 Score in %	97.78			
R in %	99.35			



Graph 1: Result Analysis



VI. CONCLUSION & FUTURE SCOPE

The brain tumour may be identified and classified according to whether it is malignant, benign, or normal utilising brain tumour classification using MRI imaging using polynomial SVM. The system employs histogram equalisation to change the image's contrast for improved image visibility. In order to make impairments more visible and create a distinct cluster that can aid in segmentation and classification, the system additionally applies a smoothening filter. For final classification, the system employs Polynomial SVM, which creates two separate clusters: one is composed of aberrant or lesion cells, while the other is composed of normal cells, which may be thought of as the image's backdrop or noise. Comparing the system's accuracy to that of CNN, BPNN, and KNN classifiers from earlier systems, the system attained a higher level. Because accuracy is crucial in the field of medical science, hybrid classifiers or models may be used in the future to improve accuracy, precision, specificity, and sensitivity. Because brain tumours are a potentially fatal condition, it is essential to achieve the highest degree of accuracy with the fewest possible false positive rates.

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