



# JOURNAL OF BIOMEDICAL ENGINEERING AND MEDICAL IMAGING



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# Parasites as Bioindicator for Health Status and Environmental quality of Freshwater Fish species in Ekiti State, Nigeria

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## ABSTRACT

The prevalence of parasites occurring on and in the internal organs of fish species was studied. Fish species (*Oreochromis niloticus*, *Clarias gariepinus*, *Tilapia zilli*) were randomly obtained from fishermen landing at the jetties of Ado, Ero, Ogbese, Ikun and Egbe dams, in Ekiti state. Samples were dissected; smears for the identification of ectoparasites were taken directly from the gills, operculum and skin. The parasites were excised, identified and counted. Nematode parasite with the total sum of 164(29.6%) was the most abundant parasite occurring in all the dams. This was followed by *Trichodina* sp (Ciliophora) with total abundance value of 147(26.5%) in all the dams. *Acanthocephalan*, *Gyrodactylus* sp, *Dactylogyrus* sp, *Diphyllbothrium latum* and *Clinostomum* sp have abundance values of 26 (4.7%), 27(4.9%), 21(3.8%), 146 (26.4%) and 23(4.2%) respectively with *Dactylogyrus* sp (Monogenea) being least abundant, 21(3.8%), in all the dams. The overall parasitic prevalence from the different dams showed that Ado dam had the highest parasitic infection with 27.3% which suggests that Ado dam has the lowest water quality. There is need for constant surveillance in rivers and reservoirs to detect early signs of parasitic infection. Viable preventive measures against fish parasitism in Nigerian freshwater bodies are highly recommended.

**Keywords:** Bioindicators, Dam, Fish, Parasites, Prevalence Freshwater,

## 1 Introduction

Parasites can be used as effective monitoring tools in environmental impact studies as they are able to accumulate certain pollutants at levels higher than those of their ambient environment [1]. Fish parasites serve as biological indicator to illustrate the ecology of their infected hosts. They have been connected to anthropogenic impact and environmental changes in freshwater habitat. Water pollution induces pathological changes in fish, thus fish species can be used as indicator of exposure to contaminants [2]. Fish is considered as one of the most significant indicators because they have long lifespan, which can be exploited in bio-accumulation studies. They occur at different trophic level in an aquatic food web (herbivores, omnivores, carnivores), they also occur ubiquitously, almost in all water, even polluted water. Fish can serve as definitive, intermediate or paratenic host in the life cycle of many species of metazoans,

protozoans and crustacean parasites. Also, due to the fish species diversity, there is an increasing interest in using parasites as bioindicator of their fish host life conditions.

Environmental stressors such as wastewater or industrial pollutants can result in an increase in fish parasites due to a decrease in immunological defenses and a lesser resistance to infections. Water quality and season are the most important factors affecting the prevalence of parasites [3]. Pollutants might promote parasitism increasingly in aquatic animals; especially fish [4, 5]. Metazoans, which are normally ectoparasites, are in constant contact with water, suggesting that poor water quality may adversely affect their diversity to a greater extent [6]

Parasites can be used to detect both heavy metals pollution and eutrophication in water [7]. Parasites are potential indicators of environmental quality due to the variety of ways in which they respond to anthropogenic pollution. They provide valuable information about the chemical state of their environment through their ability to concentrate environmental toxins within their tissues.

Conditions that predispose fish species to parasitic infections include reduced oxygen content of water, increase in organic matter in water, poor environmental condition. The effect of parasitic infection on fish species is influenced to a large extent by the type and number of parasite present.. The pathological conditions of parasites in fish includes anorexia (loss of appetite), resting at the bottom of water, changes in skin color, pale gills, increase in mucus production. Parasites of fishes include trematodes (flukes), cestodes (tapeworm), nematodes (roundworms), Acanthocephalan (thorny headed worms) and protozoans out of which trematodes and nematodes are the commonest. Knowledge of biology of parasite and its host, host-parasite relationship, and the environment can help to detect environmental change, particularly long living species (some digeneans, trematodes and cestodes life cycle stages), provide information on the seasonal migration of their host and migration habits of different age groups(feeding area/ spawning area). However, there can be a change in their abundance in the host of some of their life cycle stages which can be as a result of disappearance of their intermediate hosts, provoking disappearance of some parasite species under polluted conditions. The occurrence of heteroxenous parasites in an area affected by pollution can be related to the number of intermediate hosts at the studied sites [8].

Water quality and season are the most important factors affecting the prevalence and prevalence of parasites [3]. Pollutants might promote parasitism increasingly in aquatic animals; especially fish [4, 5]. Certain parasites are able to accumulate heavy metals than their fish host's tissues or the environment. Due to this ability, they can be sensitive bio-indicator of environmental pollutants. Fish muscles and various mussel species are commonly used as bio-indicators. Studies have shown that there is specie-specific preference for metals in parasites' propensity to accumulate different pollutants, where certain parasite specie accumulates certain metals more intensively than other parasite species and vice-versa.

Many indices are being used in aquatic environmental studies to determine the quality of water as well as fauna inhabiting the environment. Also, parasitic index in aquatic habitat can be linked with fish pathology components and other measures of fish health to assess the health status and environmental quality of fish. The use of fish parasites as bioindicators in aquatic environment has not been fully exploited like other indices hence the need for this study.



The objective of this study is to determine the prevalence of parasites found in different water bodies in Ekiti State and use the fish parasites metrics as a bio-indicator in determining the water quality.

## 2 Materials and Method

### 2.1 The Study Area

The study was carried out using fish samples collected from Ado Ekiti dam, Ero dam, Egbe dam, Ogbese River, Ikun-Ekiti dam. Ado-Ekiti dam was constructed on Ureje river in Ado Ekiti. It is located on latitude 7°37' North and Longitude 5°36' East of the equator. Ero dam is an earth filled embankment with a length of 662m and an impoundment area of 45km, located in Ikun Ekiti between Latitude 7°15' - 8°5' and Longitude 4°45' – 5°45'. Egbe dam which originated from Kwara State and flows north to south through Ode-Ekiti to Egbe- Ekiti, located at 7°36' North and Longitude 5°36' east of the equator.

### 2.2 Sample Collection:

The fish species collected and identified according to Idodo- Umeh (2003). The fish species were identified as *Clarias gariepinus*, *Tilapia zilli*, and *Oreochromis niloticus*. They were bought for 3 months from local fishermen at the water sites as soon as they landed and transported to the laboratory using big plastic container, making sure the samples were still alive. The smears for the identification of ectoparasites were taken directly from the gills, operculum and skin while dissections were done to observe the intestine for possible detection of endoparasites.

### 2.3 Experimental procedures

External surface of the fish was examined using a hand lens for ectoparasites species. The smears from the gill surface and operculum were stained using silver nitrate impregnation. They were rinsed and covered with 5% silver nitrate (AgNO<sub>3</sub>) solution and impregnated for 30 minutes in the dark. The silver nitrate was removed and the slides were covered with distilled water, exposed to light for 40-50 min and dried after exposure.

For identification purposes, Nematoda was dehydrated in a graded ethanol series and transferred to 100% glycerine (Riemann, 1988). Digenea, Monogenea and Cestoda were stained with acetic carmine, dehydrated, cleared with eugenol and mounted in Canada balsam (Palm, 2004). The identification of the parasites species on the slides were done or carried out under light based on the shape and size of the sclerotized part when they are mounted using Canada balsam and stained with Acetic carmine for monogeneans and cestodes.

For Endoparasites, the fish samples were laid on a dissecting board, the mouth, fins and dorsal cavity well clamped down with entomological pins. The dorsal side was symmetrically opened with the aid of surgical blades to show the alimentary canal. The intestine, stomach, liver, heart, gall bladder were carefully excised for parasites examination. Gonads were excised carefully using forceps into the petridish filled with 10% normal saline for 20minutes. Parasites that appear on the surface of the normal saline in the petri-dish were extracted using a dropper and was placed on a microscope slide which was viewed under a compound microscope with magnification (x40). The following parasites were identified: Nematodes, Cestodes, Trematodes, Acanthocephalan.

## 2.4 Statistical Analysis

Analysis of variance was used to determine the significance of each parasite in each dam. Relative abundance was calculated as: species of a parasites / Total specimen x100. Both descriptive and inferential statistics were employed to analyze results.

## 3 Results

### 3.1 Parasite abundance

Nematode parasite with the total sum of 164(29.6%) was the most abundant parasite occurring in all the dams. This was followed by *Trichodina* sp (Ciliophora) with total abundance value of 147(26.5%) in all the dams. *Acanthocephalan*, *Gyrodactylus* sp, *Dactylogyrus* sp, *Diphylobothrium latum* and *Clinostomum* sp have abundance values of 26 (4.7%), 27(4.9%), 21(3.8%), 146 (26.4%) and 23(4.2%) respectively with *Dactylogyrus* sp (Monogenea) being least abundant, 21(3.8%), in all the dams as shown in Table 1.

**Table 1: Represents the percentage abundance of each parasite in the study areas.**

Parasite/Study Area	Ikun Dam	Ado Dam	Ero Dam	Egbe Dam	Ogbese dam	Total
<i>Trichodina</i> sp (Ciliophora)	-	17.00 11.2%	49.00 53.8%	81.00 64.3%	-	147 26.5%
<i>Acanthocephalan</i>	-	26 17.2%	-	-	-	26 4.7%
<i>Gyrodactylus</i> sp (Monogenea)	-	3.00 2%	9.00 9.9%	15.00 11.9%	-	27 4.9%
<i>Dactylogyrus</i> sp (Monogenea)	-	5.00 3.3%	8.00 8.8%	8.00 6.3%	-	21 3.8%
<i>Clinostomum</i> sp (Trematoda)	-	4.00 2.6%	9.00 9.9%	10.00 7.9%	-	23 4.2%
<i>Diphylobothrium latum</i> (cestoda)	28 40%	22 14.6%	6.00 6.6%	3.00 2.4%	87 75%	146 26.4%
<i>Camallanus</i> sp (Nematoda)	42 60%	74 49%	10.00 11.0%	9.00 7.1%	29 25%	164 29.6%
Total	70 100%	151 100.0%	91.00 100.0%	126.00 100.0%	116 100%	554 100%

The overall parasitic prevalence from the different dams showed that Ikun dam has the least parasitic infection with 12.6% of the total and Ado dam had the highest parasitic infection with 27.3% as shown in Table 2.

**Table 2: Percentage prevalence of parasites in the study areas.**

Parasite/Study Area	Ikun Dam	Ado Dam	Ero Dam	Egbe Dam	Ogbese dam	Total
<i>Trichodina</i> sp (Ciliophora)	-	17.00	49.00	81.00	-	147
<i>Acanthocephalan</i>	-	26	-	-	-	26
<i>Gyrodactylus</i> sp (Monogenea)	-	3.00	9.00	15.00	-	27
<i>Dactylogyrus</i> sp	-	5.00	8.00	8.00	-	21

(Monogenea)						
<i>Clinostomum</i> sp (Trematoda)	-	4.00	9.00	10.00	-	23
<i>Diphylobotrium latum</i> (cestoda)	28	22	6.00	3.00	87	146
<i>Camallanus</i> sp (Nematoda)	42	74	10.00	9.00	29	164
Total parasites/percentage	70 12.6%	151 27.3%	91.00 16.42%	126.00 22.7%	116 20.9%	554 100%

#### 4 Discussion

The result showed nematode (*Camallanus* sp) to be the most prevalent parasite with abundance value of 164 (29.6%) at all locations throughout the study period. The high rate of fish susceptibility to *Camallanus* sp. observed in this research is in agreement with the findings of [9]. [10] observed that intestinal parasites appears to be more sensitive bio-accumulators of heavy metals than their fish hosts, and may serve as excellent indicators of heavy metal pollution in relation to water quality. Thus, the presence of particular parasites in a host may tell us something about the stability of the ecosystem. The high rate of *Gyrodactylus* sp susceptibility reported in study was as a result of the compromised dermal tissue that the fish samples present. This follows [11] report on excessive mucus secretion, epithelial proliferation and dermal erosion of fin fish. It could be as a result of their high population density or environmental degradation of their habitat. [12] reported that in high population density along the coasts and significant anthropogenic stress in terms of population and environmental degradation, fish can describe the environmental condition of their habitat. According to [9] the cichlids harbor majority of infection which include adult digenea infecting different tissues; trematodes metacercariae of the family clinostomatidae encysting in the tissue and adult monogenea of the families Dactylogyridae and Gynodactylidae infecting the gills and the skin of tilapia species. This shows that it could be easy to identify fish samples with compromised dermal tissues that may bear *Gyrodactylus* sp and avoid them during market purchase in order to prevent human transmission of possible infection. This research findings show the presence of *Clinostomum* sp. and *Diphylobotriumlatum* in significant quantity. This is in agreement with [13] and [9].

Nematode parasite (*Camallanus* sp) was found in large quantity 164 (29.6%) in the fish host across all the study areas, as earlier observed by [14] who reported nematodes as the most common parasites infecting 18.6% of fish population. Ado Dam has the highest population of parasites with percentage abundance value of 27.3% compared to the other dams, this can further show the level of environmental quality of water in the dam [7]. It was also observed in this study that adult fishes are more susceptible to parasitic infections than juveniles. This corroborates the findings that the longer and bigger the fish, the greater the susceptibility to parasitic infection.

#### 5 Conclusion and Recommendation

It is recommended that there should be further studies of wider scope on other fish species found in many Nigerian diets. Also in order to control parasites in fish culture system, it is recommended that lining of the ponds should be done before stocking. There should be constant surveillance and observation of fish behavior in rivers and reservoirs to detect early signs of parasitic infection. Also, weeds and various types

of vegetation should be removed to ensure digenean flukes intermediate hosts' control. Fish density control is recommended as well as selective cropping and prompt removal of infected fish to avoid rapid spread of infection.

This research showed that Ado dam has the lowest water quality as compared to the other dams. The economic impact of these parasites on fish as an article of trade would lead to better policy decision on how to protect Nigeria's aquatic resources. A comprehensive knowledge of the biology of host parasite relationship is therefore, imperative.

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# Brain Tumor Segmentation through Region-based, Supervised and Unsupervised Learning Methods: A Literature Survey

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## ABSTRACT

Image segmentation is one of the most trending fields in the domain of digital image processing. For years, researchers have shown a remarkable progress in the field of Image Segmentation, precisely, for brain tumor extraction from various medical imaging modalities including X-Ray, Computed Tomography and most importantly, Magnetic Resonance Images (MRI). In these medical imaging modalities, accurate and reliable brain tumor segmentation is extremely imperative to perform safe diagnose, healthy treatment planning and consistent treatment outcome evaluation in order to understand and cure the complexities of chronic diseases such as Cancer. This paper presents various image processing techniques that are currently being used for brain tumor extraction from medical images. Though some great work has been done in this domain but none of the techniques has been widely accepted to be brought into practice in real time clinical analysis. The paper concludes with proposing some solutions that would aid in refining the results of the techniques which will lead to clinical acceptance of these computer aided methods.

**Keywords:** Image Analysis, Image Segmentation, Brain Tumor Detection Region Based, Supervised Learning, Unsupervised Learning, Clustering, Watershed Segmentation, Convolutional Neural Networks, SVM, K-Means Clustering, MRI, CT scan

## 1 Introduction

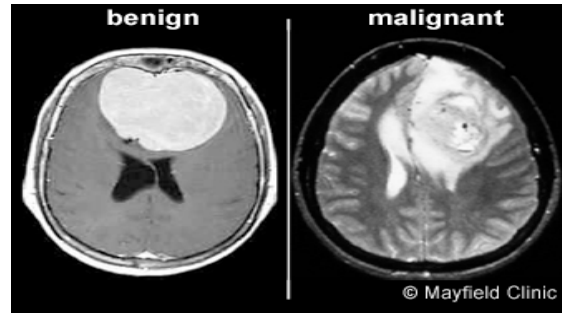
The brain is undeniably considered as the most vital organ in the human body. Although, it only weighs about 3 pounds but works as a central processing unit of human body. It controls and commands the voluntary as well as involuntary actions and reactions in a way that allows us to think and feel, retain memorable moments and feelings. Most importantly, this is the indispensable organ that completes us and turns us into a human [1]. Human brain has been under study since ages due to its complex structure and natural placement inside the skull which hinders the perfect and ultimate diagnosis of the diseases. Naturally, human brain is tightly safeguarded but prone to harms and can be affected by several fatal diseases [1]-[2]. An unwanted mass which is result of irregular growth of tissues (figure 1) - benign and malignant tumors inside the human skull, in some cases, surrounds the brain is known as Brain Tumor. Many different types of brain tumors have been observed and are categorized based on their characteristics. Depending upon their characteristics, brain tumors are categorized into two broad

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classifications: noncancerous (benign) or cancerous (malignant) types. Figure 1 shows MRI images of the brain for both of the aforementioned categories. Brain tumors mainly originate inside brain (primary brain tumors), or it can invade the brain by the means of spread of cancer through other parts of the body (metastatic or secondary brain tumors) [2-4].



**Figure 1: Benign and Malignant Tumors.**

However, the growth or spread of brain tumor can be analyzed by the various grades [4] – [5]:

- **Grade I:**

These are “low grade tumors”, which grow at a quite slower rate and exhibit benign morphology at external observation. These tumors look pretty similar as normal cells and have quite lower recurrence of tumor returning or occurrence of spread to other parts of the body. These kinds of tumors are less serious which can be treated with minimal surgery.

- **Grade II:**

These are also regarded as “low grade tumors”, which grow at a quite slower rate, look slightly abnormal and exhibit benign morphology at external observation. However, once removed from the body these tumors can more likely grow again and spread to other parts of the body, if safety precautions are not taken accordingly.

- **Grade III:**

These are “high grade tumors”, which grow at a quite faster rate and exhibit malignant morphology at external observation. These tumors look like abnormal cells and have quite higher recurrence of tumor returning and spread to other parts of the body frequently. These kinds of tumors are serious which cannot be only treated with minimal surgery, thus require extensive treatment as well such as chemotherapy or radiotherapy etc.

- **Grade IV:**

These are also regarded as “high grade tumors”, which grow at a quite faster and frequent rate, look abnormal and exhibit malignant morphology at external observation. However, once removed from the body these tumors still have higher probability to grow again and spread to other parts of the body, if safety precautions are not taken accordingly. These kinds of tumors are highly serious which require extensive treatment such as chemotherapy or radiotherapy etc.

For the treatment of brain tumor in any of the above grades, it must be detected timely and located accurately [4]-[7]. When this anomalous mass occurs inside the brain, it needs to be identified as soon as possible, and for this, various imaging modalities such as Magnetic Imaging Resonance (MRI) and Computed Tomography (CT) are widely used to detect the tumor. Among these medical imaging modalities, MRI is most widely used and highly preferred non-invasive technique in biomedical, radiology and medical imaging fields due to its capability to detect and visualize finer details in the internal structure of the body by generating three dimensional high resolution detailed anatomical images without the use of any damaging radiations. [8]-[10]. The brain MRI segmentation into several brain tissues such as gray matter (GM), white matter (WM) and cerebrospinal fluid (CSF) is highly essential for the diagnosis of various diseases. This technique is basically used to detect the detailed differences in the tissues in non-invasive fashion which have not been examined by other imaging techniques such as Computed Tomography [11].

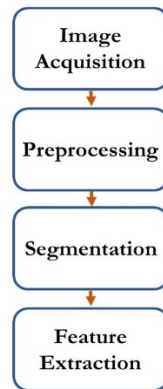
Once the MRI scanned images are obtained, the goal is to extract out the desired information in order to diagnose the tumor without any delay. One of the major problems in this whole process is separating the abnormal cells from the rest of the image content which is known as the process of segmentation. The manual segmentation is quite challenging as well as time consuming task due to complex structure of the brain and absence of well-defined boundaries among different brain tissues. Therefore, the accurate and reliable brain tumor segmentation is extremely important to perform safe diagnosis, healthy treatment planning and consistent treatment outcome evaluation in order to understand and cure the complexities of chronic diseases such as Cancer. Although the process of segmenting the desired region accurately is highly challenging and complicated but it has gained enormous importance and several studies have been conducted in improving the accuracy of this task [7][10][12].

Results obtained from the various image segmentation techniques are quite useful in obtaining features of segmented tumor region such as area, volume, perimeter, eccentricity, bounding box and tumor orientation [13]. Numerous research work has been done, and several algorithms are proposed with the goal of detecting the position and boundary of tumors automatically and eventually providing sufficient information to the clinicians, so they can carry out further diagnosis at their earliest. The study presented in this paper reviews the several prominent methods and techniques of automatic segmentation of brain tumor from the MRI images. The rest of the paper and its sections are arranged in the subsequent manner: The 2nd section presents a generic methodology adopted in the techniques implemented for brain tumor segmentation, followed by a detailed Literature survey in the 3rd section. Section V demonstrates the overall comparison and evaluation of the results followed by the final conclusions and future recommendations in the ending sections.

## 2 Methodology

The general methodology adopted in order to implement the techniques is portrayed in Figure 2. Each step of the methodology is modified according to a specific algorithm into consideration.





**Figure 2: A generalized workflow of image processing.**

- **Image Acquisition:**

To work with any kind of image processing task, it is essential to first acquire the image to apply the processing on.

- **Pre-processing:**

Generating images from various medical imaging techniques may incur unnecessary noise into the image. Hence, any MRI, CT Scan, Mammographic Image or etc., usually comes with a lot of noise. This noise can act as a hurdle when segmenting tumor region from the given input image. To eliminate this issue, an image is first preprocessed in order to remove unwanted outliers, and then sent for further processing. This preprocessing step may include techniques like noise removal, filter application, image enhancement, normalization, etc.

- **Segmentation:**

This is the most crucial and core step in applications like tumor detection. It is a simple phenomenon of dividing an image into different meaningful segments to further interpret them. Segmentation of an image can be achieved in numerous ways as already discussed in the literature.

- **Feature Extraction**

Once the image has been divided into segments, a post processing step is needed in order to sharpen any edges and blur any unwanted details. This step is called feature extraction where some of the features from the image are extracted for analysis. This will enhance the tumor region so that the area calculation or further operations can be applied on it for more effective results. Morphological operations, edge detection techniques or histogram equalization are most widely used feature extraction steps.

### **3 Segmentation Techniques**

MR Images contain a high amount of data. This makes the task of interpreting those scans laborious and tedious even for a radiologist and clinical imaging specialist. In addition, the interpretation results could be different depending upon the experience of the particular specialist [14]. Moreover, various imaging systems may introduce noise in the images, hence making it difficult to segment brain tumor and give an acceptable performance. Segmentation is important as it can help in calculation of quantitative measure of tumor affected part in the brain which is essential for treatment of patient and follow up of the disease.

The ultimate goal and foremost aim of large number of computer vision, image processing and machine learning based applications is to identify and extract the important patterns or vital features from the image data; using which as a model or reference an imperative description, valuable interpretation or fundamental understanding of the incident, situation or scene can be narrated by the machine for further detailed explanation [8][14][15]. Real time diagnosis of brain tumors and complex diseases from radiographs (as obtained through X-ray, CT scan, PET, MRI and bone-scan etc) is one of the most important but highly challenging task due to high-time consumption and distortion among these images. In order to assist clinicians in decision making, the main focus of the various research groups is to present reliable algorithms which can perform extraordinarily towards accurate segmentation and thus lead to construct a robust as well as to ensure a safe diagnosis system. Table 1 shows a few, most widely used techniques developed by the digital image processing community for image segmentation.

Segmentation Types	Techniques
Region Based	Histogram Threshold, Watershed etc.
Supervised Learning (classifiers)	Convolutional Neural Network, Support Vector Machines etc.
Unsupervised Learning	Fuzzy C-means, K-means, etc.

- **Region Based Technique**

Region-based segmentation is a way of determining and locating the desired region correctly. Simply, it combines the individual pixels in an input image to sets of pixels called regions that might correspond to an object or a meaningful part of one. Tanuja and Subhangi [16] came up with a system to segment tumors from the given MRI based on the similarity among the pixels. This idea was called region growing segmentation. It is named so as the basic idea was to define a seed pixel and move to neighboring pixels, grouping the pixels with the similar attributes. The experiments and analysis depict that this method was fast and accurate.

The two most commonly used region-based techniques include Histogram Thresholding and Watershed segmentation. The techniques are discussed in the following section.

- **Histogram Thresholding**

Histogram thresholding is widely used region-based segmentation approach that can be used to segment brain tumors among MRI images. In this approach, a threshold is established that could be used to segment the interior area or surface from other organs in the radiographic image dataset [16]-[19]. In principle, in thresholding approach every single pixel is compared to that particular established threshold. If the pixel depicts a value higher than its corresponding threshold, then the pixel is assigned a white color and called “foreground”, conversely if the pixel retrieves a value lower or equal to its corresponding threshold, a black color is assigned and titled as “background”. Most of the prevailing thresholding approaches are based on bi-level practices, in which an image is categorized into object segments and background. However, MR images, due to their high resolution and distortion issues make these methods non-applicable and difficult to perform smoothly. Consequently, it leads to loss of important information and crucial features from the images which compromise and hampers the overall diagnosis system and

may mislead the medical physicists and radiologists to perform related clinical tasks and decision making [20].

On the other hand, in the watershed approach following three major steps are followed: (i) Initially, the brain is classified into two major halves around its central axis, and finally the histogram is drawn for each of corresponding part, which helps to detect the contagious lesions of the brain. (ii) Secondly, the two histograms are compared and the corresponding threshold point values of their histograms are calculated for further evaluation. (iii) Finally, the physical dimensions and morphological features of tumors are evaluated by cropping the detected output image along its corresponding contours.

- **Watershed Segmentation**

In region-based segmentation, the mathematical morphological watershed is another most commonly used segmentation approach which segments the organs and tumor lesions based on the watershed ridge lines present in an image. In MR images or even other form of radiographs related to brain tumors, the brain tumors depict quite higher intensity and density and sometimes distortion effects, which makes it extremely difficult to segment the lesions precisely. In such cases, the watershed segmentation is one of the best and potential approach to not only segment and classify the tumors and high density/intensity tissues but also to overcome the distortion problems. Besides, the watershed approach also suffers from over-segmentation and under segmentation constraints because of distortion, noise and various other abnormalities in the radiographs or medical images. To overcome the over-segmentation limitations, the researchers have introduced a method based on the concept of controlled markers in literature. The proposed method deals with the watershed ridge lines and the catchment basins present in an image and consider them a surface wherever the light pixels are low. At first the medical images are converted into gray-scale images and some pre-processing is performed to remove noise followed by marker selection and segmentation etc. In principle, a gradient magnitude is calculated and finally the internal and external markers are calculated to distinguish the foreground of adjacent objects and extract the information from the medical images.

Watershed procedure is widely used for segmentation due to its implementation ease and space and time efficiency. It was first proposed by Digabe and Lantuejou[25], and improved by Beucher and Lantuejou [26]. The major drawback that occurs with this type of segmentation is a phenomenon called over segmentation [27]. In order to overcome this limitation, a marker-based approach was used in [28][29][30] where a brain atlas was used to detect the internal and external markers for division of foreground and background pixels. Different combination of features including color, texture, edge, orientation etc., were tested. All of them except for the combination of color and orientation, gave acceptable and good results.

Swe Zin Oo [23] used the concept of skull stripping, that is, removing any non-brain tissues from the brain image. He coupled this preprocessing technique with the conventional watershed segmentation approach. The study was not only aimed at detecting the tumor region, but also assisted in calculating the volume of the tumor resident in the brain. This volume could be essential in figuring out the grade of the tumor.

Recently, Subudhi et. al., developed an automated watershed-based lesion segmentation (WLSA) method for efficiently delineating the infarct lesions in MRI images of the brain strokes. The method incorporated

watershed transformation with guided filter through relative fuzzy connectedness (RFC) to distinguish the lesion boundaries reliably and appropriately. The suggested method achieved better results in delineating the lesions and improved the accuracy of detection of ischemic stroke lesions. The proposed technique could be extensively used for early, precise and accurate delineation of stroke lesions in clinical settings [70][71].

- **Segmentation with Classifiers (Supervised Learning)**

With the on-going research focus on automating the time-consuming segmentation methods, the classifiers are giving very encouraging results. The classifier technique basically works in two-way fashion. In first stage, it separates amongst typical and anomalous while in next stage, it organizes the kind of variation from the norm in benevolent or dangerous tumor [31]. Below discussed are the two widely used classifiers.

- **Convolutional Neural Network (CNN)**

With the ongoing advancement in computational sciences and plethora of application in machine learning, the Convolutional neural networks (CNNs) have gained enormous recognition for variety of image processing applications including automatic medical image segmentation particularly. As image segmentation is a scheme to separate an image (containing valuable information) into various different parts (such as large sets of pixels, or commonly known as super-pixels). The foremost and fundamental objective of segmentation as well as computational image processing is to design and introduce as simpler algorithms as possible which can detect and process the images and analyze the information stored in them (such as features, objects, patterns, lesions and tumors etc.) in more robust, meaningful, understandable and convenient way. Using CNN architectures as a model such valuable information from medical images and radiographs can be readily extracted out with greater accuracy and improved performance with time.

CNN is a discriminative model which directly learns from annotated images without any prior knowledge [32]. CNN based frameworks involve the use of training dataset to instruct a network; and accordingly using these trained networks not only we can predict the class labels but also extract out the important features (such as patterns, edges, lines) and further train the other set of classifiers. Using this strategy, the patches of the information (as an input) is extracted out from the MRI images which are processed through convolution based filters and local sub-sampling to obtain the highly complex features (such as patterns, edges, lines) and help to yield the location and size of the tumors based on their corresponding computed class scores. Moreover, the CNN architectures have also advantage of automatic learning the complex features related to healthy tissues as well abnormal tissues (tumors) directly acquired from MRI images.

In order to contribute to the domain, Lang, Zhao and Jiya [33] proposed an architecture of convolutional neural networks to solve manual segmentation issues. Low resolution and noisy images such as CT scan and MRI are dealt easily by powerful classification features of CNN. The study presented three different CNN architectures of 5x5, 12x12 and 28x28 patch sizes. As a result of comparison of the three architectures proposed, it was found out that the architecture with 28\*28 input patch size was the most accurate one.

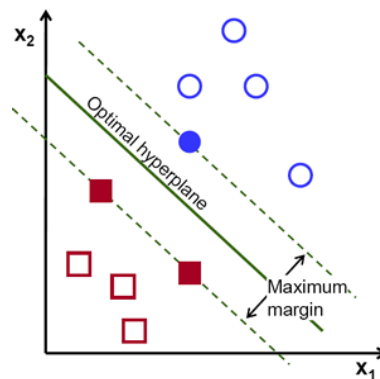
Similar architecture was adopted by Mengqiao, Yilie and Hao. They built a 22-layer deep learning architecture based on convolutional neural networks [34]. The architecture consisted of cascaded convolutional layers of size  $3 \times 3 \times 2$  instead of layers of size  $7 \times 7 \times 4$  as the effect of both is the same and the former has lesser weights. Besides convolutional layers, the architecture consisted of fully connected layers and the average pooling layers. The results were evaluated on BRATs 2015 dataset with the help of an online platform.

Recently, various machine learning as well as deep learning based approaches have been widely implemented in brain tumor segmentation based experimental and research studies after their highly dominative applications and commendable progressive success in numerous other image analysis domains, such as semantic segmentation [53] - [55], images classification [56] and objects detection [57]. Among these deep learning techniques, the newly developed CNNs based tumor segmentation method such as 2D-CNNs [58] - [61] and 3D-CNNs [62] - [64] achieved better results and overall performance compared to other reported techniques.

Most recently, Zhao et. al., developed a new deep learning based brain tumor segmentation system by involving the dynamic fusion of conditional random fields (CRFs) and fully convolutional neural networks (FCNNs) in an automated and robust unified framework. Their newly proposed method is capable of segmenting the brain images slice-by-slice, yielding better, spatially consistent and quite faster results than compared to other contemporary techniques such as image patch based segmentation [65].

- **Support Vector Machines (SVM)**

SVM has been considered as one of the consistent and best methods for classifying the features, patterns or objects present inside the images. Using SVM, the set of images are basically divided into two / various resultant classes. The classification is usually performed by finding the hyper-plane principle that freely differentiates the two classes perfectly as shown in Figure 3 below.



**Figure 3: Hyperplane Classification Principle.**

It constructs a hyper plane adopting a kernel function [13][35]. As presented in Figure.2, the feature vectors indicated on the left side of the main hyper plane belong to the class -1, whereas the feature vectors designated on the right side of the main hyper plane corresponds to the class +1.

The segmentation with SVM mainly depends upon the following phases (a) feature extraction from training image (b) the selection of SVM model (c) preparation of data-set (d) SVM training [34][36]. After performing the pre-processing and the training steps, the patterns (such as intensity and position) are

extracted out automatically from the test image which is consigned as an input to the SVM model. On the other hand, in the segmentation phase, the test feature vector is compared with the trained feature vector and based on the comparison the abnormal brain tissues (tumor) are segmented out from the consigned input image.

A. Kumar, B. Mahavir and Richika in their study [43] used Support vector machine, coupled with K-means clustering and Principle Component Analysis for extraction and classification of the tumor region inside the brain. In the methodology given the data including brain scans were trained using support vector machine while the tumor was segmented using k-means and PCA. The purpose of training the data through SVM classifier was to find out the class of the tumor detected. This approach had an accuracy of 96\% for tumor volume detection as depicted by the results. Besides segmenting the tumor region, the paper also gave a detailed information on K-means and PCA, along with the relation between them.

Similarly, G. Gupta and V. Singh also worked on SVM for classification of images, coupled with Fuzzy C-Means. Their approach was to first skull strip the input image, apply FCM to segment the image and finally use SVM to further classify the images which gave more enhanced and better results [44].

- **Segmentation with Unsupervised Learning**

Unlike supervised learning, in these algorithms we are not required to provide the prior knowledge, the algorithms are left on their own to find the insights from the given image.

More precisely, the image segmentation with unsupervised learning is a process in which an image is automatically divided into various different homogeneous sections based on their corresponding similarity measures. It is easier to segment using unsupervised learning methods as supervised increase the computational cost by demanding the machine to be trained first and then tested [52]. Clustering is one of the vital implementations of unsupervised learning, our study provides the review of 2 algorithms based on clustering i.e. Fuzzy C-Means and K-Means.

- **Fuzzy C-Means**

Clustering approach has been widely used in different computational domains such as machine learning, computer vision and image processing.

Besides, clustering technique has also recently progressed into various biomedical and healthcare applications predominantly for the detection of abnormal brain tissues (tumors) from radiographs acquired through magnetic resonance imaging (MRI) modality [37]. Dunn et al., introduced a clustering based Fuzzy c-means (FCM) approach, that was further enhanced by Bezdek et a., which greatly furthered the segmentation proficiency and was proved to be a better method compared previous reported techniques in the literature. Clustering with FCM is done in a way that it bifurcates one group of data into two or various different corresponding clusters, as presented in Figure 4.

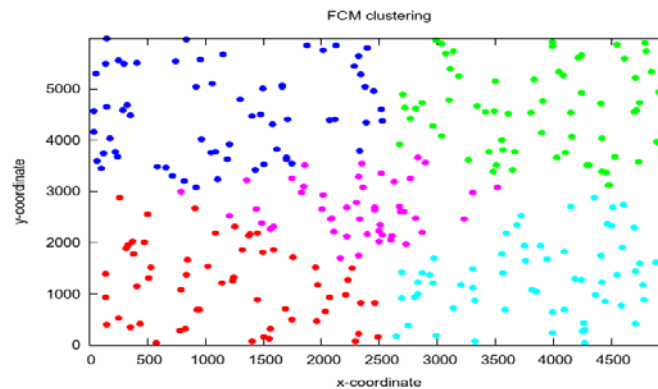


Figure 4: Clustering with FVM.

This method is basically executed for pattern recognition based applications. In this method, the membership value is allocated to each of the related data point that corresponds to each cluster center based on the distance between each of the cluster and data point. The possibility of the membership value towards any particular cluster is totally based on the closeness of the data to cluster center. The analysis of brain tumor segmentation suggests that by means of an unsupervised FCM clustering algorithm, the segmentation of brain tumor will be resulted in active cells, necrotic core and edema [4]. With respect to computational rate, the FCM algorithm will be more effective when the cluster center and membership value updating criterion is altered.

In 2016, Suganya and Shanthi reviewed Fuzzy C Means algorithm and its various applications in medical imaging, pattern recognition, bioinformatics and data mining [38]. The basic objective of the study was to analyze different algorithms based of FCM clustering and bring into light the merits and demerits of each.

Recently, Pham et. al., developed a novel clustering algorithm by integrating fuzzy entropy clustering for segmentation of the abnormal brain tissues (tumors) from MRI images. The newly suggested approach is basically based on the improved particle swarm optimization (PSO) algorithm, LHNPSO algorithm and kernelized fuzzy entropy clustering with modified fitness function to efficiently segment the brain tumors from MRI images by partially overcoming two serious issues, the first is sensitivity to noise and the other is INU artifact [66].

Kumar et. al., suggested a modified intuitionistic fuzzy c-means algorithm (MIFCM) to analytically solve the optimization problem and objective function using Lagrange method of undetermined multipliers. The proposed MIFCM method has allowed to segment brain MRI data by overcoming the limitations of noise and imprecise measurement [67].

Shanmuga Priya et. al., anticipated FCM based multilevel segmentation by combining fuzzy c-means, skull tripping and graph cut methods for detecting the tumor tissues and edema among brain MRI images. The clustering process has been enhanced by merging multiple kernels based on the spatial information to perform efficient segmentation [68].

- K-Means

The recognized clustering problem can be solved by K-means which is the simplest in unsupervised learning algorithms. There are four steps in a standard K-means algorithm, these are: initialization,

classification, computational and convergence condition. The process initializes by dividing a particular data set into stable positive K number of clusters so that k centroids can be defined as one for each cluster. As the results rely on the locations, therefore these centroids should be located in an elusive technique. One of the best ways, is to place them at a far distance from each other. The next step follows by selecting a point which is in acquaintance to a given data set and change it to the closest centroid. The first group age is complete when first step is finished which only happens when no any point exists the last. The outcome of first step results in K new centroids of the clusters which should be recalculated at the end of the first step. Once new K centroids are received, there is a need of a new connection between the same data set points and the nearest new centroid. Therefore, a loop is created which helps in analyzing the changed location of K centroids in each phase unless all centroids come to static. In other words, centroids do not move any more shown in Figure 5.

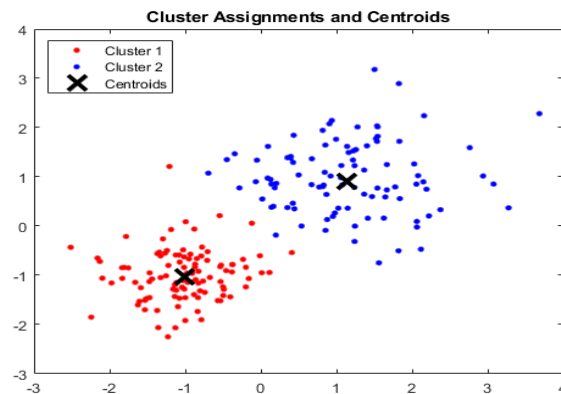


Figure 5: Clustering through K-means.

In [52] Vijay and Subhashini took into consideration the problem of labeling in image segmentation specially when it comes to automated brain tumor detection. They proposed a methodology that used morphological operations for preprocessing and traditional K-means technique with a slight modification that was to reduce the number of iterations required for proper clustering by suggesting the computed distance between a cluster center and data point under examination, which is stored in a data structure. This combination of K-means clustering, and morphological operations produced 95% accurate results on a sample space of 100 input images, as depicted in the results.

Dr. Patil, Dr. Jain and Pachpande proposed a computer aided application to segment tumor from the given MRI scans [25]. The segmentation idea adopted for the study worked with an amalgamation of K-means clustering and Fuzzy C means based clustering approaches. Four different modalities of images were tested for experiments and the results were generated based on parameters like, Mean Square Error (MSE), Contrast, Correlation, Max error, Area, etc. The study and results concluded that the method proposed was robust, accurate and time saving.

## 4 Comparative Studies

Where some researchers have been making remarkable contributions to the domain, there, on the other hand, other researchers are comparing different techniques produced and tested. In this regard, some worthy contributions are made in [40][45][51] and others.





















Said and Ibrahim [40] presented a study to compare various segmentation algorithms. A MATLAB toolbox was proposed in the paper to compare K-Means [25], Fuzzy C Means [38], Region-growing [41] and Otsu [42] techniques. Said and Ibrahim used Artificial Neural Networks (ANN) for classification and accuracy of the techniques compared. According to the results, K-Means and Fuzzy C Means algorithms had the same accuracy i.e. 96.7%, which was higher than the rest of the algorithms. Otsu had second highest accuracy of upto 90% whereas region growing method was least accurate.

Another comparative study produced by Suhasini and Vijaykumar [45] compared the studies like Support Vector Machine Classifier, Fuzzy C Means [46], K-means [47] Hybrid Clustering [48], Mathematical Morphology [49] and Integrated Bayesian Model [50] and more. These various techniques were experimentally compared based on the accuracy. Morphological filtering was regarded as the highest accuracy algorithm with an accuracy of 99% followed by Hybrid clustering, FCM and Bayesian. K-means clustering was also one of the highlights of the study as it had the greatest number of advantages in terms of implementation and usage easiness. Though morphological filtering had the most accuracy, but it required a high computational power as compared to the K-means which had minimum implementation hassle and second highest accuracy.

G. Rao and B. Srinivas in [51] contributed to the domain by contrasting Fuzzy C-Means and K-Means clustering techniques. Segmentation through FCM and K-Means was compared with respect to its Mean Square Error (MSE), Peak Signal to Noise Ratio (PSNR), Peak Time (PTime) and area calculation. Results of the study showed that FCM had a greater accuracy of approximately 93%, along with lower PTime in comparison to K-Means, which had about 76% accuracy only. accurate.

## 5 Discussion

Image Segmentation is one of the most fundamental concepts in computer vision. Segmentation aims to change/modify the representation of an image to extract meaningful information from it. This paper presents a comprehensive review of literature on various segmentation methods that are currently being worked upon for brain tumor segmentation. A number of algorithms and hybrid approaches for all these methods have been presented in Section III. Besides this, various automated and semi-automated techniques have also been discussed which on real time implementation will be able to contribute the excellence of computer technology to assist in the field of medical science. Figure 6 shows the results of a few algorithms discussed.

Technique	Input Image	Output Image
Skull Stripping and Watershed [23]		
Marker-controlled Watershed [72]		
Histogram multi-thresholding and Fusion [17]		
Convolutional Neural Network [34]		
Support Vector Machine [36]		
K-Means Clustering [52]		
K-Means, PCA, SVM [43]		
Thresholding [71]		
Hybrid clustering KFCM [48]		

**Figure 6: An overview of different brain tumor segmentation techniques.**

Despite of the huge contributions being made; none of systems has perfected to be accepted in real time clinical applications.

The study presented in the paper suggests that the supervised learning methods have better accuracy, but they are heavy in terms of computational power i.e. consume more computational cost, storage space and processing time while gradient based methods as well as unsupervised methods are accurate and require lesser resources. In other words, some methods have an easier implementation while others have a greater accuracy. One thing is achieved at the cost of another hence this domain still has a lot of area for further improvement. The researchers should aim at creating systems utilizing minimum use of resources but producing better results.

Though computer aided systems are difficult to implement and update, and it requires computer literates to operate such systems, but they are not as laborious as manual methods.

## 6 Conclusion and Future Perspectives

In this paper, a number of brain tumor segmentation algorithms are presented. It can be concluded from the study that these automated systems lack in factors like interoperability and easy handling of tools and hence are not yet clinically accepted. Also, for real life acceptance, these systems need to have more accuracy and must be able to calculate the volume of tumor to know the stage of the tumor. Moreover, they must also be able to classify multiclass tumors and estimate tumor progression.

This kind of developments in supervised learning architectures and/or the hybrids of supervised and unsupervised may aid in standardizing the current methods which will in turn help in clinical acceptance

of these automated systems, resulting in another major contribution of computer technology in medical science.

### CONFLICT OF INTEREST

The authors declare no conflict of interest.

### AUTHORS CONTRIBUTION

All authors contributed equally to this work. All authors wrote, reviewed, and commented on the manuscript. All authors have read and approved the final version of the manuscript.

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### LIST OF ABBREVIATIONS

MRI	=	Magnetic resonance imaging
CT	=	Computed tomography
GM	=	Gray matter
WM	=	White matter
CSF	=	Cerebrospinal fluid
WLSA	=	Watershed based lesion segmentation
RFC	=	Relative fuzzy connectedness
CNN	=	Convolutional neural network
FCNN	=	Fully convolutional neural network
ANN	=	Artificial neural network
CRF	=	Conditional random fields
SVM	=	Support vector machine
PCA	=	Principal component analysis
FCM	=	Fuzzy c-means
PCO	=	Particle swarm optimization
MIFCM	=	Modified intuitionistic fuzzy c-means algorithm
MSE	=	Mean square error
SNR	=	Signal to noise ratio
PSNR	=	Peak signal to noise ratio

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# Modified Self-Organizing Map Algorithm for Brain Tumor Detection and Analysis Using Magnetic Resonance Brain Images

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## ABSTRACT

medical image processing play an important role to help radiologists and support their decisions in diagnosis of the patient, magnetic resonance imaging (MRI) has ability to diagnosis the small details in the human body with a high resolution; in this paper, we propose modified self-organizing map algorithm (MSOM) for brain tumor detection and analysis using magnetic resonance brain images the significance of the (MSOM) algorithm is ability to detect tumor area in the magnetic resonance brain image (MRI) clearly with a high accuracy and best performance according of different values, the advantage of method proposed can segment and detect different types of MRI brain images FLAIR, T1 and T2-weight images with same performance and accuracy, the (MSOM) method start through input magnetic resonance brain image (MRI) and preprocessing applied to remove the noise from the image, applied modified self-organizing map (MSOM), applied tumor area, performance of the method, finally the applied of modified self-organizing map (MSOM) gave a best results us shown in the results.

**Keywords:** MRI, brain tumor detection, modified self-organizing map, accuracy values.

## 1 Introduction

At present, medical imaging, systems and medical image processing have been revolutionized by new technologies used based on advanced hardware and software architecture. There are many methods of medical imaging systems, all of which create visual representation of the human body and human organs for clinical analysis. Treatment and medical intervention. Because all systems of acquisition and treatment, medical imaging systems sometimes suffer from technical barriers such as device noise, as well as voluntary and involuntary patient movements during medical examination. Moreover, physical limitations are due to radiation sensitivity, magnetic fields and chemical products used [1]. All these factors can make it difficult to interpret and / or analyze this information and medical signals. BMRI, in particular, is one of the most commonly used techniques in imaging to visualize and analyze human head and components. However, identifying and detecting brain abnormalities remains a major challenge and an open search for improved medical diagnosis. Under these circumstances, CAD is very important for improving medical analysis and treatment. Brain tumor is one of the most common brain conditions in children and adults. Brain tumors are the cause of a quarter of cancer deaths in the world. Brain tumor is a group of cells that randomly grow inside or around the brain. There are two classes of tumors, the first

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is a non-cancerous tumor (benign) and the second is a cancerous tumor (malignant). Another abnormality in the brain called brain edema associated with brain tumors is very common and can occur and surround brain tumors [2]. Image processing algorithms and techniques provide tremendous help in this research and provide an additional opinion to improve the analysis and accurate diagnosis of radiologists. In literature, many researchers have proposed different methodologies of BMRI images to achieve the discovery of brain abnormalities with minimal human interaction.

Two fully-unsupervised methods to MR brain image segmentation using SOM-based strategies are recommended by Ortiz et al [3] segment T1-weighted images are determined by SOM (Self-Organizing Map) technique. The using the segmentation algorithm is still restricted to process T1-weighted images. Further, the sensitivity value manufactured by the algorithm requires improvement.

Logeswari and Karnan [4] reported brain tumor detection using a segmentation process based on Self-Organizing Maps (SOM). The segmentation process is curbed to segment T2weighted image sequences.

Logeswari and Karnan [5] offered brain tumor detection using segmentation process based on Hierarchical Self-Organizing Map (HSOM) technique, were T2 weighted images was segmented in an average time of 29.9708 seconds, which requires further minimization.

Yan Li and Zheru et al [6] performed MR brain image segmentation using SOM, which requires a reduction in MSE values.

Segmentation algorithms put forth by Guler et al [7], Ong et al [8], Alirezaie et al [9], Sun [10] and Jiang et al [11] have utilized unsupervised neural network technique namely, self-organizing maps (SOM).

Segmentation of T1-weighted images alone is completed employing an automated algorithm, which mixes stationary wavelets and Ayse Demirhan [12] created Self-Organizing Map (SOM).

In this paper, we proposed modified self-organizing map (MSOM) algorithm has ability to identify the tumor area using MR brain images with a best accuracy than all the methods created using self-organizing map (SOM) algorithm based on different values accuracy, nJaccard, nDice, sensitivity, specificity, recall and precision value.

## 2 Materials

Approximately 15FLAIR, T1.T2-weight brain images, we obtained from the different patients and different Age groups have been used in this paper, from the **Tianjin Medical Hospital University** details are listed below from patients:

1. The clinical image of the patient from the age of thirty-seven suffering from used meningioma.
2. The clinical image of the patient age of thirty-two suffering from and using high- quality astrocytes.
3. Images of patients with metastatic tumor and bronchial cancer, taken from Nouakchott hospital of oncology and used inventory, figure.1 shown example of different types of MR brain images with tumor.

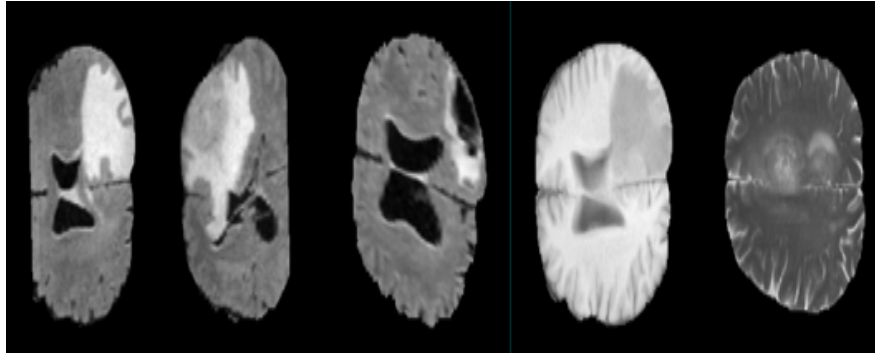


Figure.1. Flair, T1.T1 weight brain images with tumor

### 3 Methods

The methodology in this paper include five necessary stages as follows:

Stage1: Input MR Brain image (original image)

Stage2: pre-processing based on color quantization, tumor masking, and morphological cleaning.

Stage3: segmentation using modified self-organizing map (MSOM) algorithm

Stage4: detect tumor area or detect tumor alone in MR brain image

Stage5: analysis stage based on performance of the (MSOM) according of the different values product.

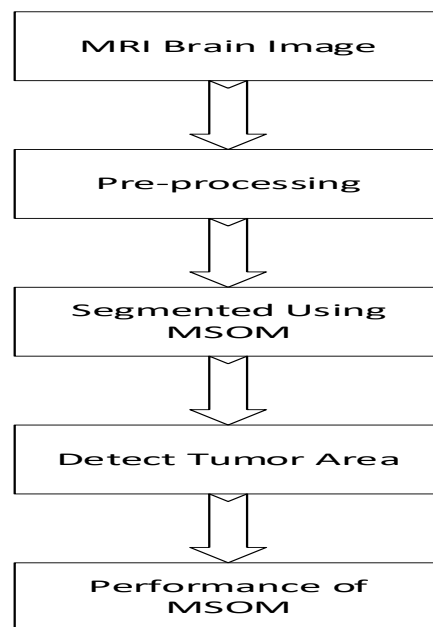


Figure.2. Flowchart explaining the (MSOM) algorithm framework

#### 3.1 Self-Organizing Map Algorithm

Self-regulating maps (SOMs) has been proposed by Willshaw and Von Der Malsburg to visualize various biological phenomena observed in animals. In particular, they have discovered that some brain regions develop structures with different regions, each of which has a high sensitivity to a particular type of input pattern.

The process behind such behavior is quite different because it is based on the rivalry between nerve units based on the principle called winner takes everything. During the training period, all units are excited with the same signal, but only one will produce the highest response. This module will automatically become a destination pool filter for this selected pattern. You will be introduced to the Kohonen model.

Kohonen SOM (also known as Kohonen network or Kohonen map) is simply represented as a two-dimensional map (for example, a square matrix or any other rectangle), but three-dimensional surfaces, such as a ball or torus, are also possible (with the existence of a suitable scale). In your case, always refer to a square matrix, where each cell is a receptive neuron characterized by a tangential weight with dimensions of input patterns:

$$X = \{\bar{x}_1, \bar{x}_2, \dots, \bar{x}_N\} \text{ where } \bar{x}_i \in \mathbb{R}^n \quad (1)$$

During both training and working phases, the winning unit is determined according to a similarity measure between a sample and each weight vector. The most common metric is the Euclidean; hence, if you consider a bi-dimensional map  $W$  with a shape  $k \times p$  so that  $W \in \mathbb{R}^{k \times p \times n}$ , the winning unit (in terms of its coordinates) is computed as follows:

$$u^* = \operatorname{argmin}_{k,p} \|W[k,p] - \bar{x}\|_2 \quad (2)$$

As explained before, it is important to avoid the premature convergence because the complete final configuration could be quite different from the initial one. Therefore, the training process is normally subdivided into two different stages.

During the first stage, the duration is usually about 10-20% of the total number of frequencies (let us call this values max), and the correction is applied to the winning unit and its neighbors (computed by adopting a decaying radius).

However, during the second stage, the radius is set to 1.0 and the correction is applied only to the winning unit.

In this way, a larger number of possible configurations can be analyzed, and the configuration associated with the least error is automatically determined. The neighborhood can have different forms. Can be square (in 3D maps closed, no longer exists), or, more easily, a radial basis function can be used based on a significantly degraded distance weight:

$$n(i,j) = e^{-\frac{\|u^* - (i,j)\|^2}{2\sigma(t)^2}} \text{ where } \sigma(t) = \sigma_0 e^{-\frac{t}{\tau}} \quad (3)$$

The relative weight of each neuron is determined by the  $\sigma(t)$ . The  $\sigma_0$  function is the initial radius and  $\tau$  is a time-constant that must be considered as a hyperactive parameter that determines the slope of the decaying weight. Suitable values are **5–10%** of the total number of iterations.

Adopting a radial basis function, it is not necessary to compute an actual neighbourhood because the multiplication factor  $n(i,j)$  becomes close to zero outside of the boundaries. A drawback is related to the computational cost, which is higher than a square neighbourhood is (as the

### 3.2 Morphological Operation

Morphological image processing consists mainly of nonlinear processes that can relate to the shape or Image morphology features, such as borders, Skeletons etc. In morphological techniques, the image is Investigation by hiding a small template or a form called the elements of structure and operations are based on this element, which identifies the area of interest or neighborhood around Pixels.

Mathematical morphology is based on set theory operations, which are defined between set of points of an image called object and the kernel called structuring element, these are some basic morphological operations:

**Dilation:** Morphological expansion is a process consists of Find the maximum between pixels belongs to Window or window. Stretching removes the existing pixel noise Object area with object size increased, here input image  $I$  with the size  $G \times H$  and structuring element  $B$  with the size  $K \times L$ , which defines the size of the window. Mathematically, it can be written as:

$$[I \oplus B](w, l) = \max[I(w - u, l - v) | (u, v) \in B] \quad (4)$$

**Erosion:** Morphological erosion is the operation, which consists of finding the minimum among the pixels belonging to the window. Erosion removes noise pixels, which are present in background with reducing size of object, here input image  $I$  with the size  $G \times H$  and  $SE B$  with the size  $K \times L$ , which defines the size of the window. Mathematically, it can be written as:

$$[I \ominus B](w, l) = \min[I(w + u, l + v) | (u, v) \in B] \quad (5)$$

### 3.3 Accuracy Value

Accuracy is the ability of the instrument to measure the accurate value. In other words, it is the closeness of the measured value to a standard or true value. The accuracy can be obtained by taking the small readings. The small reading reduces the error of the calculation. The accuracy of the system is classified in the following equation (6).

$$Accuracy = \left( \frac{k}{m \times n} \right) \times 100 \quad (6)$$

### 3.4 Dice Overlap Index (DOI) value

It is expressed with the help of the value of the Jaccard index  $J(A, B)$ . DOI identify the purpose of overlap of the input image ( $A$ ), as well the resulting segmented image ( $B$ ). DOI mentions calculating in the equation (7).

$$D(A, B) = 2 \times \frac{J(A, B)}{1 + J(A, B)} \quad (7)$$

### 3.5 Sensitivity value

(Also called the true positive rate, the recall, or probability of detection in some fields) measures the proportion of actual positives that are correctly identified as such (e.g., the percentage of sick people who are correctly identified as having the condition), It is stated in Equation (8).

$$OF = \frac{TP}{TP + FN} \quad (8)$$

### 3.6 Specificity value

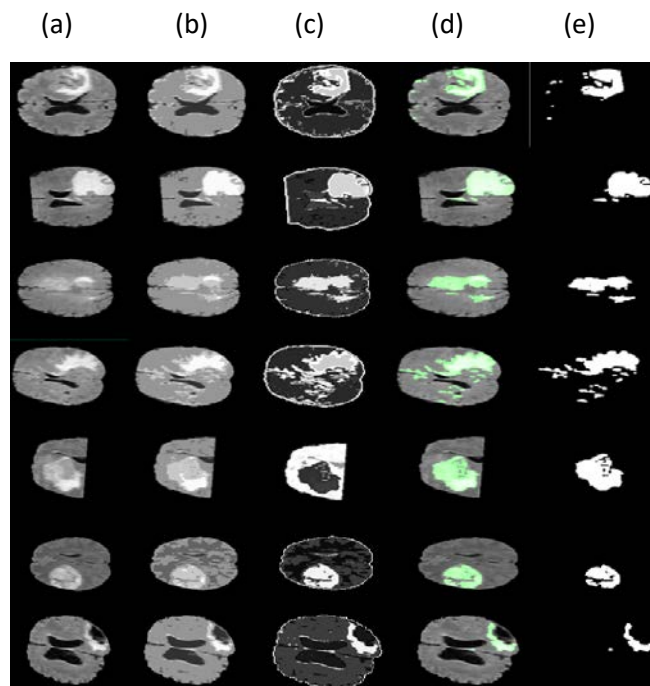
Also called the true negative rate) measures the proportion of actual negatives that are correctly identified as such (e.g., the percentage of healthy people who are correctly identified as not having the condition), specificity is shown in the Equation (9).

$$Specificity(\sigma) = \frac{TN}{TN+FP} \quad (9)$$

## 4 Results and Discussion

In this work, we divided the results of experiments to the two sections principally as follows:

*Section 1:* Results processing of seven MR brain images using modified self-organizing map (MSOM) algorithm, (MSOM) algorithm detected tumor area clearly using FLAIR, T1, and T2 weight MR brain image as shown in the figure .3.



**Figure.3 Results processing of the FLAIR, T1-T2 MR brain using modified self-organizing map (MSOM) algorithm**

Image (a): FLAIR, T1-T2 MR Brain image or original image

Image (b): Quantization

Image (c): segmentation using modified self-organizing map (MSOM)

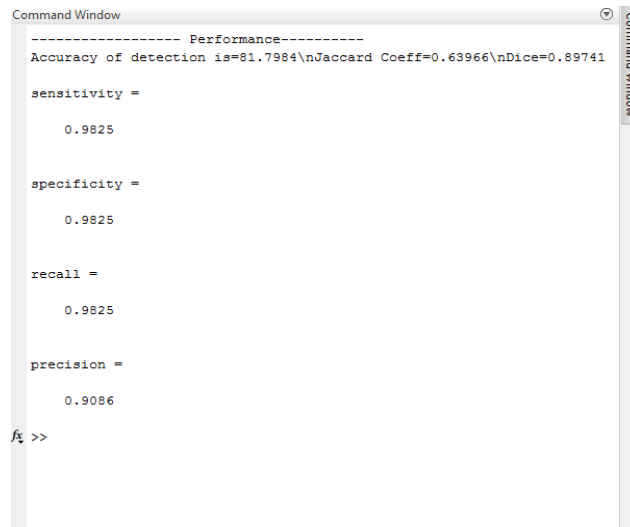
Image (d): Tumor area marked

Image (e): Tumor detection

The section 1 based on MR brain images processing to detect tumor area and it passed five important steps start it through input MR brain image and finished at detect tumor area applied (MSOM) algorithm,

the results of the processing and applied modified algorithm gave a best results and identify the tumor successfully using different types of MR brain images.

*Section 2:* Results performance of the modified self-organizing map (MSOM) algorithm, product the very good performance according of the values, the accuracy of detection is 81, 79%, nJaccard value is 63,966, Ndice is 89,741, sensitivity or true positive rate is 98,25%, specificity or true negative rate is 98,25%, recall is 98,25% and precision is 90,86% as describe in the figure4.



```
Command Window
----- Performance -----
Accuracy of detection is=81.7984\nJaccard Coeff=0.63966\nDice=0.89741

sensitivity =

    0.9825

specificity =

    0.9825

recall =

    0.9825

precision =

    0.9086

fx >>
```

**Figure.4 Performance of modified self-organizing map algorithm using values**

## 5 Conclusion

In this work, we applied method based on modified self-organizing map algorithm (MSOM) for brain tumor detection and analysis, the objective of the modification it is to improve the performance of the method, and the modification results gave a best results in terms of tumor detection using MR brain images as describe in the figure3 or performance of the modified self-organizing map (MSOM) algorithm, using different values as describe in the figure4, the results obtained from (MSOM) algorithm give the motivation to researchers in this area to continue the modification and improvement of the algorithm for more better results.

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