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Tracking Down Of Robust Cells From Computational Modeling of Tumors

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ABSTRACT

Now days we identified rapid growth in technical evolution, medical field is growing like anything. For making good sound we need cooperation from both hands. In this way, medical and engineering technology, both are joined together and developing new innovations in medical field. These innovations are enlightening the life of human by providing proper treatment. And still researchers are finding new things in the area of medical imaging. Recent bioengineering researchers involved in medical image segmentation algorithms to speed up the physician's diagnostic process. In this paper we are describing the Tracking Down Of Robust Cells From Computational Modeling Of Tumors. We implement the using the MATLAB base on the image processing. In this system we maintain the accuracy of the proposed system with help of neural network, deep CNN, Recursive least square dictionary learning algorithm.

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I. INTRODUCTION

Digital Image Processing:

What do we mean by Image Processing?
Image processing is important part of us and we continue to process images till the day we die .So even if this subject seems to be new, we have been subconsciously doing this, all these years, the human eye-brain mechanism represents the ultimate imaging systems.

Apart from our vision, we have another important trait that is common to all human beings. We like to store information, analyse it, discuss it with others and try to better it. This trait of ours is responsible for the rapid development of the human race.

Basic elements of image processing:

1. Image acquisition
2. Image storage
3. Image processing
4. Display

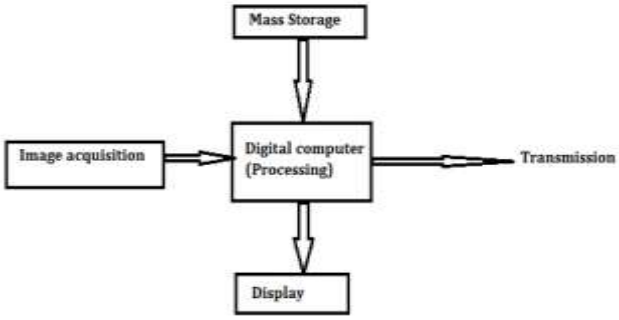

Figure 1: general flow of image processing

Image Acquisition:
Image achieved in image acquisition block. The general aim of image acquisition is to convert an optical image (real world data) into an array of numerical data which could be later manipulated on a computer. Image acquisition is achieved by suitable cameras. We use different cameras for different applications. If we need an x-ray image, we use a camera(film) that is sensitive to x -rays. If we want infrared image, we use cameras which senses the infrared radiations. For normal image(family picture etc) we use cameras(sensors) which are sensitive to the visual range.

Storage :
All images are essentially in analog form that is electrical signals convey luminance and color with continuously variable voltage. The cameras are interfaced to a computer where the processing algorithm are written. This is done by a frame grabber card. Usually a frame grabber card is printed circuit board or hard disk fitted to the host computer with its analog entrance port matching the impedance of the incoming image signal. The AD converter translates the video signals into digital values and a digital image is constructed.

Image Processing :
Image processing is a system which is connected to computer and which is most popular these days. Processing on digital image involve procedures that are usually expressed in algorithmic form due to which most image processing functions are implemented in software. The only reason for specialized image processing hardware is the need for speed in some applications or to overcome some fundamental computer limitations. The trend though is to merge general purpose small computers with image processing hardware. As stated in the earlier section, frame grabber cards play the important role of merging the image processing hardware with characterized by specific solutions.

Display:
A display is device which shows visual form of numerical values stored in a computer as an image array. Principal display devices are printer, TV monitor, and CRTs. Any erasable raster graphics display can be used as a display.

II. LITERATURE SURVEY

"A Generative Model for Brain Tumor Segmentation in Multi-Modal Images", Bjoern H. Menze, Koen Van Leemput, Danial Lashkari, this paper present a generative model for tumor appearance in multi-modal image volumes of the brain that provides channel-specific segmentations. We derive an estimation algorithm and demonstrate superior performance over standard multivariate EM segmentation. Unlike discriminative tumor segmentation methods, this model is applicable to any set of multi-modal image volumes, and is fully automatic. Further extensions of the model may consider structure of the tumor, or temporal evolution in longitudinal data sets.

"Fully Automatic Segmentation of Brain Tumor Images Using Support Vector Machine Classification in Combination with Hierarchical Conditional Random Field Regularization", Stefan Bauer, Lutz-P. Nolte, Mauricio Reyes, author presented a clinically-oriented method to segment 3D MRI images of brain tumor patients into healthy and tumor areas, including their individual subregions. For this, we propose to apply a hierarchical Support Vector Machine (SVM) -based classification and combine it with a CRF-based regularization in two stages. In contrast to those approaches, our method additionally returns all the tumor and healthy subregions while being faster in computation time. The accuracy of our automatic method lies in a similar range as the values reported for inter-observer variability of manual segmentations. However, the automatic method has advantages in longitudinal studies because the results are not biased subjectively. Due to the additional difficulty in subdividing the tumor region, Dice similarity coefficients for the individual tumor subregions are lower than for the gross tumor volume.

"Brain Tumor Segmentation based on Extremely Randomized Forest with High-Level Features", Adriano Pinto, Sergio Pereira, Higino Correia, J. Oliveira, in this paper, a new brain tumor segmentation method was proposed. This consisted of an Extra-Trees classifier based on local and context features defined on T1, T2 and Flair MRI sequences. The local features were defined as the intensity, mean intensity and gradients on the voxel under analysis, while the context features were defined as the mean intensities and gradients in nearby planes. It was also analyzed the creation of new pseudo-MRI sequences by computing the difference between sequences as well by applying nonlinearities on the original sequence. Local and context features were also computed on the pseudo sequences. This selection of classifier and features proved to be very competitive, specially in the segmentation of the core and enhanced regions of the brain tumor, allowing an overall seventh position among thirty-one methods on the test set of BraTS2013 challenge dataset.

"Brain Tumor Segmentation with Deep Neural Networks", Mohammad Havaei, Axel Davy, David Warde-Farley, in this paper, presented an automatic brain tumor segmentation method based on deep convolutional neural networks. He considered different architectures and investigated their impact on the performance. The high performance is achieved with the help of a novel two-pathway architecture (which can model both the local details and global context) as well as modeling local label dependencies by stacking two CNN's. Training is based on a two phase procedure, which we've found allows us to train CNNs efficiently when the distribution of labels is unbalanced. The resulting segmentation system is very fast, the time needed to segment an entire brain with any of these CNN architectures varies between 25 seconds and 3 minutes, making them practical segmentation methods.

III. BLOCK DIAGRAM

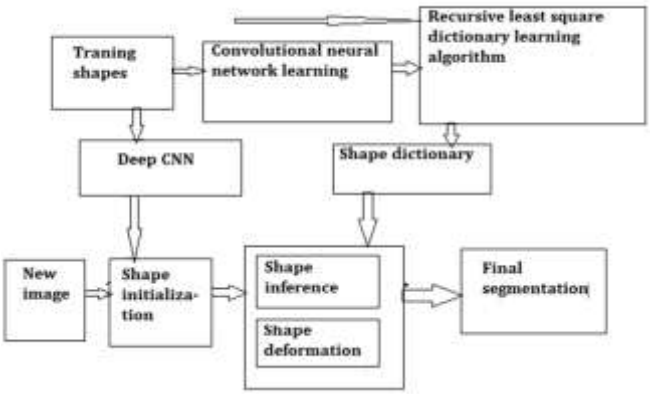

Figure 2: Block diagram of proposed framework

Training shapes:
In training shapes we are giving different images of blood cells which are affected by the brain tumor as the input to the algorithm. We are giving the images to the algorithm so that it stores the image for the further use . These images are used to compare with the new images which will be given to the algorithm at the testing stage. These images are further given to the convolutional neural network learning block and simultaneously to the deep CNN block for next process to be done.

Convolutional neural network:
CNNs are the continuation of the multi-layer perceptron. In the MLP, a unit performs a simple computation by taking the weighted sum of all other units that serve as input to it. The essence of CNNs is the convolutions. The main technique with convolutional networks that avoids or remove the problem of too many parameters is sparse connections. Every unit is not connected to every other unit in the previous layer, like in traditional neural networks.

Deep CNN:
Deep learning models are a class of machines that can learn a hierarchy of features by building high-level features from low-level ones. The convolutional neural networks (CNNs) are a type of deep models, in which trainable filters and local neighborhood pooling operations are applied alternatingly on the raw input images, resulting in a hierarchy of increasingly complex features. One property of CNN is its capability to capture highly nonlinear mappings between inputs and outputs.

Recursive least square dictionary learning algorithm:
We present the Recursive Least Squares Dictionary Learning Algorithm, RLSDLA, which can be used for learning over complete dictionaries for sparse signal representation. The Dictionary Learning Algorithms presented, for example ILS-DLA and K-SVD, update the dictionary after a batch of training vectors has been processed, usually using the whole set of training vectors as one batch. The training set is used iteratively to gradually improve the dictionary. The approach in RLS-DLA is a continuous update of the dictionary as each training vector is being processed.

Shape dictionary:
Shape dictionary is used to store the images which we have given as the training images for the comparison with the new images at the testing stage. We implement a dictionary learning algorithm (DLA) which is solved to the block-based image prediction problem.

Shape inference:
Shape inference is nothing but taking out the required part of the image. Here in our project we are taking out the infected part by recognizing through there shapes. The shapes of infected nucleus are separated from that of whole image.

Shape deformation:
In this step, we propose an efficient shape deformation method based on the chan-vese model for nucleus segmentation. The chan-vese model is formulated based on the well-known Mumford-shah functional, and consists of two region based data fitting terms one for foreground and the other for background and several regulation terms.

Algorithm Steps:
Step 1: Read Image
Step2: Convert to binary with thresolding.
Step 3: Remove noise and apply median filter.
Step 3: Dilate the Image
Step 4: Erode the Image
Step 5: Consider the large component.
Step 6: Fill Interior Gaps
Step 7: Remove Connected Objects on Border
Step 8: Smoothen the Segmented Object
Step 9: Outlined in the original image

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V. CONCLUSION

In this framework, we propose a novel nucleus segmentation framework using deep CNN and selection based sparse shape model. The approach starts with a deep learning based iterative region merging algorithm to initialize the contours, and thereafter alternately performs efficient bottom - up shape deformation and robust top-down shape inference to achieve correct nucleus segmentation. The proposed shape initialization method is robust to image noise and inhomogeneous intensity.

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